# Analysis of Variance (ANOVA)

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#### Introduction

- The "halo effect" is a hypothesized cognitive bias where a positive impression of one aspect of a person/brand leads to other aspects of that same person/brand being viewed more favorably than they should
- Today we'll look at data from the article: "Beauty is Talent: Task Evaluation as a Function of the Performer's Physical Attraction" published in *The Journal of Personality and Social Psychology* in 1974
  - 60 undergraduate males scored (from 0 to 25) an essay supposedly written by a female undergraduate
  - Attached to each essay was a photo of the supposed author that was randomly assigned from one of the following conditions: "attractive", "unattractive", or "none"

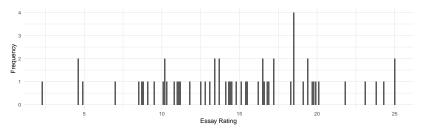


There are two types of hypotheses we might consider for this experiment:

- 1. **global hypothesis** Is an essay's rating associated with the type of photo attached to it?
- pairwise" hypotheses Do ratings in a particular condition (ie: "attractive") differ from another condition (ie: "none")?
  - There are 3 different pairwise hypotheses in this example
- The pairwise hypotheses can be evaluated using t-tests. However, type I errors are a concern
- Analysis of Variance (ANOVA) allows us to evaluate the global hypothesis with a single test



If the experimental condition and essay rating are *independent*, we'd expect the ratings in each condition to follow the same distribution. Below is the overall distribution of scores:





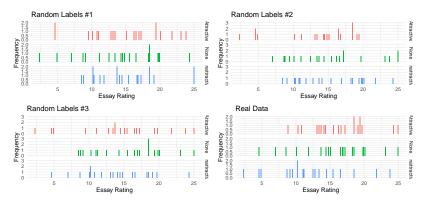
Under ANOVA's null hypothesis of *no association*, we'd expect ratings in group to be sampled from the overall distribution. Below we simulate this by randomly giving each data-point a group label (unrelated to its actual group):





## Hypothesis Testing

ANOVA aims to determine whether the observed distribution *within groups* significantly deviates from what we'd expect if group labels were random (ie: no association between "group" and "rating"):





### Predictions

 To compare the observed and expected distributions, ANOVA looks at *predicted values*

•  $\hat{y}_i$  is a prediction of  $y_i$ , the  $i^{th}$  data-point's outcome (rating)

Predictions can be made assuming H<sub>0</sub>, or using relationships in the observed data

Condition (group)	Mean Rating	Standard Deviation	n
Attractive	16.4	4.3	20
None	15.6	5.2	20
Unattractive	12.1	5.4	20
Overall	14.7	5.3	60

**Questions**: For a data-point in the "unattractive photo" group, what is its predicted rating under  $H_0$ ? What would you predict it's rating if you used the relationships present in the real data?



In ANOVA the null hypothesis reflects a *statistical model*:

 $y_i = \mu + \epsilon_i$ 

- y<sub>i</sub> is the i<sup>th</sup> data-point's observed outcome, μ is an overall mean, and ε is an error that is assumed to follow a N(0, σ) distribution
  - Because the errors have an expected value of zero, this model predicts ŷ<sub>i</sub> = µ, the overall mean, for every data-point (all i ∈ {1,2,...,n})
- This is known as the null model



ANOVA also considers an **alternative model**:

 $y_i = \mu_i + \epsilon_i$ 

- This model looks similar, but μ<sub>i</sub> is a group-specific mean that depends upon which group the i<sup>th</sup> data-point belongs to
  - Thus, this model predicts a group-specific mean for data-point, thereby reflecting an association between "group" and "rating"



Like when we first learned about regression, it is important to note the distinction between the *theoretical* and *fitted* models in ANOVA:

	Null Model	Alternative Model	
Theoretical	$y_i = \mu + \epsilon_i$	$y_i = \mu_i + \epsilon_i$	
Fitted	$\hat{y}_i = \overline{y}$	$\hat{y}_i = \overline{y}_i$	

*Note*:  $\overline{x}$  is calculated using the entire sample, but  $\overline{x}_i$  is the group-specific mean for the *i*<sup>th</sup> data-point's group

