Multivariate Analysis Lecture 8: Eigenvalues, Covariance Matrices, and MANOVA

MANOVA Sample Cov

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Outline of Lecture 08

- Eigenvalues of covariance matrices
- Examples of 2 × 2 covariance matrices
- Review of one-way ANOVA
- One-way MANOVA
- A heads up of the midterm project

Section 1

Eigenvalues and Cov Matrices

Eigenvalues

Subsection 1

Eigenvalues

Eigenvalues and Eigenvectors

- Let A be a $p \times p$ square matrix.
- We say $\nu \in \mathbb{R}^p$ is an eigenvector and $\lambda \in \mathbb{R}$ is the corresponding eigenvalue of A if

$$A\nu = \lambda \nu$$

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• The eigenvalues λ of A are roots of the characteristic equation

$$\det\left(\lambda I - A\right) = 0$$

The Spectral Decomposition of A Symmetric Matrix

• Let $A_{p \times p}$ be a symmetric matrix

$$A = \Gamma \Lambda \Gamma^T = \sum_{i=1}^{p} \lambda_i \mathbf{e}_i \mathbf{e}_i'$$

where

- $\lambda_1, \dots, \lambda_p$, often ordered from the largest to the smallest, are the eigenvalues of A
- Γ is an orthogonal matrix, i.e., $\Gamma\Gamma^T = \Gamma^T\Gamma = \mathbf{I}$ and the columns of Γ are the eigenvectors of A.
- Λ is the diagonal matrix of the eigenvalues

Covariance Matrices

Eigenvalues and Cov Matrices

Subsection 2

Covariance Matrices

A Covariance Matrix Has to be P.(S.)D.

- A covariance matrix has to be either positive definite (p.d.) or positive semidefinite (p.s.d.)
- What is the definition of p.d. or p.s.d.?
 - We say $\mathbf{A}_{p \times p}$ is p.d. (p.s.d.) is $x^T \mathbf{A} x > 0$ (> 0) for any $x \in \mathbb{R}^p$ and $x \neq 0$
- Why do covariance matrices has to be p.d. or p.s.d? The following page provides a nice explanation:

https://gowrishankar.info/blog/why-covariance-matrix-should-bepositive-semi-definite-tests-using-breast-cancer-dataset/

A Covariance Matrix Has to be P.(S.)D.

 Intuitively, the information of the pairwise covariance/correlation has to be consistent. Can the following matrix be a covariance matrix?

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$$\begin{pmatrix}
1 & 0.9 & 0.9 \\
0.9 & 1 & 0 \\
0.9 & 0 & 1
\end{pmatrix}$$

- The matrix indicates that
 - Variable 1 and 2 are highly correlated
 - Variable 1 and 3 are highly correlated
 - Variable 2 and 3 are not correlated
- The pairwise correlations do not seem to be consistent

A Covariance Matrix Has to be P.(S.)D.

• Examine the following matrix, which involves a parameter ρ

$$A = \begin{pmatrix} 1 & 0.9 & 0.9 \\ 0.9 & 1 & \rho \\ 0.9 & \rho & 1 \end{pmatrix}$$

- What values can ρ take in order for the matrix to be an appropriate covariance matrix?
- This is a linear algebra problem. In order for the matrix to be a covariance matrix, the eigenvalues should be non-negative.
- Recall that the eigenvalues are the roots to

$$0 = \det(\lambda I - A) = |\begin{pmatrix} \lambda - 1 & -0.9 & -0.9 \\ -0.9 & \lambda - 1 & -\rho \\ -0.9 & -\rho & \lambda - 1 \end{pmatrix}|$$

Subsection 3

Examples

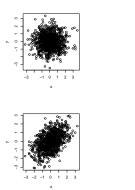
Eigenvalues of Covariance Matrix: Example 1

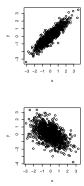
```
p=2; n=1000; rho1=0; rho2=0.9; rho3=0.5; rho4=-0.5
Sigma1=diag(1-rho1, p, p) + matrix(rho1, p, p)
Sigma2=diag(1-rho2, p, p) + matrix(rho2, p, p)
Sigma3=diag(1-rho3, p, p) + matrix(rho3, p, p)
Sigma4=diag(1-rho4, p, p) + matrix(rho4, p, p)
X1=data.frame(mvrnorm(n, rep(0,p), Sigma1)); names(X1)=c("x","y")
X2=data.frame(mvrnorm(n, rep(0,p), Sigma2)); names(X2)=c("x","y")
X3=data.frame(mvrnorm(n, rep(0,p), Sigma3)); names(X3)=c("x","y")
X4=data.frame(mvrnorm(n, rep(0,p), Sigma4)); names(X4)=c("x","y")
```

