

# **STA305 Design and Analysis of Experiments Lecture Notes (2021 Summer)**

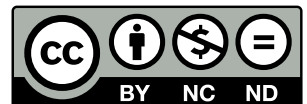
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July 31, 2021

## Preface

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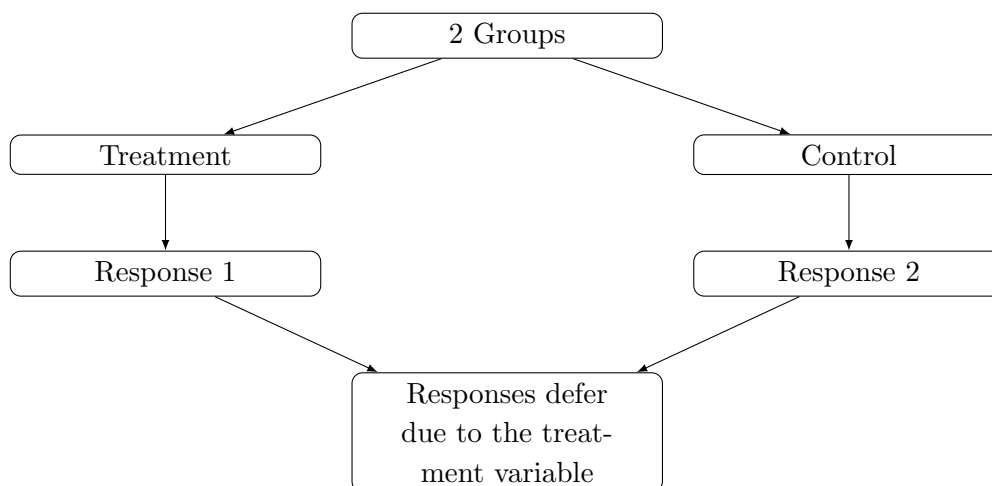
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# 1 Lec 1 Introduction to Experiments and Types of Data

## 1.1 General Outline of an Experiment

- An experiment is used to prove a scientific claim
- Simplest Setup: 2 groups
  - Apply a different Treatment to each group, i.e. Treatment and Control
  - Measure a variable of interest, i.e. Response
- Ideal situation: the groups are identical in every possible way, except for the treatment
- If the groups differ in the response, it must be due to the treatment variable.

A flowchart that generally explain this:

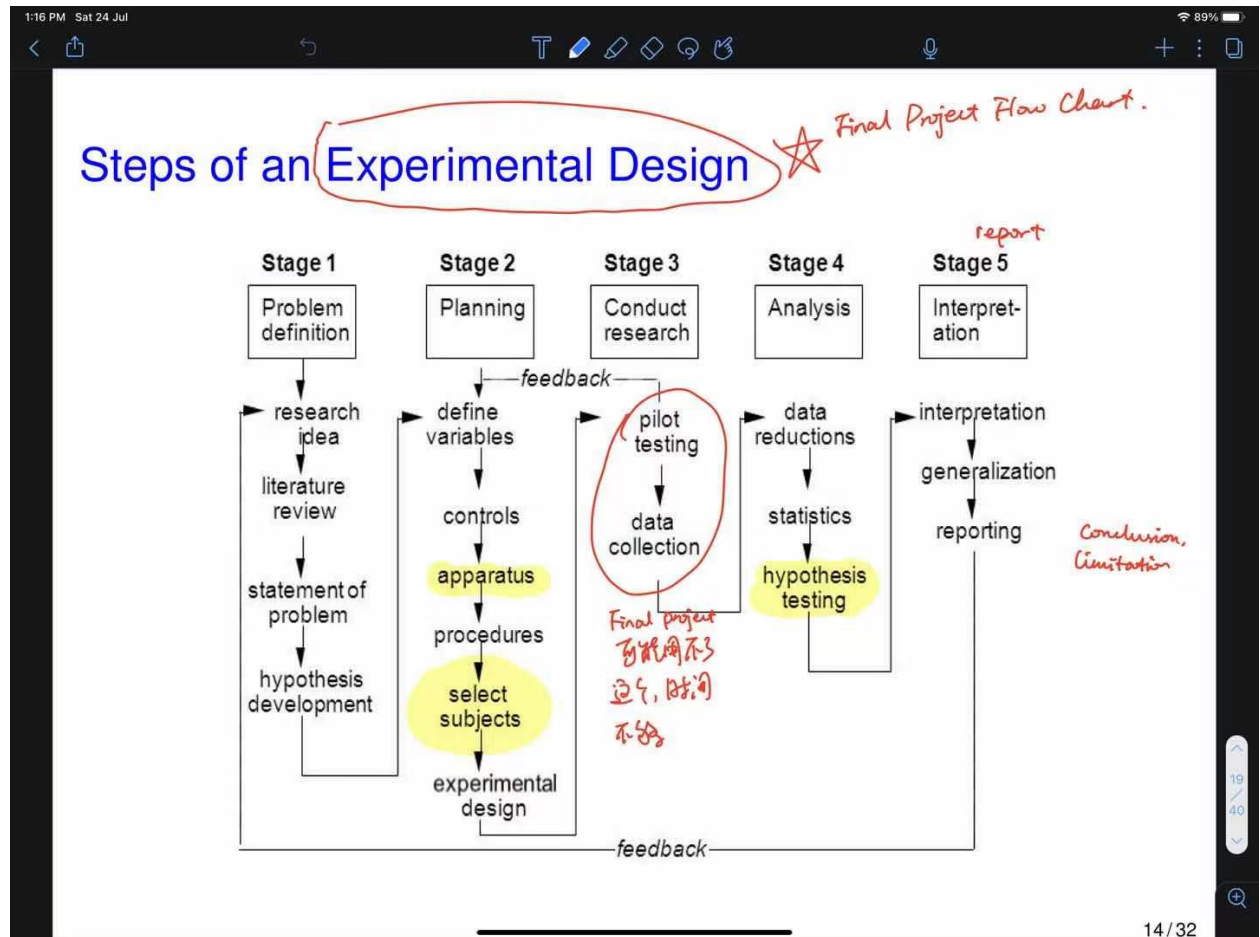


Example of different treatments: Medicine & Placebo

**Example 1.1.** A research team wants to test their effect of a new medicine on the Migraine. They have the **treatment group** of New Pill, and they have the **control group** of Placebo. Then they will test migraine sufferers. The **response** will be the pain level from 1 to 10 subjectively measured by patients themselves. The duration will be 2 hours after taking the pill. In this experiment, the **nuisance variables** are: Age, Sleep hours, Diet, Severity of migraines... The treatment has two levels: pill and placebo

## 1.2 Steps of an Experimental Design

This would be useful for Final Project



## 1.3 Definitions used in Statistical Inference

- Population: Set of units that we are interested in studying. E.g. Group of people, objects...
- Experimental Unit: The person or object on which the treatment is applied. Also called "case" or "Element" or "Subject" (When human unit).
- Sample: Subset of the population.
- Variable: A measured characteristic of a population unit. E.g., Age, weight, pain scale...
- Statistical Inference: Estimate, prediction, or generalization about population based information from a sample.

## 1.4 Experimental Study vs. Observational Study

Note this course we mainly focus on the experimental study. If you are interested in observational study you are recommended to take STA304.

### Experimental Study

- To assess the effect of a certain treatment or condition
- First randomly assign subjects to treatment, then control the rest. Randomization is important. Experimental data comes from experiments, so there are eligible reasons to conclude causation .
- Common in scientific and psychological research
- Between Subjects Design : Each experimental unit is assigned only one treatment
- Within Subjects Design : Each experimental unit is given all treatments

### Observational Study

- No randomization or controlling
- Data is collected as it comes (survey data), results from observational studies.
- Causation cannot be concluded from observational data since there may be other "confounding/lurking variables" (that were not controlled for) that is causing the relationship.
- Common in data mining, economic, and sociological research...

Note that an experiment may be hard to conduct due to ethical reasons, costs...

## 1.5 Types of Variables

There are two types of variables, which are  $X$  and  $Y$  such that  $Y$  denotes the response, that is measured by the researcher, and influenced by other variables.  $X$  is known as predictor or explanatory variable that affect the response variable, which is manipulated by researchers.

Variables can be:

- Quantitative: A variable is quantitative if it takes on numerical values for which arithmetic operations make sense. Ex: Height, marks, incomes... This type of variables can be discrete or continuous.

- Discrete: Variable can take on any one of a finite or countable list of values ( $\mathbb{Z}$ ) (there are gaps in the possible values). Ex: Number of heads in 10 tosses of a coin.
- Continuous: Variable can take on any value possible in an interval (uncountable). Ex: Time to complete a marathon... (There are infinitely many numbers in  $\mathbb{R}$ )
- Qualitative/Categorical variable: Categories, cannot be measured on numerical scales. Ex: Gender, smoking status (No, Used to, Yes)...

Variables can be measured as either qualitative or quantitative and further categorized as follows:

### Categorical:

1. Nominal: Categorize units into distinct classes.
  - Unordered categories
  - Numerical computations are not applicable
  - Ex: Gender, POSt, Favorite colors...
  - When controlled by researcher, called **Factor**
2. Ordinal: Ordered categories without natural units/distance metric
  - Natural ordering to the categories; not just the names of the categories differ
  - Professional rank, grades, satisfactions...

