

Lecture 4

Inference and Comparison of Mean Vectors

Readings: Johnson & Wichern 2007, Chapter 5.1-5.4; 6.1-6.4;
6.8

DSA 8070 Multivariate Analysis

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

Whitney Huang
Clemson University

1 Confidence Intervals/Region for Population Means

Confidence
Intervals/Region for
Population Means

2 Hypothesis Testing for Mean Vector

Hypothesis Testing for
Mean Vector

3 Multivariate Paired Hotelling's T-Square

Multivariate Paired
Hotelling's T-Square

4 Comparisons of Two Mean Vectors

Comparisons of Two
Mean Vectors

5 Multivariate Analysis of Variance

Multivariate Analysis of
Variance

This Week's Topics:

- **Single Mean Vector:** inference on μ (multivariate one-sample t -test)
- **Paired Mean Vectors:** differences between paired observations \Rightarrow reduce to one-sample Hotelling's T^2 on differences
- **Two Independent Mean Vectors:** Hotelling's T^2 two-sample test
- **Several Mean Vectors:** MANOVA (multivariate extension of ANOVA)

Analogy with Univariate Methods:

- One-sample t -test \rightarrow single μ
- Paired t -test \rightarrow paired mean vectors
- Two-sample t -test \rightarrow two mean vectors
- ANOVA \rightarrow MANOVA

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

Review: Sampling Distribution of Univariate Sample Mean \bar{X}_n

Inference and
Comparison of Mean
Vectors

CLEMSON
UNIVERSITY

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

Suppose X_1, X_2, \dots, X_n is a random sample from a univariate population distribution with mean $\mathbb{E}(X) = \mu$ and variance $\text{Var}(X) = \sigma^2$. The sample mean \bar{X}_n is a function of random sample and therefore has a distribution

- $\bar{X}_n \sim N(\mu, \frac{\sigma^2}{n})$ when the sample size n is “sufficiently” large \Rightarrow This is the central limit theorem (CLT)
- The result above is exact if the population follows a normal distribution, i.e., $X \sim N(\mu, \sigma^2)$
- The standard error $\sqrt{\text{Var}(\bar{X}_n)} = \frac{\sigma}{\sqrt{n}}$ provides a measure estimation precision. In practice, we use $\frac{s}{\sqrt{n}}$ instead where s is the sample standard deviation

Suppose X_1, X_2, \dots, X_n is a random sample from a multivariate population distribution with mean vector $\mathbb{E}(X) = \mu$ and covariance matrix $= \Sigma$.

- $\bar{X}_n \sim N(\mu, \frac{1}{n}\Sigma)$ when the sample size n is “sufficiently” large \Rightarrow This is the multivariate version of CLT
- The result above is exact if the population follows a normal distribution, i.e., $X \sim N(\mu, \Sigma)$
- Again, the estimation precision improves with a larger sample size. Like the univariate case we would need to replace Σ by its estimate S , the sample covariance matrix

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

Review: Interval Estimation of Univariate Population Mean μ

Inference and
Comparison of Mean
Vectors

CLEMSON
UNIVERSITY

The general format of a confidence interval (CI) estimate of a population mean is

Sample mean \pm multiplier \times standard error of mean.

For variable X , a CI estimate of its population mean μ is

$$\bar{X}_n \pm t_{n-1, \frac{\alpha}{2}} \frac{s}{\sqrt{n}},$$

Here the multiplier value is a function of the confidence level, α , the sample size n

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

Constructing Confidence Intervals for Mean Vector

We will still use the general recipe

Sample mean \pm multiplier \times standard error of mean.