Eigenvalues of Covariance Matrix: Example 1

Simulated data

```
par(mfrow=c(2,2),pty="s")
plot(X1);plot(X2);plot(X3);plot(X4)
```





Eigenvalues of Covariance Matrix: Example 1

The true covariance matrices.

```
Sigma1
```

```
## [,1] [,2]
## [1,] 1
## [2,] 0
```

Sigma2

```
[,1] [,2]
##
## [1,] 1.0 0.9
## [2,] 0.9 1.0
```

Eigenvalues of Covariance Matrix: Example 1

The true covariance matrices

```
Sigma3
```

```
## [,1] [,2]
## [1,] 1.0 0.5
## [2,] 0.5 1.0
```

Sigma4

```
## [,1] [,2]
## [1,] 1.0 -0.5
## [2,] -0.5 1.0
```

Eigenvalues of Covariance Matrix: Example 1

Eigenvalues of the true covariance matrices

```
eigen(Sigma1)$values
## [1] 1 1
eigen(Sigma2)$values
## [1] 1.9 0.1
eigen(Sigma3)$values
## [1] 1.5 0.5
eigen(Sigma4)$values
## [1] 1.5 0.5
```

Eigenvalues of Covariance Matrix: Example 1

Eigenvalues of the estimated covariance matrices

```
eigen(cov(X1))$values
## [1] 1.031666 1.002280
eigen(cov(X2))$values
## [1] 1.80276014 0.09575692
eigen(cov(X3))$values
## [1] 1.5450471 0.5231463
eigen(cov(X4))$values
## [1] 1.5581405 0.5016192
```

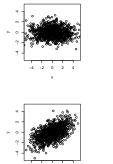
Eigenvalues of Covariance Matrix: Example 2

```
n=1000
Sigma1=diag(c(4,1), 2, 2)
Sigma2=diag(c(1,4), 2, 2)
theta=pi/6
R1=matrix(c(cos(theta), sin(theta), -sin(theta), cos(theta)), 2,2)
theta=pi/4+pi/2
R2=matrix(c(cos(theta), sin(theta), -sin(theta), cos(theta)), 2,2)
Sigma3=R1%*%Sigma1%*%t(R1)
Sigma4=R2%*%Sigma1%*%t(R2)
X1=data.frame(mvrnorm(n, rep(0,2), Sigma1)); names(X1)=c("x","y")
X2=data.frame(mvrnorm(n, rep(0,2), Sigma2)); names(X2)=c("x","y")
X3=data.frame(mvrnorm(n, rep(0,2), Sigma3)); names(X3)=c("x","y")
X4=data.frame(mvrnorm(n, rep(0,2), Sigma4)); names(X4)=c("x","y")
```

Eigenvalues of Covariance Matrix: Example 2

Simulated data

```
par(mfrow=c(2,2),pty="s")
plot(X1, xlim=c(-5,5), ylim=c(-5,5));plot(X2, xlim=c(-5,5), ylim=c(-5,5)
plot(X3, xlim=c(-5,5), ylim=c(-5,5));plot(X4, xlim=c(-5,5), ylim=c(-5,5))
```







Eigenvalues of Covariance Matrix: Example 2

The true covariance matrices

```
Sigma1
```

```
## [,1] [,2]
## [1,] 4 0
## [2,] 0 1
```

Sigma2

```
## [,1] [,2]
## [1,] 1 0
## [2,] 0 4
```

Eigenvalues of Covariance Matrix: Example 2

The true covariance matrices.

```
Sigma3
```

```
## [,1] [,2]
## [1,] 3.250000 1.299038
## [2,] 1.299038 1.750000
```

Sigma4

```
## [,1] [,2]
## [1,] 2.5 -1.5
## [2,] -1.5 2.5
```

Eigenvalues of Covariance Matrix: Example 2

Eigenvalues of the true covariance matrices

```
eigen(Sigma1)$values
## [1] 4 1
eigen(Sigma2)$values
## [1] 4 1
eigen(Sigma3)$values
## [1] 4 1
eigen(Sigma4)$values
## [1] 4 1
```

Eigenvalues of Covariance Matrix: Example 2

Eigenvalues of the estimate covariance matrices

```
eigen(cov(X1))$values
## [1] 4.041183 1.001826
eigen(cov(X2))$values
## [1] 3.950876 1.061222
eigen(cov(X3))$values
## [1] 3.9554303 0.9806424
eigen(cov(X4))$values
## [1] 4.1040168 0.9952649
```

