

Analysis of Variance

Comparing two means:

- $H_0: \mu_1 = \mu_2$; $H_1: \mu_1 \neq \mu_2$
- Compare difference between sample means vs. a pooled estimate of standard deviation.
- Use Z test or t test depending on whether σ is known or sample size > 25 .

Comparing more than two means:

- Repetitive t or Z tests between pairs of means
- Cumbersome and gives "overtesting".
- At the 5% level, 1/20 comparisons would be "significant" just by chance. Must adjust for this.

ANOVA

Compares several means at once

- $H_0: \mu_1 = \mu_2 = \mu_3 = \mu_4$; H_1 : one or more is/are different
- Compare reduction in estimate of variance for separating into more than one distribution with different means.
Compare vs. a pooled estimate of variance remaining.
- Use F test with degrees of freedom:
- d.f. total = $n - 1$
- d.f. between = number of groups (k) - 1
- d.f. within = remaining d.f. = $n - k$
- compare to F tables for d.f.'s, significance levels.

Assumptions:

- Observations are independent
- variance is homogeneous across groups.

Procedure:

- Calculate total sum of squares, $SSt = \text{Sum}(X_i^2) - (\text{Sum } X)^2/N$
- Calculate between-group sum of squares, SSb
- Derive within-group SSw by $SSt - SSb$
- Divide by appropriate d.f. to get mean squares, MSw, MSb
- $F = MSb/MSw$
- Compare to critical F values table for dfb, dfw , significance
- Reject H_0 if F is big.
- Compare differences between individual pairs of means using Tukey's HSD test as a cutoff
- Tukey's HSD is "Honestly Significant Difference"--an adjustment of the "critical t" to allow for sloppiness when doing multiple comparisons.

$$HSD = q(\alpha, k, N - k) \sqrt{\frac{MSw}{n}}$$

- q from table in appendix C
- α is significance level, 1% or %5
- k is number of groups/treatments
- N is TOTAL observations
- n is number in each group
- MSw is within (error) mean square

- The Tukey test will tell you whether your conclusion about the differences will fly.

Mechanics--just like the computational formula for variance:

- Separate observations into groups by treatment.
- Calculate totals by group and overall (grand sum)
- Calculate means by treatment. Square each mean, multiply the square * the number of observations in each mean and add all these together.
- Square the grand sum and divide by the grand number = "Correction term"
- Square every observation and sum the squares (grand sum of squares)

Calculate corrected sums of squares:

- Total sum of squares, $SSt = \text{grandsumofsquares} - CT$
- Treatment Sum of Squares,
 $SSb = \text{Sumofmeans} (*n \text{ in each}) \text{ squared} - CT$
- Within (Residual) $SSw = \text{difference, } SSt - SSb.$
- Divide by appropriate df's to get Mean Squares
- Test ratio of Treatment mean square to within mean square.
- compare to critical F (α , dfb, dfw) from appendix B tables.

Multidimensional comparisons:

- Symmetrical Square designs.
- Same thing, except you regroup the observations by another treatment as well, and go through the separation out of another treatment sum of squares.
- These other dimensions "explain" more of the error variance and reduce the residual "unexplained" variation that's attributed to Within group random error.
- Degrees of freedom are also partitioned into the new explained dimension ($df = \text{number of groups} - 1$) reducing the degrees of freedom associated with "error" variance.
- Since this further partition of variance reduces the variation attributed to random effects, it makes it easier to show differences that are attributable to causes.
- "Within group" = "Error" = "Random" = "Residual" = "Unexplained".
- "Random error" isn't without cause. It only means we can't explain (don't know) the cause.