The multiplier value also depends the strategy used for dealing with the multiple inference issue

- **One at a Time CIs:** a CI for μ_j is computed as

$$\bar{x}_j \pm t_{n-1, \frac{\alpha}{2}} \frac{s_j}{\sqrt{n}}, \quad j = 1, \dots, p$$

- **Bonferroni Method:** a CI for μ_j is computed as

$$\bar{x}_j \pm t_{n-1, \frac{\alpha}{2p}} \frac{s_j}{\sqrt{n}}, \quad j = 1, \dots, p$$

- **Simultaneous CIs:** a CI for μ_j is computed as

$$\bar{x}_j \pm \sqrt{\frac{(n-1)p}{n-p} F_{p, n-p, \alpha}} \frac{s_j}{\sqrt{n}}, \quad j = 1, \dots, p$$

Example: Mineral Content Measurements [source: Penn Stat Univ. STAT 505]

This example uses a dataset that includes mineral content measurements at two different arm bone locations for $n = 64$ women. We will determine confidence intervals for the two population means. The sample means and standard deviations for the two variables are:

Variable	Sample size	Mean	Std Dev
domradius (X_1)	$n = 64$	$\bar{x}_1 = 0.8438$	$s_1 = 0.1140$
domhumerus (X_2)	$n = 64$	$\bar{x}_2 = 1.7927$	$s_2 = 0.2835$

Let's apply the three methods we learned to construct 95% CIs

Mineral Content Measurements Example Cont'd

- **One at a Time CIs:** $\bar{x}_j \pm t_{n-1, \alpha/2} \frac{s_j}{\sqrt{n}}$, $j = 1, \dots, p$. Therefore 95% CIs for μ_1 and μ_2 are:

$$\mu_1 : 0.8438 \pm \underbrace{1.998}_{t_{63, 0.025}} \times \frac{0.1140}{\sqrt{64}} = [0.815, 0.872]$$

$$\mu_2 : 1.7927 \pm 1.998 \times \frac{0.2835}{\sqrt{64}} = [1.722, 1.864]$$

- **Bonferroni Method:** $\bar{x}_j \pm t_{n-1, \alpha/2p} \frac{s_j}{\sqrt{n}}$, $j = 1, \dots, p$.

$$\mu_1 : 0.8438 \pm \underbrace{2.296}_{t_{63, 0.0125}} \times \frac{0.1140}{\sqrt{64}} = [0.811, 0.877]$$

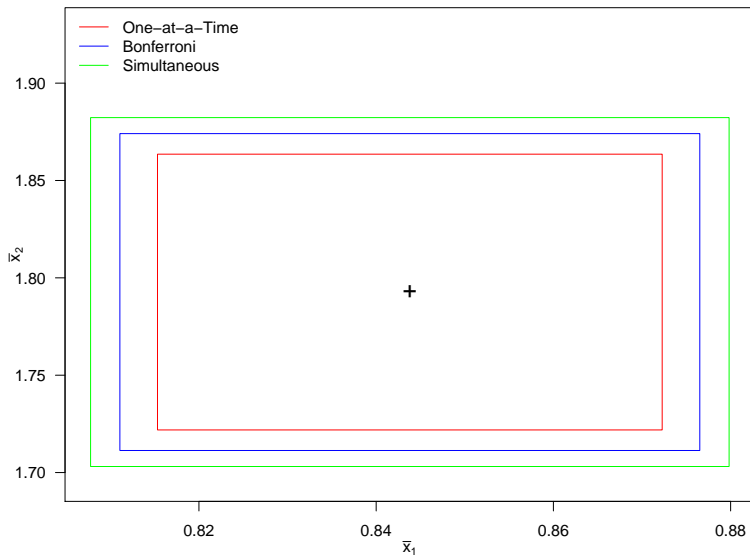
$$\mu_2 : 1.7927 \pm 2.296 \times \frac{0.2835}{\sqrt{64}} = [1.711, 1.874]$$

- **Simultaneous CIs:** $\bar{x}_j \pm \sqrt{\frac{(n-1)p}{n-p} F_{p, n-p, \alpha}} \frac{s_j}{\sqrt{n}}$, $j = 1, \dots, p$