Section 2

One-way ANOVA

One-way ANOVA: Notations and Assumptions

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- g independent samples
 - $Y_{11}, \cdots Y_{1,n_1} \stackrel{iid}{\sim} N(\mu_1, \sigma^2)$
 - $Y_{21}, \cdots Y_{2n_2} \stackrel{iid}{\sim} N(\mu_2, \sigma^2)$

 - $Y_{\sigma 1}, \cdots Y_{\sigma, n_{\sigma}} \stackrel{iid}{\sim} N(\mu_{\sigma}, \sigma^2)$
- Total sample size $n = n_1 + \cdots + n_{\sigma} = \sum_{i=1}^{g} n_i$
- Group means: \bar{Y}_i for $i = 1, \dots, g$
- Grand/overall mean: \overline{Y}

One-Way ANOVA: Partition the Sum of Squares of Total:

$$SSTO = \sum_{i=1}^{g} \sum_{j=1}^{n_{i}} (Y_{ij} - \bar{Y}_{..})^{2} = \sum_{i=1}^{g} \sum_{j=1}^{n_{i}} [(Y_{ij} - \bar{Y}_{i.}) + (\bar{Y}_{i.} - \bar{Y}_{..})]^{2}$$

$$= \sum_{i=1}^{g} \sum_{j=1}^{n_{i}} (Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{i=1}^{g} n_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2} + 2 \sum_{i=1}^{g} \sum_{j=1}^{n_{i}} [(Y_{ij} - \bar{Y}_{i.})(\bar{Y}_{i.} - \bar{Y}_{..})]$$

$$= \sum_{i=1}^{g} \sum_{j=1}^{n_{i}} (Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{i=1}^{g} n_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2} + 2 \sum_{i=1}^{g} [(\bar{Y}_{i.} - \bar{Y}_{..}) \sum_{j=1}^{n_{i}} (Y_{ij} - \bar{Y}_{i.})]$$

$$= \sum_{g} \sum_{i=1}^{g} \sum_{n_{i}} [(Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{i=1}^{g} n_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2} + 2 \sum_{i=1}^{g} [(\bar{Y}_{i.} - \bar{Y}_{..}) \cdot 0]$$

$$= \sum_{i=1}^{g} \sum_{j=1}^{n_{i}} [(Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{i=1}^{g} n_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2}$$

$$= \sum_{i=1}^{g} \sum_{j=1}^{g} [(Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{j=1}^{g} n_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2}$$

$$= \sum_{i=1}^{g} \sum_{j=1}^{g} [(Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{j=1}^{g} n_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2}$$

One-Way ANOVA: E(SSE)

- SSTO = SSE + SSTR = SSW + SSB:
- SSE is also known as SSW, the within-group variance

$$SSE = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_{i.})^2 = \sum_{i=1}^{g} (n_i - 1)s_i^2$$

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where s_i^2 is the sample variance for the *i*th group. Recall that $E[s_i^2] = \sigma^2$. Therefore,

$$E[SSE] = \sum_{i=1}^{g} (n_i - 1)\sigma^2 = (n - g)\sigma^2$$

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One-Way ANOVA: E(SSTR)

- SSTR is also known as SSB, the between-group variance
- The calculation of E(SSTR) requires the following results

$$\begin{split} E[\bar{Y}_{i.}^2] &= Var[\bar{Y}_{i.}] + \mu_i^2 = \frac{\sigma_i^2}{n_i} + \mu_i^2 \\ E[\bar{Y}_{..}^2] &= Var[\bar{Y}_{..}] + \mu_{..}^2 = \frac{\sigma^2}{n} + \mu_{..}^2 \\ E[\bar{Y}_{i.}] \bar{Y}_{..}] &= \frac{1}{n} E[\bar{Y}_{i.} \sum_{j=1}^g n_j \bar{Y}_{j.}] = \frac{1}{n} E[\bar{Y}_{i.} (n_i \bar{Y}_{i.} + \sum_{j \neq i}^g n_j \bar{Y}_{j.})] \\ &= \frac{n_i}{n} E[\bar{Y}_{i.}^2] + \mu_i \sum_{j \neq i} \frac{n_j}{n} \mu_j = \frac{1}{n} \sigma^2 + \frac{n_i}{n} \mu_i^2 + \mu_i \sum_{j \neq i} \frac{n_j}{n} \mu_j \\ &= \frac{1}{n} \sigma^2 + \mu_i \bar{\mu}_{..} \end{split}$$

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One-Way ANOVA: E(SSTR) (continued)

where $\bar{\mu}_{..} = \frac{1}{n} \sum_{i=1}^{g} n_i \mu_i$.