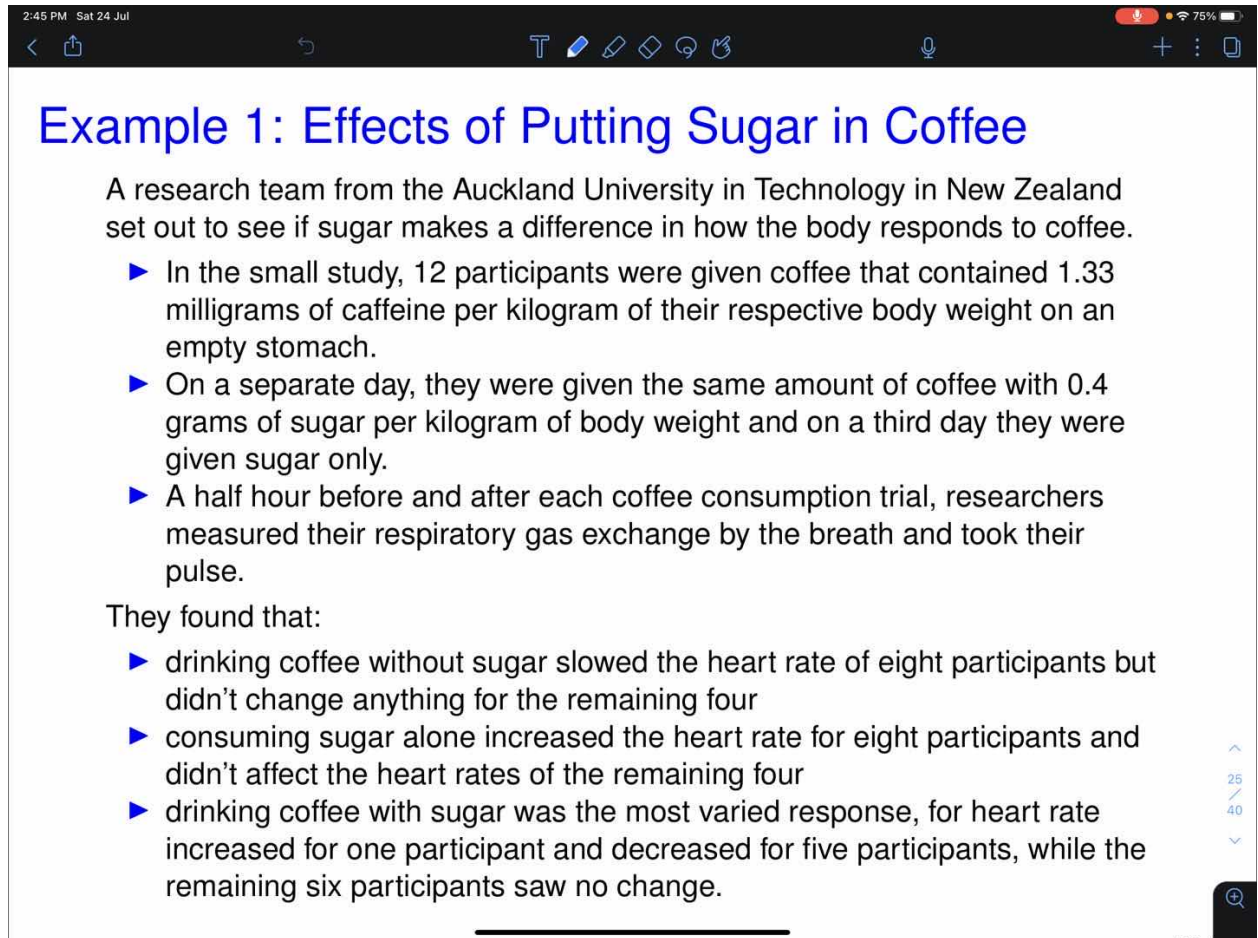
### Quantitative:

3. Interval: Numerical measurements which allow for degree of difference between values.
  - Distance is consistent but ratios are meaningless
  - Does not have a true 0 measure (0 is arbitrary)
  - Ex: Temperature ( $^{\circ}\text{C}$ ), dates...
4. Ratio: Numerical measurements on which a unique and non-arbitrary zero value exists
  - Ratios are meaningful, sensible to carry out multiplication/division
  - Ex: Temperature (K), length, time duration...

When an experiment has controls, call  $X$  a **factor** and its categories the **factor levels**. Single-Factor Experiment has one factor with levels  $\{a_1, a_2, a_3, \dots\}$ . Two-Factor Experiment has Two Factors (Factor A and B) with their own levels. A **treatment** is a combination of factor levels.



### Example 1.2. Example of Effects of Putting Sugar in Coffee



The screenshot shows a presentation slide with a dark header bar containing icons for navigation and a status bar at the top right showing the time as 2:45 PM on Saturday, July 24, and a battery level of 75%. The slide title is 'Example 1: Effects of Putting Sugar in Coffee' in blue. The text describes a research study from Auckland University of Technology in New Zealand. It lists three bullet points about the study design and findings. The slide is numbered 25 out of 40 in the bottom right corner.

## Example 1: Effects of Putting Sugar in Coffee

A research team from the Auckland University of Technology in New Zealand set out to see if sugar makes a difference in how the body responds to coffee.

- ▶ In the small study, 12 participants were given coffee that contained 1.33 milligrams of caffeine per kilogram of their respective body weight on an empty stomach.
- ▶ On a separate day, they were given the same amount of coffee with 0.4 grams of sugar per kilogram of body weight and on a third day they were given sugar only.
- ▶ A half hour before and after each coffee consumption trial, researchers measured their respiratory gas exchange by the breath and took their pulse.

They found that:

- ▶ drinking coffee without sugar slowed the heart rate of eight participants but didn't change anything for the remaining four
- ▶ consuming sugar alone increased the heart rate for eight participants and didn't affect the heart rates of the remaining four
- ▶ drinking coffee with sugar was the most varied response, for heart rate increased for one participant and decreased for five participants, while the remaining six participants saw no change.

1. For the above study, identify the following:
  - Population- All coffee drinkers
  - Experimental unit- A human participant
  - Variables (Which is response/predictor, quantitative/qualitative?)- *X: Drink with sugar or not, Y: Respiratory gas exchange and their pulse (Quantitative)*
  - Sample- 12 participants
  - A possible Statistical Inference- Generalization on effects of drinking coffee with sugar
2. Is the study observational or experimental? Justify. *The data were collected by an experiment, it has controls of giving specific amount of coffee and sugar. It is a within subjects design. Treatments are assigned.*
3. What are the factors and levels? Describe the treatments and how many there are? *Factor: Type of drink. Factor Levels (3 levels/treatments): Coffee, Coffee with sugar, sugar only (categorical)*

4. Name a confounding variable that could be present in this study. **The order of treatments applied**

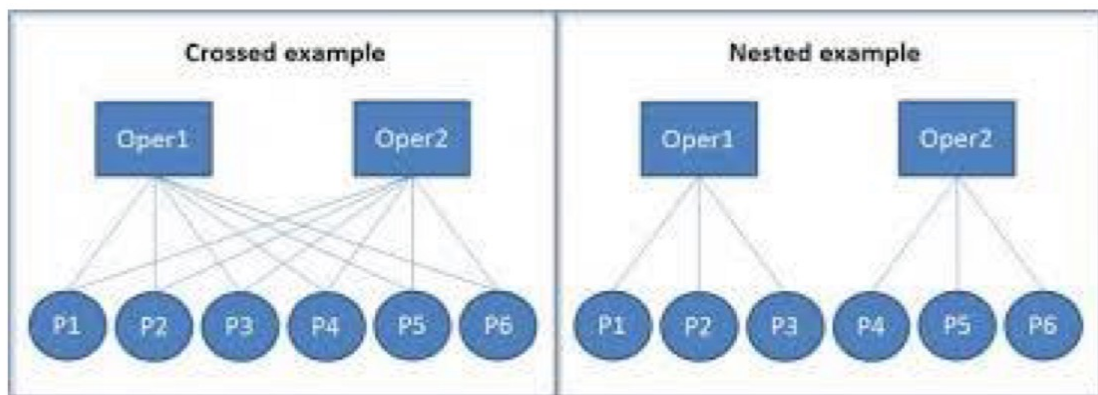
**Example 1.3. Effects of Teaching Style** Suppose you wish to conduct an experiment to see if teaching style (lecturer, facilitator, online technology) and the amount of lecture time (2 hours, 3 hours/week) affects the learning outcome of students in a large university course.

1. What are the factors and levels? **Teaching styles (Lecture, Facilitator, Online Tech); The amount of lecture time (2 hours, 3 hours)**
2. Treatments? **(Lecture & 2 hours, Lecture & 3 hours, Facilitator & 2 hours, Facilitator & 3 hours, Tech & 2 hours, Tech & 3 hours) Total 6 treatments**

## 1.6 Crossed vs. Nested Factors

When **all combinations** of factors are possible, the experiment is said to be fully **crossed**. Easier to analyze.

When each level of one factor occurs with a unique set of levels of the other factor, the design is called **nested**. Hierarchical Design.



## 1.7 Nuisance Variables

Variables other than the treatment condition that influence the response variable.

Ex: Teaching Styles:

- Different lecturers for each section

- Age of students
- Time of day for lecture section, etc...