$$\mu_1 : 0.8438 \pm 2.528 \times \frac{0.1140}{\sqrt{64}} = [0.808, 0.880]$$

$$\mu_2 : 1.7927 \pm 2.528 \times \frac{0.2835}{\sqrt{64}} = [1.703, 1.882]$$

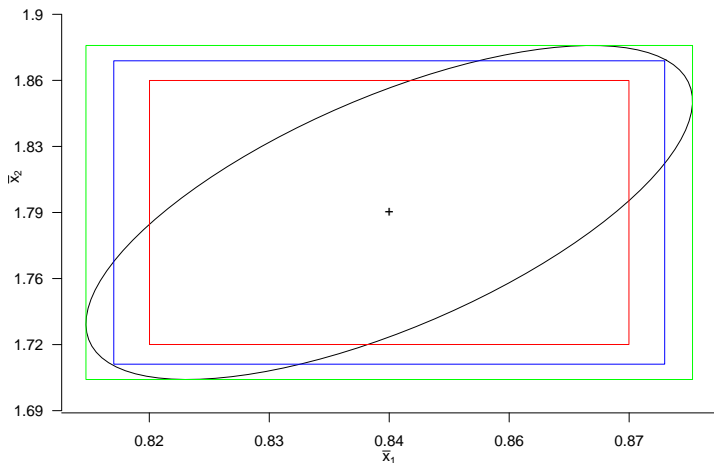
95 % CIs Based on Three Methods



Confidence Ellipsoid

A confidence ellipsoid for μ is the set of μ satisfying

$$n(\bar{X}_n - \mu)^T S^{-1}(\bar{X}_n - \mu) \leq \frac{(n-1)p}{n-p} F_{p, n-p, \alpha}$$



Hypothesis Testing for Mean

- Recall: for univariate data, t statistic

$$t = \frac{\bar{X}_n - \mu_0}{s/\sqrt{n}} \Rightarrow t^2 = \frac{(\bar{X}_n - \mu_0)^2}{s^2/n} = n(\bar{X}_n - \mu_0)(s^2)^{-1}(\bar{X}_n - \mu_0)$$

Under $H_0 : \mu = \mu_0$

$$t \sim t_{n-1}, \quad t^2 \sim F_{1,n-1}$$

- Extending to multivariate by analogy:

$$T^2 = n(\bar{\mathbf{X}}_n - \boldsymbol{\mu}_0)^T \mathbf{S}^{-1}(\bar{\mathbf{X}}_n - \boldsymbol{\mu}_0)$$

Under $H_0 : \boldsymbol{\mu} = \boldsymbol{\mu}_0$

$$\frac{(n-p)}{(n-1)p} T^2 \sim F_{p,n-p}$$

Note: T^2 here is the so-called **Hotelling's T-Square**

- 1 State the null

$$H_0 : \mu = \mu_0$$

and the alternative

$$H_a : \mu \neq \mu_0$$

- 2 Compute the test statistic

$$F = \frac{n-p}{(n-1)p} n (\bar{X}_n - \mu_0)^T S^{-1} (\bar{X}_n - \mu_0)$$

- 3 **Compute the P-value.** Under $H_0 : F \sim F_{p, n-p}$

- 4 **Draw a conclusion:** We do (or do not) have enough statistical evidence to conclude $\mu \neq \mu_0$ at α significant level

Example: Women's Dietary Intake [source: Penn Stat Univ. STAT 505]

The recommended intake and a sample mean for all women between 25 and 50 years old are given below:

Variable	Recommended Intake (μ_0)	Sample Mean (\bar{x}_n)
Calcium	1000 <i>mg</i>	624.0 <i>mg</i>
Iron	15 <i>mg</i>	11.1 <i>mg</i>
Protein	60 <i>g</i>	65.8 <i>g</i>
Vitamin A	800 μg	839.6 μg
Vitamin C	75 <i>mg</i>	78.9 <i>mg</i>