$$E[SSTR] = \sum_{i=1}^{g} n_{i} E[(\bar{Y}_{i.} - \bar{Y}_{..})^{2}]$$

$$= \sum_{i=1}^{g} n_{i} E[\bar{Y}_{i.}^{2} + \bar{Y}_{..}^{2} - 2\bar{Y}_{i.}\bar{Y}_{..}]$$

$$= \sum_{i=1}^{g} n_{i} E[\bar{Y}_{i.}^{2} + \bar{Y}_{..}^{2} - 2\bar{Y}_{i.}\bar{Y}_{..}]$$

$$= \sum_{i=1}^{g} n_{i} (\frac{\sigma^{2}}{n_{i}} + \mu_{i}^{2} + \frac{\sigma^{2}}{n} + \mu_{..}^{2} - 2[\frac{1}{n}\sigma^{2} + \mu_{i}\bar{\mu}_{..}])$$

$$= (g - 1)\sigma^{2} + \sum_{i=1}^{g} n_{i} (\mu_{i} - \bar{u}_{..})^{2}$$

One-Way ANOVA: Mean of Sum of Squares

The null hypothesis

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

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The alternative hypothesis

$$H_a$$
: $\mu_i \neq \mu_j$ for at least one pair of (i,j)

- Mean sum of squares: $MSE = \frac{1}{n-\sigma}SSE$, $MSTR = \frac{1}{\sigma-1}SSTR$
- $E(MSE) = \sigma^2$
- $E(MSTR) = \sigma^2 + \frac{1}{g-1} \sum_{i=1}^{g} n_i (\mu_i \bar{u}_{..})^2 \stackrel{H_0}{=} \sigma^2$
- Thus, a reasonable statistic is the ratio of the two mean sum of squares

One-Way ANOVA: F-statistic

The F-statistic is defined as

$$F = \frac{MSTR}{MSE}$$

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The null distribution

$$F = \frac{\frac{SSTR}{\sigma^2}/(g-1)}{\frac{SSE}{\sigma^2}/(n-g)} \stackrel{H_0}{\sim} F_{g-1,n-g}$$

- To derive the null distribution of F, we need to show that
 - $\frac{SSTR}{\sigma^2} \stackrel{H_0}{\sim} \chi_{g-1}^2$
 - $\frac{SSE}{\sigma^2} \sim \chi^2_{n-\sigma}$
 - SSTR and SSE are independent

ANOVA Table and Distributions

Source	SS	MS	DF	F
Treatment	$SSTR = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (\bar{Y}_{i.} - \bar{Y}_{})^2$	$MSTR = \frac{SSTR}{g-1}$	g-1	$F = \frac{MSTR}{MSE}$
Error	$SSE = \sum_{i=1}^{g} \sum_{j=1}^{\hat{n_i}} (Y_{ij} - \bar{Y}_{i.})^2$	$MSTR = \frac{SSTR}{g-1}$ $MSE = \frac{SSE}{n-g}$	n-g	
Total	$SSTO = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_{})^2$			

Section 3

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ANOVA to MANOVA

- So far we have learned
 - One-sample Hotelling's T²
 - Two-sample Hotelling's T^2
- The next logical extension is to multiple samples, i.e., the multivariate version anova, or MANOVA
- Compared to one-sample or two-sample multivariate analysis, there are many choices for comparing multiple samples for multivariate data

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• We will cover the following methods:

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Eigenvalues and Cov Matrices

Notations and Assumptions

g independent random samples

•
$$\mathbf{Y}_{11}, \cdots \mathbf{Y}_{1,n_1} \overset{iid}{\sim} \mathcal{N}(\boldsymbol{\mu}_1, \boldsymbol{\Sigma})$$

$$\bullet$$
 $\mathbf{Y}_{21}, \cdots \mathbf{Y}_{2,n_2} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{\mu}_2, \boldsymbol{\Sigma})$

$$ullet$$
 $\mathbf{Y}_{g1}, \cdots \mathbf{Y}_{g,n_g} \stackrel{iid}{\sim} \mathcal{N}(\mu_g, \mathbf{\Sigma})$

- n_i : the number of observations in group i
- The ith random sample is from $N(\mu_i, \Sigma)$

Notations and Assumptions (continued)

• Each $\mathbf{Y}_{ij} \in \mathbb{R}^p$, i.e,

$$\left(\begin{array}{c} Y_{ij1} \\ Y_{ij2} \\ \vdots \\ Y_{ijp} \end{array}\right)$$

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The null hypothesis

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

The alternative hypothesis

$$H_1$$
: $\mu_i \neq \mu_j$ for at least one pair of (i,j)

The Total Covariance Matrix

 The total covariance matrix is the covariance matrix if the group information is ignored. If we pool all the $n = n_1 + \cdots + n_g = \sum_{i=1}^g n_g$ observations together, what is the covariance matrix?