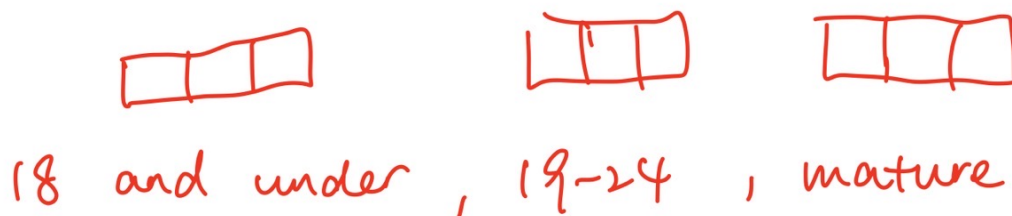
We use **Control, Blocking, Randomization, Replication** to deal with nuisance variables.

**Control:** Keep the nuisance variable constant throughout all treatment conditions (no longer a variable)

Ex: Teaching Styles: Same lecturer for each style; Same/Similar lecture times for each style...

**Blocking:** If the nuisance variable cannot be controlled, but can be observed, use Blocking to ensure each treatment has an equal amount of the variable.

Ex: Assign students to styles so each style has same age distribution.



**Randomization:** If the nuisance variable cannot be controlled or observed, use Randomization to spread out the variables and reduce the chance of confounding.

Ex: Teaching Styles: After blocking, randomly assign students to styles/sections.

**Replication:** Should not conduct the experiment with only one observation in each treatment. The resulting effect may be due to the experimental unit, and not the treatment. Cannot estimate variance within a specific treatment if it has only one observation. Use of replicates allows us to estimate **experimental error**, which is used to determine if we have statistically significant results.

Replication can occur at two different levels:

- **Treatment Level:** We take more than one observation in each treatment.
- **Experiment Level:** We wish to replicate the entire experiment to ensure results were not due other features inherent in the experiment, and so we can generalize to other populations.

Ex: Teaching Styles: Many students with similar characteristics (treatments) in each class; Another university could replicate the study.

## 1.8 Confounding

Nuisance variables that are systematically related to the treatments. Confounding variables can alter the effect of treatments even if non-systematic relationship to treatments, accidental effects can increase variability of responses and mask treatment effects.

Ex: Teaching Styles: Certain styles may be taught at different times of the day. For example, tech section always taught in night and lecture style during day. Differences cannot be uniquely credited to lecture styles.

Randomization decreases the chance that factors not accounted for in the design of the experiment will be confounded with the treatments.

## 1.9 Blinding

We use this when participants may be influenced once they know their treatment. **Placebo**.

**Double blind:** Both subjects and researcher are unaware of treatment assignment.

## 1.10 Balance

Assign the same number of experimental units to each treatment, call it **Balanced Design**. Balanced and unbalanced designs dealt with differently (calculations, etc...). Balanced design can become unbalanced easily- missing data, destroyed measurements, etc...

## 2 Lec 2 Review of Hypothesis Testing and One-Way ANOVA with 2 Levels

### 2.1 Single Factor Analysis with 2 levels

- 1 Factor with 2 levels (2 different treatments, treatment/control, etc. )
- 2 treatments in total
- Want to determine if there is a “treatment effect” (if response varies by treatment)
- Test for differences in means of the 2 populations: sample means are not enough; account for variability within each treatment group

### 2.2 Review of Hypothesis Testing

**Definition 2.1.** A Hypothesis Test is a formal statistical test that is performed to decide whether a statement about a set of parameter(s) is reasonable (to make an inference about the value of a parameter and how it relates to a specified/hypothesized numerical value).

#### Elements of a Hypothesis Test

1.  $H_0$  is Null Hypothesis and  $H_a$  is Alternative Hypothesis.
2. Test Statistic: A statistic (function of the data) that involves the parameter value and has a known distribution under  $H_0$ .
3. Distribution under  $H_0$ : Distribution of test statistic assuming that  $H_0$  is true
4. P-value: Probability of observing a test statistic as or more extreme (more contradictory to  $H_0$  i.e. favorable to  $H_a$ ) than already observed if the null hypothesis is true. Gives a measure of strength of evidence against  $H_0$ . Small P-value, test statistic value is unlikely if  $H_0$  is true (contradiction). Large p-value, test statistic value is likely if  $H_0$  is true (No contradiction)
5. Conclusion: Reject  $H_0$  (and favour  $H_a$ ) or Fail to Reject  $H_0$ . This leads to a practical conclusion about the population(s)/parameter value(s). When  $H_0$  is rejected, we say the test is statistically significant, or there is a statistically significant difference.

### 2.3 Review: Two Sample T-Tests

**Definition 2.2.** Interested in comparing two population means when we have small sample sizes. Suppose we have  $X_1, \dots, X_{n1}$  a random sample from population 1 and  $Y_1, \dots, Y_{n2}$  a random sample from population 2.

$H_0$  is  $\mu_1 - \mu_2 = D_0$ , note that this  $D_0$  is usually 0 but not always (When treatment means are equal). We have  $H_a : \mu_1 - \mu_2 \neq D_0$  OR  $\underbrace{H_a : \mu_1 - \mu_2 < D_0 \text{ OR } H_a : \mu_1 - \mu_2 > D_0}_{\text{One-tail test}}$

**Assumptions:**

1. Two samples are *iid* from Normal populations
2. Two samples are independent from each other

The test statistic is

$$t = \frac{(\bar{x} - \bar{y}) - D_0}{se(\bar{x} - \bar{y})}$$

where  $se$  is the standard error: estimated standard deviation.

## 2.4 Pooled T-Test

**Definition 2.3.** Applicable when population variances can be assumed to be equal. I.e. Population 1:  $X_1, \dots, X_{n1} \stackrel{iid}{\sim} N(\mu_1, \sigma_1^2)$  and Population 2:  $Y_1, \dots, Y_{n2} \stackrel{iid}{\sim} N(\mu_2, \sigma_2^2)$  such that  $\sigma_1 = \sigma_2$ .

**Assumptions:**

1. Two samples are random samples from the target populations and the samples are independent from each other.
2. The populations have equal standard deviations ( $\sigma_1 = \sigma_2$ )
3. Two samples are (approximately) Normal

If we can assume that the variances are equal, we use the **pooled estimate of the variance** :

$$s_p^2 = \frac{\overbrace{(n_1 - 1) s_1^2}^{\text{df of } s_1^2} + \overbrace{(n_2 - 1) s_2^2}^{\text{df of } s_2^2}}{\underbrace{n_1 + n_2 - 2}_{\text{total df}}}$$

**Hypothesis for Pooled T-Tests:**

$H_0 : \mu_1 - \mu_2 = D_0$  vs.

1.  $H_a : \mu_1 - \mu_2 \neq D_0$  OR
2.  $H_a : \mu_1 - \mu_2 < D_0$  OR
3.  $H_a : \mu_1 - \mu_2 > D_0$ .

### Test Statistic for Pooled T-Tests:

$$t = \frac{(\bar{x} - \bar{y}) - D_0}{\sqrt{s_p^2(\frac{1}{n_1} + \frac{1}{n_2})}} \sim t_{n_1+n_2-2} \text{ under } H_0$$

1.  $p = 2P(t_{n_1+n_2-2} > |t|)$  OR
2.  $p = P(t_{n_1+n_2-2} < t)$  OR
3.  $p = P(t_{n_1+n_2-2} > t)$

Conclusion: Reject  $H_0$  if  $p < \alpha$  and failed to reject otherwise.

## 2.5 Welch's T-Test Using Satterthwaite Approximation

**Definition 2.4.** Used when population variances are not equal or inconclusive evidence about equality of variances.

**Assumptions:** Two samples are random from the target populations and the samples are independent from each other; Two samples are Normal.

### Test Statistic for Welch's T-Test

$$t = \frac{(\bar{x} - \bar{y}) - D_0}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}} \sim t_\mu \text{ under } H_0$$

where

$$\mu = \frac{(s_1^2/n_1 + s_2^2/n_2)^2}{\frac{(s_1^2/n_1)^2}{n_1-1} + \frac{(s_2^2/n_2)^2}{n_2-1}}$$

The degree of freedom is calculated by Satterthwaite Approximation. Note that  $\mu$  may not be an integer, round down to the nearest integer.

### Example 2.5. Example of Arthritis Clinical Study

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Example: <sup>关节炎</sup> Arthritis Clinical Study

A clinical trial is conducted to determine the effects of auranofin therapy on the treatment of rheumatoid arthritis. The initial level of pain was measured for each of 293 patients who suffer from rheumatoid arthritis and then each was randomly assigned to receive auranofin pills or a placebo pill (3 mg to be consumed twice daily). After six months, the level of pain is once again measured for each patient to see if there is an improvement.

Question of Interest: <sup>one-sided</sup> Does auranofin therapy help to improve arthritis pain? <sup>two-sided. Is there a difference between A or B that relates to ...?</sup>

- ▶ Define the experiment: identify the factor(s) and levels, response variable(s), etc.
- ▶ State the appropriate hypotheses to test the question of interest

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There are 293 patients who suffering from RA, they are firstly measured of the initial pain. Then 146 are taking Auranofin and 147 are taking Placebo. After 6 months, we measure their pains and compare. There is a repeated measurement which is before pain and after pain. This is a between subjects design. The factor of the experiment is treatment (medicine). It has 2 levels that are Auranofin and Placebo. The responses are Pain before and Pain after 6 months. The researcher will compare the difference.  $\mu_A$  = mean difference in pain (before-after) for Auranofin users.  $\mu_P$  = mean difference in pain (before-after) for placebo users.