Here we would like to test, at $\alpha = 0.01$ level, if the $\mu = \mu_0$

Women's Dietary Intake Example Analysis

- 1 State the null

$$H_0 : \boldsymbol{\mu} = \boldsymbol{\mu}_0$$

and the alternative

$$H_a : \boldsymbol{\mu} \neq \boldsymbol{\mu}_0$$

- 2 Compute the test statistic

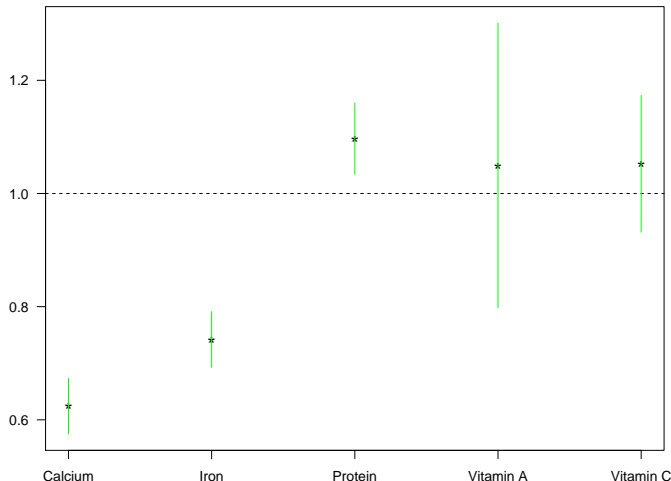
$$F = \frac{n-p}{(n-1)p} n (\bar{\mathbf{x}}_n - \boldsymbol{\mu}_0)^T \mathbf{S}^{-1} (\bar{\mathbf{x}}_n - \boldsymbol{\mu}_0) = 349.80$$

- 3 **Compute the P-value.** Under $H_0 : F \sim F_{p,n-p} \Rightarrow$ p-value
 $= \mathbb{P}(F_{p,n-p} > 349.80) = 3 \times 10^{-191} < \alpha = 0.01$

- 4 **Draw a conclusion:** We do have enough statistical
evidence to conclude $\boldsymbol{\mu} \neq \boldsymbol{\mu}_0$ at $\alpha = 0.01$ significant level

Profile Plots

- 1 Standardize each of the observations by dividing their hypothesized means
- 2 Plot either simultaneous or Bonferroni CIs for the population mean of these standardized variables



Spouse Survey Data Example

A sample ($n = 30$) of husband and wife pairs are asked to respond to each of the following questions:

- 1 What is the level of passionate love you feel for your partner?
- 2 What is the level of passionate love your partner feels for you?
- 3 What is the level of companionate love you feel for your partner?
- 4 What is the level of companionate love your partner feels for you?

Responses were recorded on a typical five-point scale: 1) None at all 2) Very little 3) Some 4) A great deal 5) Tremendous amount.

We will try to address the following question: Do the husbands respond to the questions in the same way as their wives?

Multivariate Paired Hotelling's T-Square

Let X_F and X_M be the responses to these 4 questions for females and males, respectively. Here the quantities of interest are $\mathbb{E}(\mathbf{D}) = \boldsymbol{\mu}_D$, the average differences across all husband and wife pairs.

- 1 State the null $H_0 : \boldsymbol{\mu}_D = \mathbf{0}$ and the alternative hypotheses $H_a : \boldsymbol{\mu}_D \neq \mathbf{0}$
- 2 Compute the test statistic

$$F = \frac{n-p}{(n-1)p} n \bar{\mathbf{D}}_n^T \mathbf{S}_D^{-1} \bar{\mathbf{D}}_n$$

- 3 **Compute the P-value.** Under $H_0 : F \sim F_{p, n-p}$
- 4 **Draw a conclusion:** We do (or do not) have enough statistical evidence to conclude $\boldsymbol{\mu}_D \neq \mathbf{0}$ at α significant level

- 1 State the null

$$H_0 : \mu_D = \mathbf{0}$$

and the alternative

$$H_a : \mu_D \neq \mathbf{0}$$

- 2 Compute the test statistic

$$F = \frac{n-p}{(n-1)p} n \bar{\mathbf{D}}_n^T \mathbf{S}_D^{-1} \bar{\mathbf{D}}_n = 2.942$$

- 3 **Compute the P-value.** Under $H_0 : F \sim F_{p,n-p} \Rightarrow$ p-value
 $= \mathbb{P}(F_{p,n-p} >) = 0.0394 < \alpha = 0.05$

- 4 **Draw a conclusion:** We do have enough statistical
evidence to conclude $\mu_D \neq \mathbf{0}$ at 0.05 significant level

Motivating Example: Swiss Bank Notes (Source: PSU stat 505)

Suppose there are two distinct populations for 1000 franc Swiss Bank Notes:

- The first population is the population of Genuine Bank Notes
- The second population is the population of Counterfeit Bank Notes

For both populations the following measurements were taken:

- 1 Length of the note
- 2 Width of the Left-Hand side of the note
- 3 Width of the Right-Hand side of the note
- 4 Width of the Bottom Margin
- 5 Width of the Top Margin
- 6 Diagonal Length of Printed Area

We want to determine if counterfeit notes can be distinguished from the genuine Swiss bank notes

Review: Two Sample t-Test

Suppose we have data from a single variable from population 1: $X_{11}, X_{12}, \dots, X_{1n_1}$ and population 2: $X_{21}, X_{22}, \dots, X_{2n_2}$. Here we would like to draw inference about their population means μ_1 and μ_2 .