$$\mathbf{T} = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{..}) (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{..})'$$

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

$$= \sum_{i=1}^{g} \sum_{n_i} (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{i.}) (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{i.})' + \sum_{j=1}^{g} n_i (\bar{\mathbf{Y}}_{i.} - \bar{\mathbf{Y}}_{..}) (\bar{\mathbf{Y}}_{i.} - \bar{\mathbf{Y}}_{..})' + 0$$

The Within-Group Sample Covariance Matrix

• In the definition of the total covariance matrix, we compare each observation to the grand mean vector

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 In within-group covariance matrix, we compare each observation to its group mean

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

The Within-Group Sample Covariance Matrix (continued)

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• Let $\mathbf{W}_i = \sum_{i=1}^{n_i} (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{i.}) (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{i.})'$. We can show that

$$\mathbf{W}_1, \cdots, \mathbf{W}_g \overset{independent}{\sim} \mathit{Wishart}_p(\mathit{n}_i - 1, \mathbf{\Sigma})$$

 Recall that the sum of independently distributed chi-squared distributed random variables also follows a chi-squared distribution. Similarly,

$$\textbf{W} = \sum_{i=1}^{g} \textbf{W}_{i} \sim \textbf{Wishart}_{p}(\textbf{n} - \textbf{g}, \boldsymbol{\Sigma})$$

The Between-Group Sample Covariance Matrix

 The between-group sample covariance matrix captures the difference in mean vectors between groups

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$$\mathbf{B} = \sum_{i=1}^{g} n_i (\mathbf{\bar{Y}}_{i.} - \mathbf{\bar{Y}}_{..}) (\mathbf{\bar{Y}}_{i.} - \mathbf{\bar{Y}}_{..})'$$

 When the null hypothesis is true, B follows a Wishart distribution

$$\mathbf{B} \stackrel{H_0}{\sim} Wishart_p(g-1, \mathbf{\Sigma})$$

Outline of Proof

- Let $\mathbf{Y}_{n \times p}$ denote the data matrix
- Let $X_{n \times g}$ denote the design matrix, which consists of dummy variables of the group membership. We also define $P_{n \times n} = X(X^TX)^{-1}X^T$. Then

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$$\mathbf{X} = \begin{pmatrix} \mathbf{1}_{n_1} & \mathbf{0}_{n_1} & \cdots & \mathbf{0}_{n_1} \\ \mathbf{0}_{n_2} & \mathbf{1}_{n_2} & \cdots & \mathbf{0}_{n_2} \\ \cdots & \cdots & \cdots & \cdots \\ \mathbf{0}_{n_g} & \mathbf{0}_{n_g} & \cdots & \mathbf{1}_{n_g} \end{pmatrix}, \mathbf{P} = \begin{pmatrix} \frac{1}{n_1} \mathbf{1}_{n_1} \mathbf{1}_{n_1}^T & \mathbf{0} & \cdots & \mathbf{0} \\ \mathbf{0} & \frac{1}{n_2} \mathbf{1}_{n_2} \mathbf{1}_{n_2}^T & \cdots & \mathbf{0} \\ \cdots & \cdots & \cdots & \cdots \\ \mathbf{0} & \mathbf{0} & \cdots & \frac{1}{n_g} \mathbf{1}_{n_g} \mathbf{1}_{n_g}^T \end{pmatrix}$$

Outline of Proof (continued)

It can be shown that

$$\mathbf{W} = \mathbf{Y}^T (\mathbf{I} - \mathbf{P}) \mathbf{Y}$$
, $\mathbf{B} = \mathbf{Y}^T (\mathbf{P} - \frac{1}{n} \mathbf{1}_n \mathbf{1}_n^T) \mathbf{Y}$

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• It is not difficult to verify that P, I - P, and $P - \frac{1}{n} \mathbf{1}_n \mathbf{1}_n^T$ are all projection matrices. For a projection matrix, its rank equals its trace. Thus, it is not difficult to show that

$$rank(\mathbf{P}) = g, rank(\mathbf{I} - \mathbf{P}) = n - g, rank(\mathbf{P} - \frac{1}{n}\mathbf{1}_n\mathbf{1}_n^T) = g - 1$$

• In Lecture 06 we showed that $(n-1)\mathbf{S} \sim Wishart_p(n-1, \mathbf{\Sigma})$. Use similar methods, we can show that

$$oldsymbol{\mathsf{W}} \sim \mathit{Wishart}_p(\mathsf{n}-\mathsf{g},oldsymbol{\Sigma})$$
 , $oldsymbol{\mathsf{B}} \stackrel{H_0}{\sim} \mathit{Wishart}_p(\mathsf{g}-1,oldsymbol{\Sigma})$ $oldsymbol{\mathsf{W}} \perp oldsymbol{\mathsf{B}}$