Question: Does the Auranofin therapy improve the pain? i.e., Is  $\mu_A > \mu_P$ ? Note that  $\mu$  is a mean difference, not the pain level!



The R Output is following:

```
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< [share] [undo] [text] [draw] [eraser] [laser] [undo] [redo] [microphone] [plus] [three dots] [device]

R Code

# Read in csv file
> arthdata <- read.csv("arthritisdata.csv")
> arthdata
  Treatment Baseline Month6
1  TAuranofin      3      1
2   Placebo      2      2
3  TAuranofin      3      2
.
.
.
292  Placebo      3      4
293 TAuranofin      3      4

# Attach data to R's search path
> attach(arthdata)

# Create Response Variable: Difference in Pain
> DiffinPain <- Baseline-Month6
> DiffinPain
[1] 2 0 1 0 0 0 -1 -1 -1 . . .

# Difference in Pain for each treatment
> DiffinPainA <- DiffinPain[Treatment=="TAuranofin"]
> DiffinPainP <- DiffinPain[Treatment=="Placebo"]

# Sample sizes, means, and variances by treatment group

> tapply(DiffinPain, Treatment, length)
TAuranofin Placebo
146         147
<-- n_A n_P

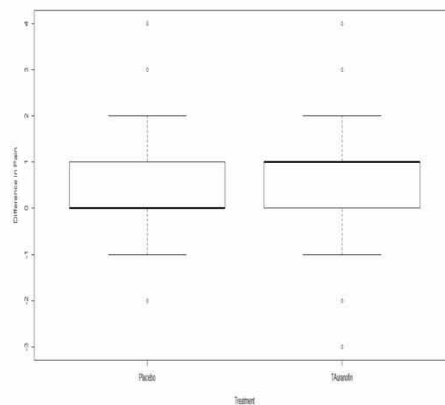
> tapply(DiffinPain, Treatment, mean)
TAuranofin Placebo
0.6575342 0.3401361
       $\bar{y}_A$   $\bar{y}_P$ 

> tapply(DiffinPain, Treatment, var)
TAuranofin Placebo
1.1784601 0.9383096
       $s_A^2$   $s_P^2$ 

> tapply(DiffinPain, Treatment, length) > tapply(variable, group, stat)
```

## Boxplots

```
# Side by Side boxplots by treatment  
> boxplot(DiffinPain ~ Treatment,  
          xlab="Treatment", ylab="Difference in Pain")
```



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## R Code: Pooled t-test

$\sigma_A^2 = \sigma_P^2$   
 $\text{response} \sim \text{factor}$   
 alternative = "greater"

```

1 > t.test(DiffinPainA, DiffinPainP, var.equal=TRUE)

# OR (same output as above)
2 > t.test(DiffinPain ~ Treatment, var.equal=TRUE)

Two Sample t-test

data: DiffinPainA and DiffinPainP

t = 2.641, df = 291, p-value = 0.008713

alternative hypothesis: true difference in means
is not equal to 0

95 percent confidence interval:
0.08086399 0.55393239

sample estimates:
mean of x mean of y
0.6575342 0.3401361
  
```

$\bar{y}_A$        $\bar{y}_P$

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Using hand calculation:

Pooled T-Test:  $H_0 : \mu_A = \mu_P$  vs.  $H_a : \mu_A > \mu_P$ , we have data:

Treatment	$n$	$\bar{y}$	$s^2$
Auranofin	146	0.6575	1.1785
Placebo	147	0.3401	0.9383

With the data above, we have

$$s_{\text{pooled}}^2 = \frac{145(1.1785) + 146(0.9383)}{293 - 2} \doteq 1.0660$$

$$t = \frac{(\bar{y}_A - \bar{y}_P) - 0}{\sqrt{s_{\text{pooled}}^2 \left( \frac{1}{n_A} + \frac{1}{n_P} \right)}} = \frac{0.6575 - 0.3401}{\sqrt{1.0660 \left( \frac{1}{146} + \frac{1}{147} \right)}} \doteq 2.64 \sim t_{291} \text{ under } H_0$$

By using T-Table:  $p = P(t_{291} > 2.64) = 0.0041 < 0.05$ .

Since  $p < 0.05$ , we reject  $H_0$ . Strong evidence to conclude that Auranofin therapy helps improve pains.

Using R, we can get the exact p-value that the 2 sided  $p = 0.0087$ , so the exact p value will be  $p = P(t_{291} > 2.64) = \frac{0.0087}{2} = 0.00435$ . The conclusion still holds.

## 2.6 Analysis of Variance (ANOVA)

**Definition 2.6.** Analysis of Variance (ANOVA) is a collection of statistical models and procedures for comparing factor level means in a factor. We wish to test if there is a statistically significant difference between the factor level/group means.

One-Way ANOVA: One factor; Two-Way ANOVA: Two factors, etc.

It's called "Analysis of Variance" because we compare the **within-group** variance to the **between-group** variance.

**Main Idea:** If the variation between groups is significantly bigger than the variation within groups, there is a statistically significant difference in the group means.

Note: An ANOVA is equivalent to using a linear regression model with categorical predictors (with indicator variables).

## 2.7 Recall: Multiple Linear Regression

$$Y_i = \beta_0 + \beta_1 X_{1,i} + \beta_2 X_{2,i} \dots + \beta_{p-1} X_{p-1,i} + \epsilon_i \text{ for } i = 1, 2, \dots, n$$

where

- $Y_i$ : Response for the  $i$ th case (Quantitative variable)
- $X_{1,i} \dots$ : Predictors for  $i$ th case (Quantitative or categorical)
- $\epsilon_i$ : Error term for the  $i$ th case, where  $\epsilon_i \stackrel{iid}{\sim} N(0, \sigma^2)$
- $\beta_0, \beta_1, \dots, \beta_{p-1}$ : Regression coefficients/parameters,  $\beta_0$ : Intercept
- $n$ : Number of cases/sample size

Matrix form:  $Y = X\beta + \epsilon$ ; where

- $Y$  is an  $(n \times 1)$  vector
- $X$  is  $(n \times p)$  matrix, each column contains values for each predictor. If model uses intercept, first column of  $X$  is all 1s.

- $\beta$  is a  $(p \times 1)$  vector of regression coefficients
- $\epsilon$  is a  $(n \times 1)$  vector of error terms

**Example 2.7.**  $y = \begin{pmatrix} y_1 \\ \vdots \\ y_n \end{pmatrix}$ ,  $X = \begin{pmatrix} 1 & x_{11} & x_{21} & \dots & x_{p-1} & 1 \\ 1 & x_{12} & x_{22} & \dots & x_{p-1} & 2 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ 1 & x_{1n} & x_{2n} & \dots & x_{p-1} & n \end{pmatrix}$ ,  $\beta = \begin{pmatrix} \beta_0 \\ \vdots \\ \beta_{p-1} \end{pmatrix}$ ,  $\epsilon = \begin{pmatrix} \epsilon_1 \\ \vdots \\ \epsilon_n \end{pmatrix}$

## 2.8 Indicator Variables

**Definition 2.8.** A factor has  $a$  levels, when using a model with intercept, use only  $(a-1)$  indicators. The level without an indicator is the default or reference level. For  $k = 1, 2, \dots, a-1$

$$I_{k,i} = \begin{cases} 1, & \text{if } i\text{th case belongs in factor level } k \\ 0, & \text{otherwise} \end{cases}$$

**Example 2.9. Linear Regression Model for Arthritis Example**

$$Y_i = \beta_0 + \beta_1 I_{A,i} + \epsilon_i \text{ for } i = 1, 2, \dots, n$$

where  $Y_i$  = difference in pain after 6 months for  $i$ th patient.

$$E(Y_i) = \begin{cases} \beta_0 + \beta_1, & \text{if } i\text{th patient received Auranofin (1)} \\ \beta_0, & \text{if } i\text{th patient received placebo (0)} \end{cases}$$

Therefore,  $\beta_0$  is the mean difference in pain for placebo group,  $\beta_1$  is the difference in the mean difference of pain (response) between Auranofin and Placebo groups.

$\beta_1 = 0$ : No difference

$\beta_1 > 0$ : Mean in Auranofin group  $>$  Placebo Group

$\beta_1 < 0$ : Mean in Auranofin Group  $<$  Placebo Group

## 2.9 One-Way ANOVA with 2 levels

When there are 2 factor levels (comparing 2 means like in Arthritis Example), use t-test to test if  $\beta_1 = 0$ . Alternatively, use F-Test: Can be used to compare many factor level means. T-Test using linear regression is equivalent to F-Test using ANOVA when comparing 2 means.  $t^2 = F$ .

## Example 2.10. Example of Arthritis Clinical Study- Linear Regression Model

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### Example: Arthritis Clinical Study - Linear Model

Test the question of interest using the linear model approach.  
Include all the steps in the hypothesis test and a practical conclusion.

```
# Regression Model with intercept
> arth.regmodel <- lm(DiffinPain ~ Treatment)
> summary(arth.regmodel)
```

Call:  
lm(formula = DiffinPain ~ Treatment)

Residuals:

Min	1Q	Median	3Q	Max
-3.6575	-0.6575	-0.3401	0.6599	3.6599

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.34014	0.08484	4.009	7.74e-05 ***
TreatmentTAuranofin	0.31740	0.12018	2.641	0.00871 **

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.029 on 291 degrees of freedom  
Multiple R-squared: 0.02341, Adjusted R-squared: 0.02005  
F-statistic: 6.975 on 1 and 291 DF, p-value: 0.008713

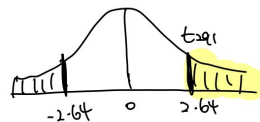
Note: R automatically makes the first level alphabetically/numerically as the default (no indicator).