Assumptions:

- **Homoscedasticity**: The data from both populations have common variance σ^2
- **Independence**: The subjects from both populations are independently sampled $\Rightarrow \{X_{1i}\}_{i=1}^{n_1}$ and $\{X_{2j}\}_{j=1}^{n_2}$ are independent to each other
- **Normality**: The data from both populations are normally distributed (not that crucial for “large” sample)

Here we are going to consider testing $H_0 : \mu_1 = \mu_2$ against $H_a : \mu_1 \neq \mu_2$

Review: Two Sample t-Test

We define the sample means for each population using the following expression:

$$\bar{x}_1 = \frac{\sum_{j=1}^{n_1} x_{1j}}{n_1}, \quad \bar{x}_2 = \frac{\sum_{j=1}^{n_2} x_{2j}}{n_2}.$$

We denote the sample variance

$$s_1^2 = \frac{\sum_{j=1}^{n_1} (x_{1j} - \bar{x}_1)^2}{n_1 - 1}, \quad s_2^2 = \frac{\sum_{j=1}^{n_2} (x_{2j} - \bar{x}_2)^2}{n_2 - 1}.$$

Under the **homoscedasticity** assumption, we can “pool” two samples to get the pooled sample variance

$$s_p^2 = \frac{(n_1 - 1)s_1^2 + (n_2 - 1)s_2^2}{n_1 + n_2 - 2}$$

Test statistic

$$t = \frac{\bar{x}_1 - \bar{x}_2}{\sqrt{s_p^2 \left(\frac{1}{n_1} + \frac{1}{n_2} \right)}} \stackrel{H_0}{\sim} t_{n_1 + n_2 - 2}$$

We can use this result to construct confidence intervals and to perform hypothesis tests

The Two Sample Problem: The Multivariate Case

Now we would like to use two independent samples

$\{X_{11}, \dots, X_{12}, \dots, X_{1n_1}\}$ and $\{X_{21}, \dots, X_{22}, \dots, X_{2n_2}\}$, where

$$X_{ij} = \begin{bmatrix} X_{ij1} \\ X_{ij2} \\ \vdots \\ X_{ijp} \end{bmatrix}$$

to infer the relationship between μ_1 and μ_2 , where

$$\mu_i = \begin{bmatrix} \mu_{i1} \\ \mu_{i2} \\ \vdots \\ \mu_{ip} \end{bmatrix}$$

Assumptions

- Both populations have common covariance matrix, i.e.,
 $\Sigma_1 = \Sigma_2$
- Independence**: The subjects from both populations are independently sampled
- Normality**: Both populations are normally distributed

The Multivariate Two-Sample Problem

Here we are testing

$$H_0 : \begin{bmatrix} \mu_{11} \\ \mu_{12} \\ \vdots \\ \mu_{1p} \end{bmatrix} = \begin{bmatrix} \mu_{21} \\ \mu_{22} \\ \vdots \\ \mu_{2p} \end{bmatrix}, \quad H_a : \mu_{1k} \neq \mu_{2k} \text{ for at least one } k \in \{1, 2, \dots, p\}$$

Under the **common covariance** assumption we have

$$\mathbf{S}_p = \frac{(n_1 - 1)\mathbf{S}_1 + (n_2 - 1)\mathbf{S}_2}{n_1 + n_2 - 2},$$

where

$$\mathbf{S}_i = \frac{1}{n_i - 1} \sum_{j=1}^{n_i} (\mathbf{x}_{ij} - \bar{\mathbf{x}}_i)(\mathbf{x}_{ij} - \bar{\mathbf{x}}_i)^T, \quad i = 1, 2$$

The Two-Sample Hotelling's T-Square Test Statistic

The two-sample t test is equivalent to

$$t^2 = (\bar{x}_1 - \bar{x}_2)^T \left[s_p^2 \left(\frac{1}{n_1} + \frac{1}{n_2} \right) \right]^{-1} (\bar{x}_1 - \bar{x}_2).$$

Under H_0 , $t^2 \sim F_{1, n_1 + n_2 - 2}$. We can use this result to perform a hypothesis test

We can extend this to the multivariate situation:

$$T^2 = (\bar{\mathbf{x}}_1 - \bar{\mathbf{x}}_2)^T \left[\mathbf{S}_p \left(\frac{1}{n_1} + \frac{1}{n_2} \right) \right]^{-1} (\bar{\mathbf{x}}_1 - \bar{\mathbf{x}}_2)$$

Under H_0 , we have

$$F = \frac{n_1 + n_2 - p - 1}{p(n_1 + n_2 - 2)} T^2 \sim F_{p, n_1 + n_2 - p - 1}$$

We can use this result to perform inferences for multivariate cases

Two-Sample Test for Swiss Bank Notes

```
> (xbar1 <- colMeans(dat[real, -1]))
      V2      V3      V4      V5      V6      V7
214.969 129.943 129.720   8.305  10.168 141.517
> (xbar2 <- colMeans(dat[fake, -1]))
      V2      V3      V4      V5      V6      V7
214.823 130.300 130.193  10.530  11.133 139.450
> Sigma1 <- cov(dat[real, -1])
> Sigma2 <- cov(dat[fake, -1])
> n1 <- length(real); n2 <- length(fake); p <- dim(dat[, -1])[2]
> Sp <- ((n1 - 1) * Sigma1 + (n2 - 1) * Sigma2) / (n1 + n2 - 2)
> # Test statistic
> T.squared <- as.numeric(t(xbar1 - xbar2) %*% solve(Sp * (1 / n1 + 1 / n2)) %*% (xbar1 - xbar2))
> Fobs <- T.squared * ((n1 + n2 - p - 1) / ((n1 + n2 - 2) * p))
> # p-value
> pf(Fobs, p, n1 + n2 - p - 1, lower.tail = F)
[1] 3.378887e-105
```

Conclusion

The counterfeit notes can be distinguished from the genuine notes on at least one of the measurements \Rightarrow which ones?

Simultaneous Confidence Intervals

$$\bar{x}_{1k} - \bar{x}_{2k} \pm \sqrt{\frac{p(n_1 + n_2 - 2)}{n_1 + n_2 - p - 1} F_{p, n_1 + n_2 - p - 1, \alpha}} \sqrt{\left(\frac{1}{n_1} + \frac{1}{n_2}\right) s_{k,p}^2},$$

where $s_{k,p}^2$ is the pooled variance for the variable k

Variable	95% CI
Length of the note	(-0.04, 0.34)
Width of the Left-Hand note	(-0.52, -0.20)
Width of the Right-Hand note	(-0.64, -0.30)
Width of the Bottom Margin	(-2.70, -1.75)
Width of the Top Margin	(-1.30, -0.63)
Diagonal Length of Printed Area	(1.81, 2.33)

Assumptions:

- **Homoscedasticity:** The data from both populations have common covariance matrix Σ

Will return to this in next slide

- **Independence:**

This assumption may be violated if we have clustered, time-series, or spatial data

- **Normality:**

Multivariate QQplot, univariate histograms, bivariate scatter plots

Testing for Equality of Mean Vectors when $\Sigma_1 \neq \Sigma_2$

- Bartlett's test can be used to test if $\Sigma_1 = \Sigma_2$ but this test is sensitive to departures from normality
- As a crude rule of thumb: if $s_{1,k}^2 > 4s_{2,k}^2$ or $s_{2,k}^2 > 4s_{1,k}^2$ for some $k \in \{1, 2, \dots, p\}$, then it is likely that $\Sigma_1 \neq \Sigma_2$
- Life gets difficult if we cannot assume that $\Sigma_1 = \Sigma_2$. However, if both n_1 and n_2 are "large", we can use the following approximation to conduct inferences:

$$T^2 = (\bar{\mathbf{X}}_1 - \bar{\mathbf{X}}_2)^T \left[\frac{1}{n_1} \mathbf{S}_1 + \frac{1}{n_2} \mathbf{S}_2 \right]^{-1} (\bar{\mathbf{X}}_1 - \bar{\mathbf{X}}_2) \stackrel{H_0}{\sim} \chi_p^2$$