MANOVA Table

• We can also construct a table

Source	Sample Cov	DF
Treatment	$\mathbf{B} = \sum_{i=1}^{g} n_i (\mathbf{\bar{Y}}_{i.} - \mathbf{\bar{Y}}_{}) (\mathbf{\bar{Y}}_{i.} - \mathbf{\bar{Y}}_{})'$	g-1
Error	$\mathbf{W} = \sum_{i=1}^{\overline{g}} \sum_{j=1}^{n_i} (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{i.}) (\mathbf{Y}_{ij} - \bar{\mathbf{Y}}_{i.})'$	n-g
Total	$\mathbf{T} = \sum_{i=1}^g \sum_{j=1}^{ ilde{n}_i} (\mathbf{Y}_{ij} - \mathbf{ar{Y}}_{\cdot\cdot}) (\mathbf{Y}_{ij} - \mathbf{ar{Y}}_{\cdot\cdot})'$	

Section 4

MANOVA Test Statistics

MANOVA Test Statistics

• In one-way ANOVA, we understand that SSB should be large relative to SSW when the null hypothesis is not true

- In the multivariate version, we also expect that **B** should be large relative to **W** when the null hypothesis is not true
- However, B and W are matrices. How to define "large"?

Iris Data: B and W

```
#rearrange the data such as the response matrix is
\#an \ n-by-p \ matrix
Y=cbind(SepalL=c(iris3[,1,1],iris3[,1,2],iris3[,1,3]),
SepalW=c(iris3[,2,1],iris3[,2,2],iris3[,2,3]),
PetalL=c(iris3[,3,1],iris3[,3,2],iris3[,3,3]),
PetalW=c(iris3[,4,1],iris3[,4,2],iris3[,4,3]))
#for unknown reasons, data.frame won't work but chind works
#alternatively, we can use the following way to define y
#Y=aperm(iris3, c(1,3,2)); dim(y)=c(150,4)
#define the covariate variable X, which is vector of labels
iris.type=rep(c("Setosa","Versicolor","Virginica"),each=50)
T=(150-1)*cov(Y)
W=(50-1)*cov(iris3[,,1]) + (50-1)*cov(iris3[,,2]) + (50-1)*cov(iris3[,,3])
B=T-W
```

Iris Data: B and W

В

```
## SepalL SepalW PetalL PetalW
## SepalL 63.21213 -19.95267 165.2484 71.27933
## SepalW -19.95267 11.34493 -57.2396 -22.93267
## PetalL 165.24840 -57.23960 437.1028 186.77400
## PetalW 71.27933 -22.93267 186.7740 80.41333
```

W

```
## Sepal L. Sepal W. Petal L. Petal W. ## Sepal L. 38.9562 13.6300 24.6246 5.6450 ## Sepal W. 13.6300 16.9620 8.1208 4.8084 ## Petal L. 24.6246 8.1208 27.2226 6.2718 ## Petal W. 5.6450 4.8084 6.2718 6.1566
```

MANOVA in R

```
summary.manova {stats} R Documentation
Summary Method for Multivariate Analysis of Variance
Description
A summary method for class "manova".
Usage
## S3 method for class 'manova'
summary(object,
        test = c("Pillai", "Wilks", "Hotelling-Lawley", "Roy"),
        intercept = FALSE, tol = 1e-7, ...)
```

MANOVA - Method 1: Pallai Trace

$$V = trace(B(B+W)^{-1}) = trace(BT^{-1})$$

MANOVA - Method 2: Wilk's Lambda

• Wilk's Lambda distribution Let $\mathbf{A} \sim Wishart_p(m_1, \mathbf{I})$ and $\mathbf{B} \sim Wishart_p(m_2, \mathbf{I})$ be independent with $m_1 > p$. We say

$$\Lambda = \frac{|\mathbf{A}|}{|\mathbf{A} + \mathbf{B}|} \sim \Lambda(p, m_1, m_2)$$

MANOVA Sample Cov

Test Statistic

$$\Lambda^* = \frac{|\mathbf{W}|}{|\mathbf{B} + \mathbf{W}|} = \frac{|\mathbf{W}|}{|\mathbf{T}|} = \frac{|\mathbf{\Sigma}^{-1/2} \mathbf{W} \mathbf{\Sigma}^{-1/2}|}{|\mathbf{\Sigma}^{-1/2} (\mathbf{B} + \mathbf{W}) \mathbf{\Sigma}^{-1/2}|}$$