*Handwritten notes:*  
- "linear model" under `lm`  
- "response ~ Factor" under `Treatment`  
- "placebo → default" and "TAuranofin" with an arrow pointing to the treatment level in the coefficients table.  
- "Two sided p-value" with an arrow pointing to the p-value 0.00871.  
-  $H_0: \beta_1 = 0$  and  $H_a: \beta_1 \neq 0$  with an arrow pointing to the p-value.

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Hand Calculation:

We have  $H_0 = \beta_1 = 0$ ,  $H_a = \beta_1 > 0$ , according to R output we have  $t = 2.64 \sim t_{291}$  under  $H_0$ .



$p = P(t_{291} > 2.64) = \frac{0.00871}{2} \doteq 0.0044 < 0.05$ . So, we have strong evidence to conclude that the Auranofin is effective in reducing pain.

## 2.10 Variation and Sum of Squares

When we apply a treatment, we consider a possible treatment effect. In the two sample case, it is the difference between two groups in the population, i.e.,  $\mu_1 - \mu_2$ . This difference may not be due to the treatment effect alone, but experimental error as well.

Generally,

$H_0 : \mu_1 = \mu_2 = \dots = \mu_a$  (No difference in group means,  $a$  groups in total)

$H_a : \mu_i \neq \mu_j$  for at least one  $i \neq j$  where  $i, j = 1, 2, \dots, a$  (Difference in group means)

$$X = \frac{\text{variation between groups}}{\text{variation between subjects, within groups}}$$

If  $X > 1$ , then more evidence we have a difference in group means (more significant).

## 2.11 One-Way ANOVA Notation

- One factor, Factor A with  $a$  levels
- $j$  denotes the group/level and  $i$  denotes the position of the subject within that level/group ( $j = 1, \dots, a; i = 1, \dots, n_j$ )
- $y_{ij}$  denotes the  $i$ th observation from the  $j$ th group ( $i$  is the index,  $j$  is the group/level)
- $n_j$  is the number of subjects in the  $j$ th group
- $N = \sum_{j=1}^a n_j$  is the total sample size
- $\bar{Y}_T$  is the grand sample mean regardless of group
- $\bar{y}_j$  is the sample mean in the  $j$ th group

### Definition 2.11. Total Sum of Squares

SST measures the total sample variability (total deviation from the grand mean):

$$SST = \underbrace{\sum_{j=1}^a \sum_{i=1}^{n_j}}_{\text{All observations}} (y_{ij} - \bar{y}_T)^2$$

- $(N - 1)$  df
- $SST = SSReg + SSE$

**Definition 2.12. Sum of Squares for A**

Sum of Squares for Factor A (SSA) measures the variability of the factor level means:

$$SSA = \sum_{j=1}^a \sum_{i=1}^{n_j} (\hat{y}_{ij} - \bar{y}_T)^2 = \sum_{j=1}^a n_j (\bar{y}_j - \bar{y}_T)^2$$

Note that  $\hat{y}_{ij}$  is the estimated observation,  $\bar{y}_T$  is the grand sample mean. It has  $(a-1)$  df, df=number of parameters in the model -1. It's normally called *SSA* or *SSReg*.

**Definition 2.13. Residual/Error Sum of Squares**

SSE measures the variability that is left unexplained by the model:

$$SSE = \sum_{j=1}^a \sum_{i=1}^{n_j} (y_{ij} - \hat{y}_{ij})^2 = \sum_{j=1}^a \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_j)^2$$

Note that it has  $(N-a)$  df, df=number of data - number of parameters in the model. SSE decreases as more predictors are added to the model. We want SSE to be low!

**2.12 One-Way ANOVA Table**

Note that SS refers to Sum of Squares, MS refers to Mean Square =  $\frac{SS}{df}$

ANOVA Table			
Source	df	SS	MS
Group/Regression	$a - 1$	$SSA$	$\frac{SSA}{a-1} = MSA$
Error	$N - a$	$SSE$	$\frac{SSE}{N-a} = MSE$
Total	$N - 1$	$SST$	

$$MSE = \frac{\sum_{j=1}^a \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_j)^2}{N - a} = \frac{\sum_{j=1}^a (n_j - 1) s_j^2}{N - a} = s_{\text{pooled}}^2$$

*SSA* is often called “between group” SS: measures the variability of observations between factor levels

*SSE* is often called “within group” SS: measures the variability of observations within factor levels.



## 2.13 F-Test

$$F = \frac{MSA}{MSE} \sim F_{a-1, N-a}$$

### Hypothesis Test Using ANOVA

Hypothesis:  $H_0 : \mu_1 = \mu_2 = \dots = \mu_a$  vs.  $H_a : \text{at least one of the means } \mu_j \text{ differs from others; for } j = 1, 2, \dots, a.$

Test Statistic and its Distribution under  $H_0$ :

$$F = \frac{MSA}{MSE} = \frac{\frac{SSA}{a-1}}{\frac{SSE}{N-a}} \sim F_{a-1, N-a} \text{ under } H_0$$

P-value:  $p = P(F_{a-1, N-a} > F)$

### Example 2.14. Arthritis Clinical Study-ANOVA

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## Example: Arthritis Clinical Study -ANOVA

Test the question of interest using ANOVA.

```
# Regression Model with intercept
> arth.regmodel <- lm(DiffinPain ~ Treatment)

# ANOVA table
> anova(arth.regmodel)
```

Analysis of Variance Table

Response: DiffinPain

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Treatment	1	7.379	7.3792	6.9749	0.008713 **
Residuals	291	307.870	1.0580		

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

**Q:** What do you notice about your results from the pooled t-test, t-test for slope in linear regression, and from ANOVA F-test?

**A:**

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In this study we have levels  $a = 2$ ,  $H_0 : \mu_A = \mu_P$ ,  $H_a : \mu_a \neq \mu_P$ .

$$F = \frac{MSA}{MSE} = \frac{MS_{\text{Treatment}}}{MS_{\text{Residual}}} = 6.97 \sim F_{1,291}$$

So the p-value will be  $p = P(F_{1,291} > 6.97) = 0.008713 < 0.05$ . Therefore, we reject  $H_0$ , we have very strong evidence that the Auranofin and Placebo therapy differs. Since  $a = 2$ , we can have an easy follow up for two sample t-test with  $H_a : \mu_A > \mu_P$ . Once  $\bar{y}_A > \bar{y}_P$  then we can conclude that Auranofin is effective.

### 3 Lec 3 One-Way ANOVA with $a$ Levels

#### 3.1 Computational Formulae for SS

- $SST = \sum_{j=1}^a \sum_{i=1}^{n_j} y_{ij}^2 - N\bar{y}_T^2 = [Y] - [T]$
- $SSA = \sum_{j=1}^a n_j \bar{y}_j^2 - N\bar{y}_T^2 = [A] - [T]$
- $SSE = \sum_{j=1}^a \sum_{i=1}^{n_j} y_{ij}^2 - \sum_{j=1}^a n_j \bar{y}_j^2 = [Y] - [A]$

$[Y]$  denotes the individual observations ( $n$ ),  $[A]$  denotes the actual groups means,  $[T]$  denotes the grand means. The actual formula for these are:

- $[Y] = \sum_{j=1}^a \sum_{i=1}^{n_j} y_{ij}^2$
- $[A] = \sum_{j=1}^a n_j \bar{y}_j^2$
- $[T] = N\bar{y}_T^2$

#### 3.2 Proofs of Computational Formulae for SS

*Proof.*

$$\begin{aligned} SST &= \sum_{j=1}^a \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_T)^2 \\ &= \sum_i \sum_j y_{ij}^2 - 2\bar{y}_T \sum_i \sum_j y_{ij} + \sum_i \sum_j \bar{y}_T^2 \\ &= [Y] - 2\bar{y}_T N\bar{y}_T + N\bar{y}_T^2 \\ &= [Y] - N\bar{y}_T^2 \\ &= [Y] - [T] \end{aligned}$$

$$\begin{aligned}
SSA &= \sum_{j=1}^a n_j (\bar{y}_j - \bar{y}_T)^2 \\
&= \sum_j n_j \bar{y}_j^2 - 2 \sum_j \bar{y}_T n_j \bar{y}_j + \sum_j n_j \bar{y}_T^2 \\
&= [A] - 2\bar{y}_T \sum_j n_j \bar{y}_j + N\bar{y}_T^2 \\
&= [A] - N\bar{y}_T^2 \\
&= [A] - [T]
\end{aligned}$$

$$\begin{aligned}
SSE &= \sum_i \sum_j (y_{ij} - \bar{y}_j)^2 \\
&= \sum_i \sum_j y_{ij}^2 - \sum_i \sum_j 2\bar{y}_j y_{ij} + \sum_i \sum_j \bar{y}_j^2 \\
&= [Y] - 2 \sum_j \bar{y}_j \sum_i y_{ij} + \sum_j n_j \bar{y}_j^2 \\
&= [Y] - \sum_j n_j \bar{y}_j^2 \\
&= [Y] - [A]
\end{aligned}$$

$$\begin{aligned}
SST &= [Y] - [T] \\
&= [A] - [T] + [Y] - [A] \\
&= SSA + SSE
\end{aligned}$$

□

### Example 3.1. Example of Fertilizers in Farming

The image is a screenshot of a presentation slide. At the top, there is a status bar with the time '10:13 PM', date 'Sun 25 Jul', and battery level '100%'. Below this is a navigation bar with icons for back, forward, search, and other presentation controls. The main content area has a title 'Example: Fertilizers in Farming' in blue. The text describes an experiment with four fertilizers (A, B, C, D) and 16 plots. It asks the user to define the experiment, state hypotheses, and check if the design is balanced. The slide is numbered '11 / 16' in the bottom right corner.

