## **STAT** 500

Inference about Means for Several Populations

#### **Scenario**

- Observational Studies
  - More than Two Populations
- Experiments
  - One Factor with more than two levels
- Compare observations of variable (quantitative) for multiple treatment groups or populations.

#### **Notation**

- Population or Treatment Group Parameters
  - Number:  $i=1,2,\ldots,r$
  - Population or Treatment means  $=\mu_1,\mu_2,\ldots,\mu_r$
  - Population or Treatment Variances  $=\sigma_1^2,\sigma_2^2,\ldots,\sigma_r^2$
  - Population or Treatment Std. Dev.  $= \sigma_1, \sigma_2, \ldots, \sigma_r$

#### **Notation**

- Data and Summary Statistics
  - $n_i=$  sample size for ith sample or treatment group
  - $-\ Y_{ij}=j$ th observation in the ith sample or treatment group, where  $j=1,2,\ldots,n_i$
  - Mean for ith sample or treatment group  $ar{Y}_i = ar{Y}_i. = rac{1}{n_i} \sum\limits_{i=1}^{n_i} Y_{ij}$
  - Variance for *i*th sample or treatment group  $1 \quad n_i \quad \quad 2$

$$S_i^2 = rac{1}{n_i-1}\sum\limits_{j=1}^{n_i}(Y_{ij}-ar{Y}_{i\cdot})^2$$

#### **Notation**

- Summary Statistics
  - Total number of observations:  $N = \sum\limits_{i=1}^{r} n_i$
  - Overall mean:  $ar{Y} = ar{Y}_{\cdot \cdot \cdot} = rac{1}{N} \sum_{i=1}^r \sum_{j=1}^{n_i} Y_{ij}$
  - Pooled variance estimate:

$$S_p^2 = rac{\Sigma_{i=1}^r (n_i-1) S_i^2}{N-r}$$
 with df  $=\sum\limits_{i=1}^r (n_i-1) = N-r$ 

# Inference about Means for Several Populations

- Basic linear model
- Analysis of Variance (ANOVA)
  - F-tests
  - Contrasts
- Model diagnostics
- Nonparametric tests

## **Research Question**

- Do the populations or treatment groups have the same mean values for the variable?
- Two sources of variation
  - Variability among observations within each treatment group (or within each population)
  - Variability among mean responses for treatments (or between populations)
- Question:
  - Are differences among group means large relative to variation within groups?
  - Do all populations have the same mean?

#### **Cell Means Model**

$$Y_{ij} = \mu_i + \epsilon_{ij}$$

- ullet Each observation  $Y_{ij}$  can be described by two components:
  - Fixed mean value  $\mu_i$
  - Random error term  $\epsilon_{ij}$
- ullet Gives an equation for each of the  $N=\Sigma_{i=1}^r\,n_i$  observations

#### Cell Means Model in Matrix Notation

We can write this system of N equations in matrix notation

$$\begin{pmatrix} Y_{11} \\ Y_{12} \\ \vdots \\ Y_{1n_1} \\ Y_{21} \\ \vdots \\ Y_{2n_2} \\ Y_{31} \\ \vdots \\ Y_{rn_r} \end{pmatrix} = \begin{pmatrix} \mu_1 + \epsilon_{11} \\ \mu_1 + \epsilon_{12} \\ \vdots \\ \mu_1 + \epsilon_{1n_1} \\ \mu_2 + \epsilon_{21} \\ \vdots \\ \mu_2 + \epsilon_{2n_2} \\ \mu_3 + \epsilon_{31} \\ \vdots \\ \mu_r + \epsilon_{rn_r} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 & \cdots & 0 \\ 1 & 0 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 0 & \cdots & 0 \\ 0 & 1 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ 0 & 1 & 0 & \cdots & 0 \\ 0 & 0 & 1 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & 1 \end{pmatrix} \begin{pmatrix} \mu_1 \\ \mu_2 \\ \mu_3 \\ \vdots \\ \mu_r \end{pmatrix} + \begin{pmatrix} \epsilon_{11} \\ \epsilon_{12} \\ \vdots \\ \epsilon_{1n_1} \\ \epsilon_{21} \\ \vdots \\ \epsilon_{2n_2} \\ \epsilon_{31} \\ \vdots \\ \epsilon_{rn_r} \end{pmatrix}$$

#### Cell Means Model in Matrix Notation

Let

$$\mathbf{Y} = \begin{bmatrix} Y_{11} \\ Y_{12} \\ \vdots \\ Y_{1n_1} \\ Y_{21} \\ \vdots \\ Y_{2n_2} \\ Y_{31} \\ \vdots \\ Y_{rn_r} \end{bmatrix}, \mathbf{X} = \begin{bmatrix} 1 & 0 & 0 & \cdots & 0 \\ 1 & 0 & 0 & \cdots & 0 \\ 1 & 0 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 0 & \cdots & 0 \\ 0 & 1 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ 0 & 1 & 0 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & 1 \end{bmatrix}, \boldsymbol{\beta} = \begin{bmatrix} \mu_1 \\ \mu_2 \\ \mu_3 \\ \vdots \\ \mu_r \end{bmatrix}, \quad \text{and } \boldsymbol{\epsilon} = \begin{bmatrix} \epsilon_{11} \\ \epsilon_{12} \\ \vdots \\ \epsilon_{1n_1} \\ \epsilon_{21} \\ \vdots \\ \epsilon_{2n_2} \\ \epsilon_{31} \\ \vdots \\ \epsilon_{rn_r} \end{bmatrix}$$