By the definition of Wilk's Lambda distribution,

$$\Lambda^* \stackrel{H_0}{\sim} \Lambda(p, n-g, g-1)$$

MANOVA - Method 3: Lawley-Hotelling Trace

$$T_0^2 = trace(\mathbf{BW}^{-1})$$

MANOVA - Method 4: Roy's Largest Root

Two equivalent test statistics have been used

$$\lambda_{ extit{max}}(\mathsf{BW}^{-1}) \ \lambda_{ extit{max}}(\mathsf{B}(\mathsf{B}+\mathsf{W})^{-1})$$

Test Statistics and the Eigenvalues of BW^{-1}

 One interesting observation is that all the test statistics can be expressed in terms of eigenvalues of BW^{-1} . Let $\lambda_1, \dots, \lambda_p$ denote the eigenvalues, from the largest to the smallest. We have

MANOVA Sample Cov

Pillai trace

$$tr(\mathbf{B}(\mathbf{B}+\mathbf{W})^{-1}) = \sum_{i=1}^{min(p,g-1)} \frac{\lambda_i}{1+\lambda_i}$$

Wilk's Lambda

$$\Lambda^* = \prod_{i=1}^{min(p,g-1)} rac{1}{1+\lambda_i}$$

Test Statistics and the Eigenvalues of ${\sf BW}^{-1}$

Lawley-Hotelling trace

$$\mathit{trace}(\mathsf{BW}^{-1}) = \sum_{i=1}^{\mathit{min}(p,g-1)} \lambda_i$$

MANOVA Sample Cov

Roy's largest root

$$\lambda_{max}(\mathbf{B}\mathbf{W}^{-1}) = \lambda_1 \ \lambda_{max}(\mathbf{B}(\mathbf{B} + \mathbf{W})^{-1}) = rac{\lambda_1}{1 + \lambda_1}$$

Proof

 If you are curious about how to prove these results, I provide an example.

MANOVA Sample Cov

$$(\Lambda^*)^{-1} = rac{|\mathbf{B} + \mathbf{W}|}{|\mathbf{W}|} = |\mathbf{I} + \mathbf{W}^{-1}\mathbf{B}|$$

$$= \prod_{i=1}^p [\text{the ith eigenvalue of } \mathbf{I} + \mathbf{W}^{-1}\mathbf{B}]$$

$$= \prod_{i=1}^p [1 + \text{the ith eigenvalue of } \mathbf{W}^{-1}\mathbf{B}]$$

$$= \prod_{i=1}^p (1 + \lambda_i) \stackrel{1}{=} \prod_{i=1}^{min(p,g-1)} (1 + \lambda_i)$$

As a result, $\Lambda^* = \prod_{i=1}^p \frac{1}{1+\lambda_i}$

Proof (continued)

- The last step implies that $rank(\mathbf{W}^{-1}\mathbf{B}) = min(p, g 1)$
- Why is it true?
 - First, $rank(\mathbf{W}^{-1}\mathbf{B}) = rank(\mathbf{B})$
 - Second, because $\mathbf{B} \stackrel{H_0}{\sim} Wishart_n(g-1, \mathbf{\Sigma})$, there exists $\mathbf{Y}_{(\varrho-1)\times\varrho}$ such that

$$\mathbf{B} = \mathbf{Y}^T \mathbf{Y}$$
 ,

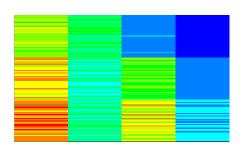
MANOVA Sample Cov

and **Y** is a random sample of size g-1 from $N(\mathbf{0}, \mathbf{\Sigma})$. The rank of **Y** is $rank(\mathbf{Y}) = min(p, g - 1)$.

• Third, $rank(\mathbf{B}) = rank(\mathbf{Y}^T\mathbf{Y}) = rank(\mathbf{Y}) = min(p, g-1)$.

Example - Iris Data

```
mycolors=rainbow(12)[9:1]
image(t(iris[150:1, 1:4]),col = mycolors, xaxt="n", yaxt="n")
```



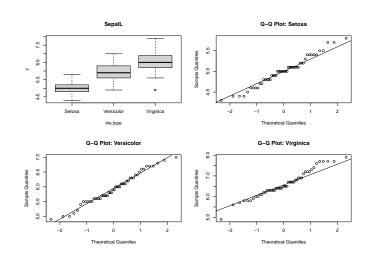
Iris Data: Univariate One-way ANOVA: Data Formatting

```
#rearrange the data in the (X,Y) format
y=c(iris3[,1,1], iris3[,1,2], iris3[,1,3])
#alternatively, you may use:
#y=aperm(iris3[,1,], c(1,2)); dim(y)=c(150,1)
#define the covariate variable X,
#which is vector of labels
iris.type=rep(c("Setosa", "Versicolor", "Virginica"), each=50)
```

