# ANOVA

## Young W. Lim

### 2019-09-06 Fri

Young W. Lim

ANOVA

2019-09-06 Fri 1 / 26

2

・ロト ・ 四ト ・ ヨト ・ ヨト





- One-way ANOVA
- one-way ANOVA Model
- Two-way ANOVA
- Within Groups Variance Estimate  $S_W^2$

#### "Understanding Statistics in the Behavioral Sciences" R. R. Pagano

I, the copyright holder of this work, hereby publish it under the following licenses: GNU head Permission is granted to copy, distribute and/or modify this document under the terms of the GNU Free Documentation License, Version 1.2 or any later version published by the Free Software Foundation; with no Invariant Sections, no Front-Cover Texts, and no Back-Cover Texts. A copy of the license is included in the section entitled GNU Free Documentation License.

CC BY SA This file is licensed under the Creative Commons Attribution ShareAlike 3.0 Unported License. In short: you are free to share and make derivative works of the file under the conditions that you appropriately attribute it, and that you distribute it only under a license compatible with this one.

(日)

Analysis of variance (ANOVA) is

 a collection of <u>statistical models</u> and
 their associated <u>estimation procedures</u>
 (such as the "variation" among and between groups)
 used to analyze the differences
 among group means in a sample.

• The ANOVA is based on the law of total variance, where the <u>observed variance</u> in a particular variable is <u>partitioned</u> into components attributable to different sources of variation.  In its simplest form, ANOVA provides a statistical test of whether two or more population means are <u>equal</u>, and therefore generalizes the t-test beyond two means.  The analysis of variance can be used as an exploratory tool to explain observations. A dog show provides an example.

- A dog show is <u>not</u> a <u>random</u> sampling of the breed: it is typically limited to dogs that are adult, pure-bred, and exemplary.
- A <u>histogram</u> of dog <u>weights</u> from a show might plausibly be rather complex, like the yellow-orange distribution shown in the illustrations.
- Suppose we wanted to <u>predict</u> the <u>weight</u> of a dog based on a certain set of characteristics of eachdog.

 one-way analysis of variance (one-way ANOVA) is a technique that can be used to <u>compare means</u> of two or more samples (using the F distribution).  can be used only for numerical response data, the "Y", usually one variable, and numerical or (usually) categorical input data, the "X", always one variable, hence "one-way"

- The ANOVA tests the <u>null hypothesis</u> that samples in all groups are drawn from populations with the same mean values.
- To do this, two estimates are made of the population variance.
- These estimates rely on various assumptions

 The ANOVA produces an F-statistic, the <u>ratio</u> of the <u>variance</u> calculated <u>among the means</u> to the <u>variance</u> within the samples.

э

< 4<sup>3</sup> ► <

- If the group means are drawn from populations with the same mean values, the variance between the group means should be lower than the variance of the samples, following the central limit theorem.
- A <u>higher ratio</u> therefore implies that the samples were drawn from populations with different mean values.

## • The normal linear model describes treatment groups with probability distributions which are identically bell-shaped (normal) curves with different means.

- Thus fitting the models requires only the <u>means</u> of each treatment <u>group</u> and a <u>variance</u> calculation (an <u>average variance</u> within the treatment <u>groups</u> is used).
- Calculations of the <u>means</u> and the <u>variance</u> are performed as part of the hypothesis test.

# One-way ANOVA Model (3)

- The commonly used normal linear models for a completely randomized experiment are:
- i = 1, ..., I is an index over experimental units
- $j = 1, \ldots, J$  is an index over treatment groups
- $I_j$  is the <u>number</u> of experimental units in the <u>j-th</u> treatment group
- $I = \sum_{j} I_{j}$  is the total number of experimental units

- *y<sub>i,j</sub>* are observations
- $\mu_j$  is the <u>mean</u> of the observations for the jth treatment group
- $\mu$  is the grand mean of the observations
- $\tau_j$  is the jth treatment effect, a deviation from the grand mean
- $\sum \tau_j = 0$
- $\mu_j = \mu + \tau_j$
- $\varepsilon \sim N(0, \sigma^2)$ ,
- $\varepsilon_{i,j}$  are <u>normally</u> distributed <u>zero-mean</u> random errors.

- the two-way analysis of variance (ANOVA) is an extension of the one-way ANOVA that examines the influence of <u>two different categorical independent variables</u> on one continuous dependent variable.
- The two-way ANOVA not only aims at assessing the main <u>effect</u> of each <u>independent variable</u> but also if there is any interaction between them.

Suppose a data set

for which a dependent variable may be influenced by two factors which are potential sources of variation.

• The first factor has I levels  $i \in \{1, ..., I\}$ ) and the second has J levels  $j \in \{1, ..., J\}$ )

- Each combination (*i*, *j*) defines a treatment, for a total of *I* × *J* treatments.
- We represent the number of replicates for treatment (i, j) by  $n_{ij}$ ,
- and let k be the index of the replicate in this treatment  $(k \in \{1, ..., n_{ij}\})$ .

• From these data, we can build a contingency table, where

• 
$$n_{i+} = \sum_{j=1}^J n_{ij}$$

• 
$$n_{+j} = \sum_{i=1}^{l} n_{ij}$$

• 
$$n = \sum_{i,j} n_{ij} = \sum_i n_{i+1} = \sum_j n_{+j}$$

the total number of replicates

3

< ロ > < 同 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < 回 > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ >

- The experimental design is balanced if each treatment has the same number of replicates, *K*.
- the design is also said to be orthogonal allowing to fully distinguish the effects of both factors.

• We hence can write 
$$\forall i, j \ n_{ij} = K$$
, and  
 $\forall i, j \ n_{ij} = \frac{n_{i+} \cdot n_{+j}}{n}$ 

- In the classical approach, testing null hypotheses (that the factors have no effect) is achieved via their significance which requires calculating sums of squares.
- Testing if the interaction term is significant can be difficult because of the potentially large number of degrees of freedom.

- Following Gelman and Hill, the assumptions of the ANOVA, and more generally the general linear model, are, in decreasing order of importance:
  - the data points are relevant with respect to the scientific question under investigation;
  - the mean of the response variable is influenced <u>additively</u> (if not interaction term) and linearly by the factors;
  - It the errors are independent;
  - the errors have the same variance;
  - **o** the errors are normally distributed.

 weighted estimate of H<sub>0</sub> population variance, σ<sup>2</sup> weighted average of s<sub>1</sub><sup>2</sup> and s<sub>2</sub><sup>2</sup>

$$s_W^2 = \frac{SS_1 + SS_2}{(n_1 - 1) + (n_2 - 1)} = \frac{SS_1 + SS_2}{N - 2}$$

 weighted estimate of H<sub>0</sub> population variance, σ<sup>2</sup> weighted average of \$s\_1<sup>2</sup>, s\_2<sup>2</sup>, ...\$ and s<sup>2</sup><sub>k</sub>

$$s_W^2 = \frac{SS_1 + SS_2 + \dots + SS_k}{(n_1 - 1) + (n_2 - 1) + \dots + (n_k - 1)}$$
$$= \frac{SS_1 + SS_2 + \dots + SS_k}{N - k}$$