**Example: Fertilizers in Farming**

There are four different fertilizers (brands A, B, C, D). We wish to determine if there a difference in the crop yield due to the fertilizer.

There are 16 plots of lands available to be tested. Each fertilizer is randomly assigned to four plots. The crop yield (in kg) of each plot is then measured after the fertilizer is used.

- ▶ Define the experiment: identify the experimental units, factor(s) and levels, response variable(s), etc.
- ▶ State the appropriate hypotheses to test the question of interest
- ▶ Is the design balanced or unbalanced?

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R output as follows:

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## R Code - Farming Example

```
> farmdata <- read.csv("farmingfertilizerdata.csv")
> farmdata
  Fertilizer CropYield
1          A         65
2          A         54
.
5          B         55
6          B         58
.
9          C         64
10         C         67
.
13         D         60
.
16         D         70
```

$a=4$

```
> attach(farmdata)

> tapply(CropYield, Fertilizer, (length) n)
A B C D
4 4 4 4

> tapply(CropYield, Fertilizer, (mean)  $\bar{y}$ )
      A      B      C      D 
58.75 60.00 68.75 65.50

> tapply(CropYield, Fertilizer, (sd) s)
      A      B      C      D 
4.856267 4.396969 4.272002 4.434712
```

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## Boxplots

$y \sim \text{Factor}$

```
> boxplot(CropYield ~ Fertilizer,
main="Farming Fertilizer Experiment",
xlab="Fertilizer",
ylab="Crop Yield (kg)")
```

highest

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## Example: Farming - ANOVA

Test the question of interest using ANOVA. Include all the steps.

```
# Regression Model with intercept
> farm.regmodel<- lm(CropYield ~ Fertilizer)

# ANOVA table
> anova(farm.regmodel)
```

*Linear model*      *response ~ Factors*

Analysis of Variance Table

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Fertilizer	3	264.5	88.167	4.3629	0.02694 *
Residuals	12	242.5	20.208		

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

*a=4*  
*N=16*

*p-value*

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The experimental unit is a plot. The factor is the brand of fertilizer which are A, B, C, D (4 levels/treatments). The response is the crop field (in kg). In the experiment,  $N = 16$  plots,  $n = 4$ ,  $n_A = n_B = n_C = n_D$  that means it's a balanced design. The hypothesis test is  $H_0 : \mu_A = \mu_B = \mu_C = \mu_D$  vs.  $H_a : \mu_i \neq \mu_j$  for at least one pair of  $i, j$  that  $i \neq j$ , and  $i, j = A, B, C, D$ .

$$F = \frac{MS_{\text{Fertilizer}}}{MSE} = 4.3629 \sim F_{3,12} \text{ under } H_0$$

$$p = P(F_{3,12} > 4.3629) = 0.0269$$

Since p value is less than 0.05, reject  $H_0$ . Strong evidence to conclude that the brands of fertilizer differ in crop yield.

Question for the next chapter:

Which fertilizer is the best? - We can examine the boxplot or look at all pairwise comparisons between fertilizers.

Is Fertilizer A better than B, etc.? Does using Fertilizer A result in double the crop yield of using B, etc.? - Test Contrasts.

### 3.3 What happens if we reject $H_0$ ?

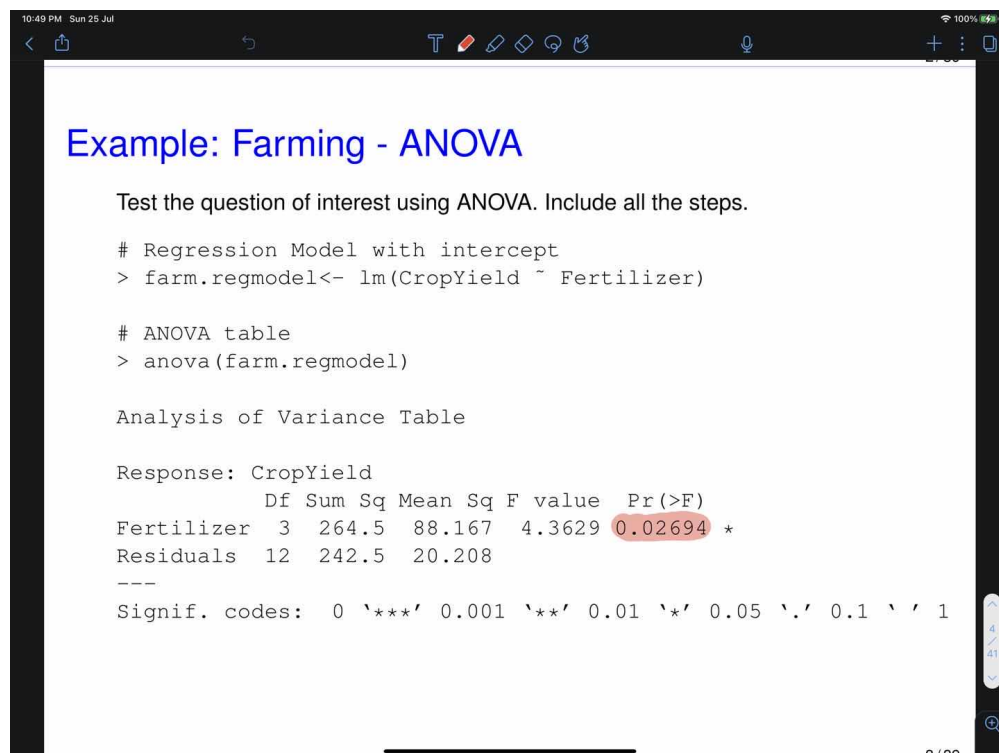
If we reject  $H_0$ , ANOVA tells us that there is statistically significant difference between group means. When there are more than 2 groups, it can be because:

- One group has a different mean than others
- Some groups have different means
- All groups have different means from each other

## 4 Lec 4 Follow-Up Comparisons for One-Way ANOVA: Testing Contrasts and Post-Hoc Analysis

### 4.1 Recall Example of Fertilizers in Farming- ANOVA

**Example 4.1.** The R output as follows:



The screenshot shows a presentation slide with a dark background. The title 'Example: Farming - ANOVA' is in blue. Below it, the text 'Test the question of interest using ANOVA. Include all the steps.' is followed by R code. The code defines a linear model and runs an ANOVA. The output shows the ANOVA table with a p-value of 0.02694, which is highlighted in red. A significance code legend is at the bottom.

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Example: Farming - ANOVA

Test the question of interest using ANOVA. Include all the steps.

# Regression Model with intercept
> farm.regmodel<- lm(CropYield ~ Fertilizer)

# ANOVA table
> anova(farm.regmodel)

Analysis of Variance Table

Response: CropYield
      Df Sum Sq Mean Sq F value Pr(>F)
Fertilizer  3  264.5   88.167   4.3629 0.02694 *
Residuals 12  242.5   20.208
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

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## 4.2 Pairwise Comparisons

**Definition 4.2.** Compare one group mean to another group mean (1 df since there are 2 groups). Conducted like a two-sample t-test but use pooled estimate of variance which is calculated using all groups:

$$s_p^2 = \frac{1}{N-a} \sum_{j=1}^a (n_j - 1) s_j^2 = MSE$$

where  $s_j^2$  is the sample variance in the  $j$ th group and  $\sum_{j=1}^a n_j = N$ . This  $s_p^2$  is a better estimate of the variance than just using observations from 2 groups, which means more power. The idea is to set up (linear) contrasts to make comparisons.

**Definition 4.3. Linear Contrasts** A linear contrast is a linear combination of the group means:  $\psi = \sum_{j=1}^a c_j \mu_j$ . Restrict sum of the coefficients in the contrast to be 0:  $\sum_{j=1}^a c_j = 0$ .

We use  $H_0 : \mu_j = \mu_k$  for  $j \neq k$ ,  $j, k = 1, 2, \dots, a$ . Results in coefficients such as  $c = \{1, -1, 0, 0\}$ , or  $c = \{0, 1, 0, -1\}$ , etc. Usually make the sign of  $\psi$  positive to reflect direction of the difference we predict. If group 1 is predicted to be higher than group 2, then  $c = \{1, -1, 0, 0\}$  rather than  $c = \{-1, 1, 0, 0\}$ . Test  $\psi = 0$  using  $\psi = \sum_{j=1}^a c_j \bar{Y}_j$ .

Advantages of using Contrasts:

- Compare more than 2 groups or more than one set of equalities.
- General procedure that cover many possible research questions.
- Simple to check if contrasts are orthogonal/linearly independent.