Comparing More Than Two Populations: Romano-British Pottery Example (source: PSU stat 505)

- Pottery shards are collected from four sites in the British Isles:
 - Llanedyrn (L)
 - Caldicot (C)
 - Isle Thorns (I)
 - Ashley Rails (A)
- The concentrations of five different chemicals were be used
 - Aluminum (Al)
 - Iron (Fe)
 - Magnesium (Mg)
 - Calcium (Ca)
 - Sodium (Na)
- **Objective:** to determine whether the chemical content of the pottery depends on the site where the pottery was obtained

Review: (Univariate) Analysis of Variance (ANOVA)

- $H_0 : \mu_1 = \mu_2 = \cdots = \mu_g$
 $H_a : \text{At least one mean is different}$

Source	df	SS	MS	F statistic
Treatment	$g - 1$	$SSTr$	$MSTr = \frac{SSTr}{g-1}$	$F = \frac{MSTr}{MSE}$
Error	$N - g$	SSE	$MSE = \frac{SSE}{N-g}$	
Total	$N - 1$	$SSTo$		

- Test Statistic: $F^* = \frac{MSTr}{MSE}$. Under H_0 , $F^* \sim F_{df_1=g-1, df_2=N-g}$
- **Assumptions:**
 - The distribution of each group is normal with equal variance (i.e. $\sigma_1^2 = \sigma_2^2 = \cdots = \sigma_g^2$)
 - Responses for a given group are independent to each other

One-way Multivariate Analysis of Variance (One-way MANOVA)

Subject \ Group	1	2	...	g
1	$\mathbf{Y}_{11} = \begin{bmatrix} Y_{111} \\ Y_{112} \\ \vdots \\ Y_{11p} \end{bmatrix}$	$\mathbf{Y}_{21} = \begin{bmatrix} Y_{211} \\ Y_{212} \\ \vdots \\ Y_{21p} \end{bmatrix}$...	$\mathbf{Y}_{g1} = \begin{bmatrix} Y_{g11} \\ Y_{g12} \\ \vdots \\ Y_{g1p} \end{bmatrix}$
2	$\mathbf{Y}_{12} = \begin{bmatrix} Y_{121} \\ Y_{122} \\ \vdots \\ Y_{12p} \end{bmatrix}$	$\mathbf{Y}_{22} = \begin{bmatrix} Y_{221} \\ Y_{222} \\ \vdots \\ Y_{22p} \end{bmatrix}$...	$\mathbf{Y}_{g2} = \begin{bmatrix} Y_{g21} \\ Y_{g22} \\ \vdots \\ Y_{g2p} \end{bmatrix}$
\vdots	\vdots	\vdots	...	\vdots
n_i	$\mathbf{Y}_{1n_i} = \begin{bmatrix} Y_{1n_i1} \\ Y_{1n_i2} \\ \vdots \\ Y_{1n_ip} \end{bmatrix}$	$\mathbf{Y}_{2n_i} = \begin{bmatrix} Y_{2n_i1} \\ Y_{2n_i2} \\ \vdots \\ Y_{2n_ip} \end{bmatrix}$...	$\mathbf{Y}_{gn_i} = \begin{bmatrix} Y_{gn_i1} \\ Y_{gn_i2} \\ \vdots \\ Y_{gn_ip} \end{bmatrix}$

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

- **Notation:** \mathbf{Y}_{ij} is the vector of variables for subject j in group i ; n_i is the sample size in group i ;
 $N = n_1 + n_2 + \dots + n_g$ the total sample size
- **Assumptions:** 1) common covariance matrix Σ ; 2) Independence; 3) Normality

Test Statistics for MANOVA

- We are interested in testing the null hypothesis that the group mean vectors are all equal

$$H_0 : \mu_1 = \mu_2 = \cdots = \mu_g.$$

The alternative hypothesis:

