# Lecture 6 Comparisons of Several Mean Vectors Readings: Johnson & Wichern 2007, Chapter 6.3-6.5

DSA 8070 Multivariate Analysis

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Comparisons of Two Mean Vectors

# Agenda

Comparisons of Several Mean Vectors



Comparisons of Two Mean Vectors

Multivariate Analysis of Variance



# Comparisons of Two Mean Vectors



# 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:

- Length of the note
- Width of the Left-Hand side of the note
- Width of the Right-Hand side of the note
- Width of the Bottom Margin
- Width of the Top Margin
- Diagonal Length of Printed Area

We want to determine if counterfeit notes can be distinguished from the genuine Swiss bank notes Comparisons of Several Mean Vectors



Comparisons of Two Mean Vectors

# **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  $\mu_1$  and  $\mu_2$ .

### Assumptions:

- Homoscedasticity: The data from both populations have common variance  $\sigma^2$
- 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  $H_a: \mu_1 \neq \mu_2$ 

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Comparisons of Two Mean Vectors

#### **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

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

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





Comparisons of Two Mean Vectors

The Two Sample Problem: The Multivariate Case Now we would like to use two independent samples  $\{X_{11}, \cdots X_{12}, \cdots X_{1n_1}\}$  and  $\{X_{21}, \cdots X_{22}, \cdots X_{2n_2}\}$ , where

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

to infer the relationship between  $\mu_1$  and  $\mu_2$ , where

$$\boldsymbol{\mu}_i = \begin{bmatrix} \mu_{i1} \\ \mu_{i2} \\ \vdots \\ \mu_{ip} \end{bmatrix}$$

# 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

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Comparisons of Two Mean Vectors

#### The Multivariate Two-Sample Problem

### Here we are testing

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

Under the common covariance assumption we have

$$S_p = \frac{(n_1 - 1)S_1 + (n_2 - 1)S_2}{n_1 + n_2 - 2},$$

where

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





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### 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:

$$T^{2} = (\bar{x}_{1} - \bar{x}_{2})^{T} \left[ S_{p} \left( \frac{1}{n_{1}} + \frac{1}{n_{2}} \right) \right]^{-1} (\bar{x}_{1} - \bar{x}_{2})$$

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





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#### **Two-Sample Test for Swiss Bank Notes**

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



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#### 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

<u> </u>	0.50/ 01
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)





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# **Checking Model Assumptions**

# Assumptions:

• Homoscedasticity: The data from both populations have common covariance matrix  $\boldsymbol{\Sigma}$ 

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

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

- Bartlett's test can be used to test if Σ<sub>1</sub> = Σ<sub>2</sub> but this test is sensitive to departures from normality
- As as 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:

$$T^{2} = (\bar{\boldsymbol{X}}_{1} - \bar{\boldsymbol{X}}_{2})^{T} \left[ \frac{1}{n_{1}} \boldsymbol{S}_{1} + \frac{1}{n_{2}} \boldsymbol{S}_{2} \right]^{-1} (\bar{\boldsymbol{X}}_{1} - \bar{\boldsymbol{X}}_{2}) \stackrel{H_{0}}{\sim} \chi_{p}^{2}$$

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Comparisons of Two Mean Vectors

# 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





Comparisons of Two Mean Vectors

# Review: (Univariate) Analysis of Variance (ANOVA)

• 
$$H_0: \mu_1 = \mu_2 = \dots = \mu_g$$
  
 $H_a:$  At least one mean is different

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

• Test Statistic: 
$$F^* = \frac{\text{MSTr}}{\text{MSE}}$$
. 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





Comparisons of Two Mean Vectors

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

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





Comparisons of Two Mean Vectors

- Notation:  $Y_{ij}$  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

$$H_0: \boldsymbol{\mu}_1 = \boldsymbol{\mu}_2 = \dots = \boldsymbol{\mu}_g$$

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:  $\bar{y}_{i.} = \frac{1}{n_i} Y_{ij}, \quad i = 1, \dots, g$
- Grand Mean Vector:  $\bar{\boldsymbol{y}}_{..} = \frac{1}{N} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \boldsymbol{Y}_{ij}$
- Total Sum of Squares:

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





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#### **MANOVA Decomposition and MANOVA Table**

$$T = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - y_{..}) (Y_{ij} - \bar{y})^T$$
  
=  $\sum_{i=1}^{g} \sum_{j=1}^{n_i} [(Y_{ij} - \bar{y}_{i.}) + (\bar{y}_{i.} - \bar{y}_{..})] [(Y_{ij} - \bar{y}_{i.}) + (\bar{y}_{i.} - \bar{y}_{..})]^T$   
=  $\underbrace{\sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \bar{y}_{i.}) (Y_{ij} - \bar{y}_{i.})^T}_{E} + \underbrace{\sum_{i=1}^{g} n_i (\bar{y}_{i.} - \bar{y}_{..}) (\bar{y}_{i.} - \bar{y}_{..})^T}_{H}}_{H}$ 

MANOVA Table

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

Reject  $H_0: \mu_1 = \mu_2 = \dots = \mu_g$  if the matrix H is "large" relative to the matrix E





Comparisons of Two Mean Vectors

# **Test Statistics for MANOVA**

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

Wilks Lambda

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

Reject  $H_0$  if  $\Lambda^*$  is "small"

Hotelling-Lawley Trace

$$T_0^2 = \operatorname{trace}(\boldsymbol{H}\boldsymbol{E}^{-1})$$

Reject  $H_0$  if  $T_0^2$  is "large"

• Pillai Trace

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

Reject  $H_0$  if V is "large"





Comparisons of Two Mean Vectors

#### **Romano–British Pottery Example**

```
> dat <- read.table("pottery.txt", header = F)</pre>
> out <- manova(cbind(V2, V3, V4, V5, V6) \sim V1, data = dat)
> summary(out, test = "Wilks")
         Df Wilks approx F num Df den Df Pr(>F)
          3 0.012301 13.088 15 50.091 1.84e-12 ***
V1
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
```

#### $\Rightarrow$ at least one of the chemicals differs among the sites





Comparisons of Two Mean Vectors

# Summary

In this lecture, we learned about:

• Hypothesis Testing for Two Mean Vectors

MANOVA

In the next lecture, we will learn about Multivariate Linear Regression





Comparisons of Two Mean Vectors