### Example 4.4. Farming- Pairwise Comparisons Using Contrasts

Set up a table with coefficients to compare these pairwise means:

- Is Fertilizer A different from B?-  $\psi_1$
- Is Fertilizer B different from C?-  $\psi_2$
- Is Fertilizer C different from D?-  $\psi_3$

Group	A	B	C	D
$\psi_1$	1	-1	0	0
$\psi_2$	0	1	-1	0
$\psi_3$	0	0	1	-1



**Table Including Means and Contrast Estimates**

Group	A	B	C	D	Contrast Estimate
Means	58.75	60	68.75	65.5	$\hat{\psi}$
$\psi_1$	1	-1	0	0	-1.25
$\psi_2$	0	1	-1	0	-8.75
$\psi_3$	0	0	1	-1	3.23

Example of the calculation:

$$\begin{aligned}
 \hat{\psi}_1 &= \sum_{j=1}^4 c_j \bar{y}_j = 1(\bar{y}_A) - 1(\bar{y}_B) + 0(\bar{y}_C) + 0(\bar{y}_D) \\
 &= \bar{y}_A - \bar{y}_B \\
 &= 58.75 - 60 = -1.25
 \end{aligned}$$

**Example 4.5. Farming- More Complex Comparisons**

Set up a table with coefficients to compare the following:

- Compare the two farms:  $\psi_4$
- Compare the two types of element-based fertilizers:  $\psi_5$

Farm	2	2	1	1
Element	P	P	N	P
$\psi_4$	1	1	-1	-1
$\psi_5$	1	1	-3	1

**Table Including All Comparisons**

Group	A	B	C	D	Contrast Estimate
Farm	2	2	1	1	$\hat{\psi}$
Element	P	P	N	P	
Means	58.75	60	68.75	65.5	
$\psi_1$	1	-1	0	0	-1.25
$\psi_2$	0	1	-1	0	-8.75
$\psi_3$	0	0	1	-1	3.23
$\psi_4$	1	1	-1	-1	-15.5
$\psi_5$	1	1	-3	1	-22

Note that those numbers are table of  $C_j$ 's.

### 4.3 Testing Using Contrasts

Under certain conditions,  $\hat{\psi} \sim N\left(\sum_{i=1}^a c_j \mu_j, \sigma^2 \sum_{i=1}^a \frac{c_j^2}{n_j}\right)$

#### Definition 4.6. Contrast Sum of Squares

The Contrast Sum of Squares is explained variation by the contrast:

$$SS_{\psi} = \frac{\hat{\psi}^2}{\sum_{j=1}^a (c_j^2/n_j)}$$

- Since 1 df,  $MS_{\psi} = SS_{\psi}$
- Compare to MSE to see if contrast explains more of the variation than noise:

$$F = \frac{MS_{\psi}}{MSE}$$

- Equivalent to a t-test:  $t = \frac{\hat{\psi}}{\sqrt{MSE \sum_{j=1}^a (c_j^2/n_j)}} \sim t_{N-a}$  under  $H_0$
- t Confidence Intervals for contrasts:

$$\hat{\psi} \pm t_{N-a;\alpha/2} \sqrt{MSE \sum_{j=1}^a (c_j^2/n_j)}$$

### Example 4.7. Farming R output:

```
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+ : [fullscreen]

Farming R Output - ANOVA

# Regression Model with intercept
> farm.regmodel<- lm(CropYield ~ Fertilizer)

# ANOVA table
> anova(farm.regmodel)

Analysis of Variance Table

Response: CropYield
      Df Sum Sq Mean Sq F value Pr(>F)
Fertilizer 3  264.5   88.167   4.3629 0.02694 *
Residuals 12  242.5   20.208
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

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```

```
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+ : [fullscreen]

R Code - Contrasts


$$Y_i = \beta_1 I_{A,i} + \beta_2 I_{B,i} + \beta_3 I_{C,i} + \beta_4 I_{D,i} + \epsilon$$

# Model without intercept
> model2 <- lm(CropYield ~ Fertilizer - 1)

# Coefficients of contrasts
> L <-matrix(c(
+ 1, -1, 0, 0,
+ 0, 1, -1, 0,
+ 0, 0, 1, -1,
+ 1, 1, -1, -1,
+ 1, 1, -3, 1), nrow=5, byrow=T)

# Required package
> require(multcomp)

No intercept
β1 = μA
⋮
β4 = μD

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```

```

R Output - Testing Contrasts

> summary(glht(model2, L), test=adjusted("none"))

Simultaneous Tests for General Linear Hypotheses

Fit: lm(formula = CropYield ~ Fertilizer - 1)

Linear Hypotheses:
Estimate Std. Error t value Pr(>|t|)
1 == 0    -1.250      3.179  -0.393  0.70104
2 == 0    -8.750      3.179  -2.753  0.01751 *
3 == 0     3.250      3.179   1.022  0.32675
4 == 0   -15.500      4.495  -3.448  0.00482 **
5 == 0   -22.000      7.786  -2.826  0.01530 *
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.'
(Adjusted p values reported -- none method)

```

State the conclusions based on the contrasts that were tested.

- Compare brands of fertilizers
- Compare farms
- Compare element bases
- Which fertilizer brand would you recommend? Why?

Conclusions:

- Moderate (to strong) evidence that Fertilizer C yields more than Fertilizer B
- Strong evidence that Farm 1 yields more than Farm 2
- Moderate (to strong) evidence that Nitrogen-based fertilizers are better than Phosphorous-based.

#### 4.4 Orthogonal Contrasts

**Definition 4.8.** This is useful for dividing sums of squares into different components. Any SS can be decomposed into as many independent sums as there are degrees of freedom. Two contrasts are

said to be **orthogonal** if: for  $\psi_1$  and  $\psi_2$  with coefficient vectors  $\vec{c}_1$  and  $\vec{c}_2$  respectively,

$$\sum_{j=1}^a \frac{c_{1j}c_{2j}}{n_j} = 0$$

Note that when sample sizes equal, then orthogonality simplifies to dot product of 0:

$$\vec{c}_1 \cdot \vec{c}_2 = \sum_{j=1}^a c_{1j}c_{2j} = 0$$

Orthogonality is a property of a pair of contrasts. If there are  $k$  contrasts, then there are  $\binom{k}{2}$  pairs.

**Definition 4.9.** If all contrasts in a set are orthogonal to one another, then it's called **mutually orthogonal**. There can be no more mutually orthogonal contrasts than the treatment df (i.e.  $a - 1$ ). Any set of  $a - 1$  mutually orthogonal contrasts can be used to express the SS for the treatment/factor (for balanced designs at least):

$$SSA = \sum_{k=1}^{a-1} SS_{\psi_k}$$

If orthogonal, each SS is independent and each contrast test can be interpreted additively (no problems with multi-collinearity).

#### **Example 4.10. Example: Farming - Orthogonal Contrasts**

How many contrasts are orthogonal amongst the 5 contrasts we tested?

- df for Fertilizer is 3. We are limited to 3 orthogonal contrasts
- Can we find a set of 3? Which contrasts?

In this example, we can easily find that  $a = 4$ , df = 3 (calculated from  $a - 1$ )

Table to check Orthogonality					
Group	A	B	C	D	
$\psi_1$	1	-1	0	0	
$\psi_2$	0	1	-1	0	
$\psi_3$	0	0	1	-1	
$\psi_4$	1	1	-1	-1	
$\psi_5$	1	1	-3	1	Sum
$\psi_1$ vs $\psi_2$	0	-1	0	0	-1
$\psi_1$ vs $\psi_3$	0	0	0	0	0
$\psi_1$ vs $\psi_4$	1	-1	0	0	0
$\psi_1$ vs $\psi_5$	1	-1	0	0	0
$\psi_2$ vs $\psi_3$	0	0	-1	0	-1
$\psi_2$ vs $\psi_4$	0	1	1	1	2
$\psi_2$ vs $\psi_5$	0	1	3	0	4
$\psi_3$ vs $\psi_4$	0	0	-1	1	0
$\psi_3$ vs $\psi_5$	0	0	-3	-1	-4
$\psi_4$ vs $\psi_5$	1	1	3	-1	4

We have  $\binom{5}{2} = 10$  pairs of contrast. In this example, we have  $a = 4$ , so there are  $a - 1 = 3$  mutually orthogonal contrasts. We will choose  $\psi_1, \psi_3, \psi_4$  to be mutually orthogonal.

#### 4.5 Method for Constructing Mutually Orthogonal Set

We have  $a$  groups, we will compare 1st group to average of all other groups. For all other contrasts, make 1st group's coefficient 0. Compare 2nd group to average of the latter groups. Continue this pattern. Make 2nd group's coefficient 0. Compare 3rd group to average of the latter groups, and so on...

##### Constructing Helmert Contrasts

Start with  $c_1 = 1$  and  $c_j = -\frac{1}{a-1}$  for  $j > 1$ . For  $j = 2, \dots, a - 1$ , let  $c_k = 0$  when  $k \leq j - 1$  and  $c_j = 1$  and  $c_k = -\frac{1}{a-j}$  when  $k \geq j + 1$ . These contrasts are known as **Helmert Contrasts**.

##### Example 4.11. Farming: Helmert Contrasts

Construct the Helmert Contrasts. Verify that their contrasts are mutually orthogonal.

In this example, we have group  $a = 4$ , it implies to there are 3 contrasts are mutually orthogonal. Note that the following vector has the order of  $(A, B, C, D)$ :

$$\psi_1 : \vec{c}_1 = (1, -1/3, -1/3, -1/3) \implies A \text{ to average of others}$$

$$\psi_2 : \vec{c}_2 = (0, 1, -1/2, -1/2) \implies B \text{ to average of latter groups}$$

$$\psi_3 : \vec{c}_3 = (0, 0, 1, -1) \implies C \text{ to average of latter groups}$$

$$\vec{c}_1 \cdot \vec{c}_2 = \vec{c}_1 \cdot \vec{c}_3 = \vec{c}_2 \cdot \vec{c}_3 = 0 \implies \psi_1, \psi_2, \psi_3 \text{ are mutually orthogonal.}$$

## 4.6 Balanced Designs

ANOVA is the best for balanced designs, there are potential problems with unbalanced designs. MSE in unbalanced designs:

- MSE is a pooled variance, weighted by size of each group. Larger groups dominate MSE towards their variance.
- MSE is still unbiased for  $\sigma^2$ .

In balanced designs:

- MSE gets an equal contribution from each group.
- SSA can be decomposed into the sum of the contrast SS.
- Usually more powerful.

ANOVA can still be used for unbalanced designs. The more unequal the group sizes, the poorer the performance. So, we need to attempt to design balanced experiment. Sometimes necessary to use unbalanced designs (e.g. Larger control group).

## 4.7 Problem with Multiple Comparisons

Inflation of Type I Error rate: Chance of making a Type I Error is very high when conducting many simultaneous tests.

We are often interested in a family/group of different hypothesis tests or confidence intervals

- All pairwise differences in group means, there are  $\binom{a}{2}$  pairwise comparisons.
- Meaningful contrasts

The **familywise error rate**,  $\alpha_{FW}$  is the probability of making at least one Type I Error in the set of tests. Suppose we conduct  $k$  independent tests, each at significance level  $\alpha$ , then:

$$\begin{aligned}\alpha_{FW} &= P(\text{reject at least one } H_0 | \text{all } H_0 \text{ are true}) \\ &= 1 - (1 - \alpha)^k\end{aligned}$$

#### Example 4.12. Familywise Type I Error Rate

If we conduct  $k = 5$  tests (e.g., 5 contrasts such as in the Farming example) with  $\alpha = 0.05$ , then

$$\alpha_{FW} = P(\text{at least one Type I Error}) = 1 - (1 - 0.05)^5 = 0.2262$$

The probability of committing a Type I Error somewhere in these tests is now 22.62%. Inflated Type I Error: will increase even more as we increase number of tests. If tests are dependent,  $\alpha_{FW}$  will be lower, but not lower than the Type I Error rate for any individual test. We need to control for this.

#### 4.8 Pre-planned Tests/Primary Research Questions

Most often, an experiment is set up to answer one primary research question. Could also have follow-up questions.

- If research question/tests are pre-planned/ad-hoc (before looking at the results of the analysis), it can be tested without correcting Type I Error
- Why before and not after? *Fishing for significant results!*
- If question is important, do not conduct it in a family of less powerful tests
- Need to correct if we test secondary research questions or make comparisons after seeing results of analysis (post-hoc)

#### 4.9 Simple Solution to Correct Type I Error

- Decrease the family wise Type I Error rate ( $\alpha_{FW}$ ) by decreasing the Type I Error rate for each individual test ( $\alpha$ )
- Adjust Type I Error for the worst case (independent tests)
- This adjustment results in less power/higher chance of making Type II Error

Are there other solutions? Yes:

1. Bonferroni
2. Sidak-Bonferroni
3. Tukey's HSD
4. Scheffe



## 4.10 Bonferroni Method

**Definition 4.13.** Bonferroni inequality:  $P(A \cup B) \leq P(A) + P(B)$ .

Let  $A_i$  be the event of making a Type I Error on  $i$ th test, then:  $\alpha_{FW} = P(\bigcup_{i=1}^k A_i) \leq \sum_{i=1}^k P(A_i)$ . Note that  $\alpha_i = P(A_i)$  for  $i = 1, \dots, k$ .

### Bonferroni Method

For each of the  $k$  tests, use a significance level of  $\alpha_{FW}/k$ . Then CI coverage rate or Type I Error rate is at most  $100(1 - \alpha_{FW})\%$  or  $\alpha_{FW}$ .

- Conservative procedure
- Since the chance of making at least one Type I Error can be much lower than  $\alpha$ , it is likely that the  $k$  tests may result in Type II Error (less powerful)
- Type I Errors are not always more serious than Type II (depends on the context)

**Example 4.14.**

**R Code: Bonferroni - Pairwise Comparisons**

```
p.adj = "bonf"

With this command p-values in 'R' are multiplied by k. Reject H0 if adjusted p-value is < αFW.
```

#Pairwise comparisons without adjustment

```
> pairwise.t.test(CropYield, Fertilizer, p.adj="none")
```

Pairwise comparisons using t tests with pooled SD

data: CropYield and Fertilizer

A	B	C
B	0.7010	-
C	0.0084	0.0175
D	0.0552	0.1092

P value adjustment method: none

# Bonferroni - Pairwise comparisons

```
> pairwise.t.test(CropYield, Fertilizer, p.adj="bonf")
```

Pairwise comparisons using t tests with pooled SD

data: CropYield and Fertilizer

A	B	C
B	1.000	-
C	0.051	0.105
D	0.331	0.655

P value adjustment method: bonferroni

$\alpha_{FW} = 0.05$

**Handwritten Notes:**

- $a = 4$
- $\binom{4}{2} = 6$  pairwise comparisons
- $\alpha_i = \frac{\alpha_{FW}}{k}$  for  $i = 1, \dots, k$
- Reject H<sub>0</sub> if  $p < \frac{0.05}{6}$  (adjusted p-value)
- Reject H<sub>0</sub> if  $p < \frac{\alpha_{FW}}{k}$  (adjusted)
- Reject H<sub>0</sub> if  $k p < \alpha_{FW}$  (R does this)
- Reject H<sub>0</sub> if  $p < 0.05$  (unadjusted)

**R Code: Bonferroni - Contrasts**

```

# Model without intercept
> model <- lm(CropYield ~ Fertilizer - 1)

# Coefficients of contrasts
> L <- matrix(c(
+ 1, -1, 0, 0,
+ 0, 1, -1, 0,
+ 0, 0, 1, -1,
+ 1, 1, -1, -1,
+ 1, 1, -3, 1), nrow=5, byrow=T)

# Required package
> require(multcomp)

> summary(glht(model, L), test=adjusted("bonf"))
Simultaneous Tests for General Linear Hypotheses

Fit: lm(formula = CropYield ~ Fertilizer - 1)

Linear Hypotheses:
      Estimate Std. Error t value Pr(>|t|)
1 == 0   -1.250      3.179  -0.393  1.0000
2 == 0   -8.750      3.179  -2.753  0.0876 .
3 == 0    3.250      3.179   1.022  1.0000
4 == 0  -15.500      4.495  -3.448  0.0241 *
5 == 0  -22.000      7.786  -2.826  0.0765 .
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Adjusted p values reported -- bonferroni method)

```

Handwritten notes:

- $\psi = \sum_{j=1}^a c_j \mu_j$
- $k = 5$  tests / contrasts

#### 4.11 Sidak-Bonferroni Procedure

Bonferroni procedure is simple to use (by hand). We can improve procedure if using computer for calculations by directly working with:  $\alpha_{FW} = 1 - (1 - \alpha)^k$

##### Definition 4.15. Sidak Bonferroni Method

Adjust the individual Type I Error as follows:

$$\alpha = 1 - (1 - \alpha_{FW})^{1/k}$$

For example,  $k = 5$  and  $\alpha_{FW} = 0.05$  compare Bonferroni and Sidak-Bonferroni:

- $\alpha_B = \frac{0.05}{5} = 0.01$
- $\alpha = 1 - (1 - 0.05)^{1/5} = 0.0102$  This is slight higher due to lower individual Type I Error rate.

Still have same family-wise rate, but more power for individual tests.

## 4.12 Confidence Intervals

For tests, we adjust the p-values or individual significance levels. For CIs, adjust the critical value. Suppose we want to construct  $k$  confidence intervals with familywise coverage of  $100(1 - \alpha_{FW})\%$ , then use

$$\hat{\theta} \pm t_{df, \frac{\alpha_{FW}}{2k}} SE(\hat{\theta})$$

## 4.13 Tukey's HSD Procedure

- Based on “Studentized Range Distribution” which is based on  $\max_{j,k \in \{1,2,\dots,a\}} \{\bar{y}_j - \bar{y}_k\}$
- Gives a simultaneous Type I Error rate of  $\alpha_{FW}$  or confidence level of  $100(1 - \alpha_{FW})\%$
- Conservative if unbalanced design
- Less conservative than Bonferroni method, especially if group sample sizes are equal (approx.)

**Definition 4.16. Tukey's Honestly Significant Difference (for differences between pairs of groups)**

Confidence interval for  $\mu_k - \mu_j$  is:

$$(\bar{y}_j - \bar{y}_k) \pm \frac{q_{\alpha_{FW}; a, N-a}}{\sqrt{2}} \sqrt{MSE(\frac{1}{n_j} + \frac{1}{n_k})}$$

where  $q_{\alpha; a, N-a}$  is the appropriate critical value of the Studentized Range Distribution.

To avoid redundancy, compare the largest and smallest means and if we failed to reject  $H_0$ , then stop. If  $H_0$  is rejected, compare next largest to next smallest means and continue on...

Code:

```
> TukeyHSD(aov(model_name), factor = "factor_name")
```

‘R’ Gives CIs for all pairwise means. Reject  $H_0$  for the given pair if the adjusted p-value  $< \alpha$  or CI does not contain 0.