#### **Linear Model**

Cell Means Model is an example of a **linear model** in matrix form:

$$Y = X\beta + \epsilon$$

- ullet The vector Y is length N and is the vector of observations.
- The matrix X is size  $N \times r$  and is called the design matrix. It relates the observations to the parameters according to the model. It is fixed (non-random).
- ullet The vector eta is length  $oldsymbol{r}$  and is the vector of parameter values.
- ullet The vector  $\epsilon$  is length N and is the vector of random error terms.

## **Expected Values: Linear Model**

ullet Assuming  $E(\epsilon)=0$ , we have

$$egin{array}{lll} E(\mathrm{Y}) &=& E(Xeta+\epsilon) \ &=& Xeta+E(\epsilon) \ &=& Xeta+0 \ &=& Xeta \end{array}$$

#### **Cell Means Model**

Expected Value for the cell means model:

$$E(\mathbf{Y}) = egin{pmatrix} 1 & 0 & 0 & \cdots & 0 \ 1 & 0 & 0 & \cdots & 0 \ \vdots & \vdots & \vdots & \vdots & \vdots \ 1 & 0 & 0 & \cdots & 0 \ 0 & 1 & 0 & \cdots & 0 \ \vdots & \vdots & \vdots & \vdots & \vdots \ 0 & 1 & 0 & \cdots & 0 \ 0 & 0 & 1 & \cdots & 0 \ \vdots & \vdots & \vdots & \vdots & \vdots \ 0 & 0 & 0 & \cdots & 1 \end{pmatrix} egin{pmatrix} \mu_1 \ \mu_2 \ \mu_3 \ \vdots \ \mu_r \end{pmatrix} = egin{pmatrix} \mu_1 \ \mu_1 \ \mu_2 \ \vdots \ \mu_2 \ \vdots \ \mu_r \end{pmatrix}$$

## **Least Squares Estimation**

- ullet Using our data, we will estimate the parameters in the eta vector using the method of least squares.
- Least squares estimation: Find the estimates of the population parameters that minimize the sum of squared deviations between the observed outcomes and the estimates of the expected outcomes. For cell means model, find  $\hat{\mu}_1$ ,  $\hat{\mu}_2$ ,  $\cdots$   $\hat{\mu}_r$  that minimize

$$\sum\limits_{i=1}^{r}\sum\limits_{j=1}^{n_i}(Y_{ij}-\hat{\mu}_i)^2$$

or, equivalently,

$$(\mathbf{Y} - X\hat{\boldsymbol{\beta}})^T (\mathbf{Y} - X\hat{\boldsymbol{\beta}})$$

## **Least Squares Estimation**

- ullet If the design matrix  $oldsymbol{X}$  is of full column rank, then
  - value of the parameter vector  $oldsymbol{eta}$  that minimizes the squared errors is

$$\hat{\beta} = (X^T X)^{-1} X^T Y$$

- value  $\hat{\beta}$  is unique since  $(X^TX)^{-1}$  is unique.

#### **Cell Means Model**

- ullet Design matrix  $oldsymbol{X}$  has full column rank.
- Unique least squares estimator for the parameter vector  $\beta$  is:

$$\hat{\beta} = (X^T X)^{-1} X^T Y$$

$$=egin{bmatrix} n_1 & 0 & 0 & \cdots & 0 \ 0 & n_2 & 0 & \cdots & 0 \ 0 & 0 & n_3 & \cdots & 0 \ dots & dots & dots & dots & dots & dots \ 0 & 0 & 0 & \cdots & n_r \end{bmatrix}^{-1} egin{pmatrix} \Sigma_{j=1}^{n_1} Y_{1j} \ \Sigma_{j=1}^{n_2} Y_{2j} \ dots \ \Sigma_{j=1}^{n_r} Y_{rj} \end{pmatrix} = egin{pmatrix} ar{Y}_1. \ ar{Y}_2. \ dots \ ar{Y}_r. \end{pmatrix}$$

 This is the least squares estimator for the population parameters (population means)

#### Predicted Values: Linear Model

Using the least squares estimator  $\hat{\beta}$ , the predicted value for Y is:

$$\hat{\mathbf{Y}} = X\hat{\boldsymbol{\beta}}$$

$$= X(X^TX)^{-1}X^TY$$

$$= P_XY$$

where  $P_X = X(X^TX)^{-1}X^T$  is the orthogonal projection operator onto the column space of matrix X.