Iris Data: Univariate One-way ANOVA: Exploratory

```
#visual checking
par(mfrow=c(2,2))
#box plot
boxplot(y~iris.type, main="SepalL")
#alternatively, you may use: boxplot(iris3[,1,],
#main="SepalL")
#qq plots
qqnorm(iris3[,1,1], main="Q-Q Plot: Setosa");
qqline(iris3[,1,1])
qqnorm(iris3[,1,2], main="Q-Q Plot: Versicolor");
qqline(iris3[,1,2])
qqnorm(iris3[,1,3], main="Q-Q Plot: Virginica");
qqline(iris3[,1,3])
```

Iris Data: Univariate One-way ANOVA: Exploratory



Iris Data: Univariate One-way ANOVA: Analysis

```
obj.aov=aov(y~iris.type)
summary(obj.aov)
```

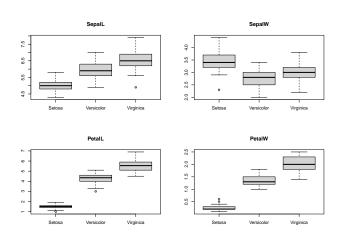
Iris Data: MANOVA: Data Formatting

```
#rearrange the data such as the response matrix is
\#an \ n-by-p \ matrix
Y=cbind(SepalL=c(iris3[,1,1],iris3[,1,2],iris3[,1,3]),
SepalW=c(iris3[,2,1],iris3[,2,2],iris3[,2,3]),
PetalL=c(iris3[,3,1],iris3[,3,2],iris3[,3,3]),
PetalW=c(iris3[,4,1],iris3[,4,2],iris3[,4,3]))
#for unknown reasons, data.frame won't work but chind works
#alternatively, we can use the following way to define y
#Y=aperm(iris3, c(1,3,2)); dim(y)=c(150,4)
#define the covariate variable X, which is vector of labels
iris.type=rep(c("Setosa", "Versicolor", "Virginica"), each=50)
```

Iris Data: MANOVA: Exploratory

```
#visual investigation
par(mfrow=c(2,2))
boxplot(iris3[,1,],main="SepalL")
boxplot(iris3[,2,],main="SepalW")
boxplot(iris3[,3,],main="PetalL")
boxplot(iris3[,4,],main="PetalW")
```

Iris Data: MANOVA: Exploratory



[1] O

Conducting MANOVA "Manually"

```
T=(150-1)*cov(Y)
W=(50-1)*cov(iris3[,,1]) + (50-1)*cov(iris3[,,2])+(50-1)*cov(iris3[,,3])
B=T-W
Lambda=prod(1/(1+ eigen(B%*%solve(W))$values))
(150-3-2)/3*(1-sqrt(Lambda))/sqrt(Lambda)
## [1] 267.3711
# Using relationship between Wilk's lambda and F-distribution
# (see wikipedia about "Wilks's lambda distribution")
1-pf((150-3-2)/3*(1-sqrt(Lambda))/sqrt(Lambda), 2*3, 150-3-2)
```

Conducting MANOVA using "manova" in R

```
obj=manova(Y~iris.type)
obj.aov
## Call:
      aov(formula = y ~ iris.type)
##
##
## Terms:
##
                   iris.type Residuals
                   63.21213 38.95620
## Sum of Squares
                            2
  Deg. of Freedom
                                    147
##
## Residual standard error: 0.5147894
## Estimated effects may be unbalanced
```

Conducting MANOVA using "manova" in R

```
summary(obj, test="Pillai")
##
             Df Pillai approx F num Df den Df Pr(>F)
## iris.type 2 1.1919 53.466 8
                                        290 < 2.2e-16 ***
## Residuals 147
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
summary(obj, test="Wilks")
##
             Df
                  Wilks approx F num Df den Df Pr(>F)
## iris.type 2 0.023439 199.15 8
                                          288 < 2.2e-16 ***
## Residuals 147
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Conducting MANOVA using "manova" in R

```
summary(obj, test="Hotelling-Lawley")
             Df Hotelling-Lawley approx F num Df den Df Pr(>F)
##
                         32.477 580.53
## iris.type
                                             8
                                                  286 < 2.2e-16 ***
## Residuals 147
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
summary(obj, test="Roy")
##
             Df
                  Roy approx F num Df den Df Pr(>F)
## iris.type 2 32.192
                          1167 4 145 < 2.2e-16 ***
## Residuals 147
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Too Many Choices?

• Due to the nature of multivariate analysis, we have seen many choices for conducting one-way MANOVA

- Do they work equally well?
- Does their performance depend on the true distribution?
- You will be asked to compare the methods in your midterm project