$H_a : \mu_{ik} \neq \mu_{jk}$ for at least one $i \neq j$ and at least one variable k

- **Mean vectors:**

- **Sample Mean Vector:** $\bar{\mathbf{y}}_{i.} = \frac{1}{n_i} \mathbf{Y}_{ij}, \quad i = 1, \dots, g$
- **Grand Mean Vector:** $\bar{\mathbf{y}}_{..} = \frac{1}{N} \sum_{i=1}^g \sum_{j=1}^{n_i} \mathbf{Y}_{ij}$

- **Total Sum of Squares:**

$$\mathbf{T} = \sum_{i=1}^g \sum_{j=1}^{n_i} (\mathbf{Y}_{ij} - \bar{\mathbf{y}}_{..})(\mathbf{Y}_{ij} - \bar{\mathbf{y}}_{..})^T$$

MANOVA Decomposition and MANOVA Table

$$\begin{aligned} \mathbf{T} &= \sum_{i=1}^g \sum_{j=1}^{n_i} (\mathbf{Y}_{ij} - \mathbf{y}_{..})(\mathbf{Y}_{ij} - \bar{\mathbf{y}})^T \\ &= \sum_{i=1}^g \sum_{j=1}^{n_i} [(\mathbf{Y}_{ij} - \bar{\mathbf{y}}_{i.}) + (\bar{\mathbf{y}}_{i.} - \bar{\mathbf{y}}_{..})][(\mathbf{Y}_{ij} - \bar{\mathbf{y}}_{i.}) + (\bar{\mathbf{y}}_{i.} - \bar{\mathbf{y}}_{..})]^T \\ &= \underbrace{\sum_{i=1}^g \sum_{j=1}^{n_i} (\mathbf{Y}_{ij} - \bar{\mathbf{y}}_{i.})(\mathbf{Y}_{ij} - \bar{\mathbf{y}}_{i.})^T}_{\mathbf{E}} + \underbrace{\sum_{i=1}^g n_i (\bar{\mathbf{y}}_{i.} - \bar{\mathbf{y}}_{..})(\bar{\mathbf{y}}_{i.} - \bar{\mathbf{y}}_{..})^T}_{\mathbf{H}} \end{aligned}$$

MANOVA Table

Source	df	SS
Treatment	$g - 1$	\mathbf{H}
Error	$N - g$	\mathbf{E}
Total	$N - 1$	\mathbf{T}

Reject $H_0 : \mu_1 = \mu_2 = \cdots = \mu_g$ if the matrix \mathbf{H} is “large” relative to the matrix \mathbf{E}

There are several different test statistics for conducting the hypothesis test:

- Wilks Lambda

$$\Lambda^* = \frac{|E|}{|H + E|}$$

Reject H_0 if Λ^* is “small”

- Hotelling-Lawley Trace

$$T_0^2 = \text{trace}(HE^{-1})$$

Reject H_0 if T_0^2 is “large”

- Pillai Trace

$$V = \text{trace}(H(H + E)^{-1})$$

Reject H_0 if V is “large”

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance

Romano-British Pottery Example

```
> dat <- read.table("pottery.txt", header = F)
> out <- manova(cbind(V2, V3, V4, V5, V6) ~ V1, data = dat)
> summary(out, test = "Wilks")
              Df      Wilks approx F num Df den Df    Pr(>F)
V1              3 0.012301   13.088     15 50.091 1.84e-12 ***
Residuals 22
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> summary(out)
              Df Pillai approx F num Df den Df    Pr(>F)
V1              3 1.5539   4.2984     15     60 2.413e-05 ***
Residuals 22
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

⇒ at least one of the chemicals differs among the sites

Summary

In this lecture, we learned about:

- Confidence Intervals/Regions for Mean Vector
- Hypothesis Testing for Mean Vector
- Multivariate Version of Paired Tests
- Hypothesis Testing for Two Mean Vectors
- MANOVA

In the next two lectures, we will learn about **Multivariate Regression**

Confidence
Intervals/Region for
Population Means

Hypothesis Testing for
Mean Vector

Multivariate Paired
Hotelling's T-Square

Comparisons of Two
Mean Vectors

Multivariate Analysis of
Variance