**Questions**: How could you *summarize* how well each model fits the observed data?



### Summarizing a Model

Under the any model each subject deviates from their prediction by a **residual**:

 $\begin{aligned} r_i &= y_i - \hat{y}_i \text{ (Definition of a residual)} \\ &= y_i - \overline{y} \text{ (Residuals for the null model)} \end{aligned}$ 

We can *summarize* the total variability of the null model's predictions using a **sum of squares**:

$$SST = \sum_{i} r_i^2$$
 for the null model

We call this *SST* (sum of squares total) because it is the *largest possible* sum of squares (of any justifiable model)



The alternative model can also be summarized using a **sum of squares**:

 $SSE = \sum_{i} r_i^2$  for the alternative model where  $r_i = y_i - \overline{y}_i$  (Residuals for the alt model)

We call this *SSE* because it summarizes the unexplained variability (errors) of the model that uses "group"



### Creating a Test Statistic

 In ANOVA we ask: "does the grouping variable improve model fit beyond what might be expected due to random chance?"
This can be assessed using the F-test statistic:

$$F = \frac{(SST - SSE)/(d_1 - d_0)}{\text{Std. Error}}$$



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► d<sub>1</sub> and d<sub>0</sub> denote the number of parameters estimated in model, in our example d<sub>0</sub> = 1 (the single overall mean) and d<sub>1</sub> = 3 (each group's mean)



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- The F statistic is the standardized drop in the sum of squares per additional parameter used by the alternative model



- We started off by considering what the data might look like if we randomly assigned group labels (slide 5/6)
  - If we did this many times, we could get an idea of how the F-statistic is distributed under the null hypothesis
  - This StatKey menu allows us to perform such a simulation

Here is a link to the data:

https://remiller1450.github.io/data/halo\_effect.csv



- Under the null hypothesis (ie: presuming the null model is true), this F-statistic follows an F-distribution that depends upon two different degrees of freedom (df) parameters
  - The numerator df is  $d_1 d_0$
  - The denominator df is  $n d_1$
- ▶ We can use StatKey to view various *F*-distribution curves
  - The observed F-statistic is compared to the F-distribution to determine the p-value



#### What is the Standard Error?

- We've seen that standard errors tend to look like a measure of variability divided by the sample size
- In the ANOVA setting:

Std. Error = 
$$\frac{SSE}{n-d_1}$$

- ► This is the sum of squares of the alternative model divided by its *degrees of freedom*,  $df = n d_1$
- Using this standard error, the F statistic can be expressed:

$$F = \frac{(SST - SSE)/(d_1 - d_0)}{SSE/(n - d_1)}$$



## Simplifying the *F*-statistic

Because this test statistic looks complex, statisticians define the "sum of square groups" as: SSG = SST - SSE, making the *F*-statistic:

$$F = \frac{SSG/(d_1 - d_0)}{SSE/(n - d_1)}$$

This is further simplified by denoting a sum of squares defined by its degrees of freedom as a **mean square**:

$$F = \frac{MSG}{MSE}$$

Here MSG is the mean square of "groups", MSE is the mean square of "error"



Calculating the *F*-statistic and the corresponding *p*-value are too tedious to perform by hand, but you are expected to be familiar with **ANOVA tables**, which are how software like R will report the results of an ANOVA test:

Source	df	Sum Sq.	Mean Sq.	F-statistic	<i>p</i> -value
"Group"	$d_1 - d_0$	SSG	MSG	MSG/MSE	Use $F_{d_1-d_0,n-d_1}$
Residuals	$n-d_1$	SSE	MSE		
Total	$n-d_0$	SST			

You might be asked to fill in the missing components of such a table, interpret a printed table, or relate an ANOVA table to visualizations or models



# What Should You Know? (cont.)

In addition to understanding the components of ANOVA table output, you should have a high-level understanding of the *F*-test:

- The ANOVA F-test involves a nominal categorical explanatory variable and a quantitative response variable
- The null model states that a single, overall mean is sufficient (indicative of independence), while the alternative model uses group-specific means (indicative of an association)
  - The F-statistic compares the performance of these two models on the sample data using sums of squares
- Under the null hypothesis of independence, the *F*-statistic should follow an *F*-distribution, which is used to determine the *p*-value

A small p-value provides evidence of an association

