

Chapter 16 (continued): Analysis of Variance

Note: In a single-factor study, the levels are also known as treatments.

Partitioning the Total Sum of Squares

SSTO =

SSTR =

SSTR measures the variability

SSE =

SSE measures the variability

- **SSE estimates the “natural” variation in the data (variation not due to the different treatments).**
- **If the treatment means differ greatly, then SSTR will be**

Note that $SSTO = SSTR + SSE$.

Proof:

- The associated degrees of freedom are also additive, and are used to obtain the mean squares.

Expected Values of the Mean Squares (see pg. 696-698 for details):

Note: • MSE is an unbiased estimator of the error variance σ^2 .

• If all treatments have the same population mean, then

$E(\text{MSTR}) =$

• If the treatments have different population means, then MSTR should be _____ than MSE, on average.

F-test for Equality of Treatment Population Means

• A natural test statistic to use is:

• If $F^* \gg 1 \rightarrow$ evidence supports

If F^* near 1 \rightarrow evidence supports

• Under H_0 , F^* has a

• We reject H_0 if

Theoretical Justification of F-test
(Proof of distribution of F^* under H_0)

- We need to use Cochran's Theorem:

Suppose our observations $Y_1, \dots, Y_n \sim N(\mu, \sigma^2)$. Then if we break SSTO into k sums of squares SS_1, \dots, SS_k (having degrees of freedom (df_1, \dots, df_k) , then for $j = 1, \dots, k$,

random variables, provided that

ANOVA situation:

- We have n_T observations. Under H_0 , they are

ANOVA Table

Kenton Food Example (from SAS):

- **Do the four package designs have the same population mean sales?**

We test:

Factor Effects Model

- This is an alternative formulation of the ANOVA model.
- If μ_{\cdot} is a type of overall population mean response, then let
- τ_i is the effect of the i -th factor level, or the i -th treatment effect.
- μ_{\cdot} is often the simple average of $\mu_1, \mu_2, \dots, \mu_r$, but it could be defined as a weighted average of $\mu_1, \mu_2, \dots, \mu_r$.
- Note our model equation is:
- Our hypothesis of $H_0: \mu_1 = \mu_2 = \dots = \mu_r$ is equivalent to
- If μ_{\cdot} is an “overall mean response” then τ_1, \dots, τ_r measure how much the treatment means deviate from the overall mean, and
- We can use a regression approach to estimate the parameters of this model.

Regression Approach to the ANOVA Model

- Since $\tau_r = -\tau_1 - \tau_2 - \dots - \tau_{r-1}$, we need not estimate τ_r . We only estimate μ_{\cdot} , τ_1 , τ_2 , \dots , τ_{r-1} .
- Recall example when $r = 3$ and $n_1 = 1$, $n_2 = 3$, $n_3 = 2$. Then let:

- Then the factor-effects model can be stated as

For example:

- If we use the indicator variables

then this will produce the X matrix above, and we can fit the factor-effects ANOVA model via regression.

SAS Example (Kenton Foods, Table 16.1):

- **To fit the cell-means model via regression, we should let**

- **Thus we can define indicator variables**

which will produce the above X matrix.

- **We can fit the cell-means model via regression (specifying NO intercept term in this case!).**

Kenton Foods Example: