

# Canonical Correlation Analysis

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STAT 4690—Applied Multivariate Analysis

# Introduction

- Canonical Correlation Analysis (CCA) is a dimension reduction method that is similar to PCA, but where we simultaneously reduce the dimension of **two** random vectors  $\mathbf{Y}$  and  $\mathbf{X}$ .
- Instead of trying to explain overall variance, we try to explain the covariance  $\text{Cov}(\mathbf{Y}, \mathbf{X})$ .
  - Note that this is a measure of **association** between  $\mathbf{Y}$  and  $\mathbf{X}$ .
- Examples include:
  - Arithmetic speed and power ( $\mathbf{Y}$ ) and reading speed and power ( $\mathbf{X}$ )
  - College performance metrics ( $\mathbf{Y}$ ) and high-school achievement metrics ( $\mathbf{X}$ )

# Population model i

- Let  $\mathbf{Y}$  and  $\mathbf{X}$  be  $p$ - and  $q$ -dimensional random vectors, respectively.
  - We will assume that  $p \leq q$ .
- Let  $\mu_Y$  and  $\mu_X$  be the mean of  $\mathbf{Y}$  and  $\mathbf{X}$ , respectively.
- Let  $\Sigma_Y$  and  $\Sigma_X$  be the covariance matrix of  $\mathbf{Y}$  and  $\mathbf{X}$ , respectively, and let  $\Sigma_{YX} = \Sigma_{XY}^T$  be the covariance matrix  $\text{Cov}(\mathbf{Y}, \mathbf{X})$ .
  - Assume  $\Sigma_Y$  and  $\Sigma_X$  are positive definite.
- Note that  $\Sigma_{YX}$  has  $pq$  entries, corresponding to all covariances between a component of  $\mathbf{Y}$  and a component of  $\mathbf{X}$ .

- **Goal of CCA:** Summarise  $\Sigma_{YX}$  with  $p$  numbers.
  - These  $p$  numbers will be called the *canonical correlations*.

# Dimension reduction i

- Let  $U = a^T \mathbf{Y}$  and  $V = b^T \mathbf{Y}$  be linear combinations of  $\mathbf{Y}$  and  $\mathbf{X}$ , respectively.
- We have:
  - $\text{Var}(U) = a^T \Sigma_Y a$
  - $\text{Var}(V) = b^T \Sigma_X b$
  - $\text{Cov}(U, V) = a^T \Sigma_{YX} b$ .
- Therefore, we can write the correlation between  $U$  and  $V$  as follows:

$$\text{Corr}(U, V) = \frac{a^T \Sigma_{YX} b}{\sqrt{a^T \Sigma_Y a} \sqrt{b^T \Sigma_X b}}.$$

## Dimension reduction ii

- We are looking for vectors  $a \in \mathbb{R}^p, b \in \mathbb{R}^q$  such that  $\text{Corr}(U, V)$  is **maximised**.

# Definitions

- The *first pair of canonical variates* is the pair of linear combinations  $U_1, V_1$  with unit variance such that  $\text{Corr}(U_1, V_1)$  is maximised.
- The  **$k$ -th pair of canonical variates** is the pair of linear combinations  $U_k, V_k$  with unit variance such that  $\text{Corr}(U_k, V_k)$  is maximised among all pairs that are uncorrelated with the previous  $k - 1$  pairs.
- When  $U_k, V_k$  is the  $k$ -th pair of canonical variates, we say that  $\rho_k = \text{Corr}(U_k, V_k)$  is the  $k$ -th *canonical correlation*.

# Derivation of canonical variates i

- Make a change of variables:
  - $\tilde{a} = \Sigma_Y^{1/2} a$
  - $\tilde{b} = \Sigma_X^{1/2} b$
- We can then rewrite the correlation:

$$\begin{aligned}\text{Corr}(U, V) &= \frac{a^T \Sigma_{YX} b}{\sqrt{a^T \Sigma_Y a} \sqrt{b^T \Sigma_X b}} \\ &= \frac{\tilde{a}^T \Sigma_Y^{-1/2} \Sigma_{YX} \Sigma_X^{-1/2} \tilde{b}}{\sqrt{\tilde{a}^T \tilde{a}} \sqrt{\tilde{b}^T \tilde{b}}}.\end{aligned}$$



## Derivation of canonical variates ii

- Let  $M = \Sigma_Y^{-1/2} \Sigma_{YX} \Sigma_X^{-1/2}$ . We have

$$\max_{a,b} \text{Corr}(a^T \mathbf{Y}, b^T \mathbf{Y}) \iff \max_{\tilde{a}, \tilde{b}: \|\tilde{a}\|=1, \|\tilde{b}\|=1} \tilde{a}^T M \tilde{b}$$

- The solution to this maximisation problem involves the **singular value decomposition** of  $M$ .
- Equivalently, it involves the **eigendecomposition** of  $MM^T$ , where

$$MM^T = \Sigma_Y^{-1/2} \Sigma_{YX} \Sigma_X^{-1} \Sigma_{XY} \Sigma_Y^{-1/2}.$$

## CCA: Main theorem i

- Let  $\lambda_1 \geq \dots \geq \lambda_p$  be the eigenvalues of  $\Sigma_Y^{-1/2} \Sigma_{YX} \Sigma_X^{-1} \Sigma_{XY} \Sigma_Y^{-1/2}$ .
  - Let  $e_1, \dots, e_p$  be the corresponding eigenvector with unit norm.
- Note that  $\lambda_1 \geq \dots \geq \lambda_p$  are also the  $p$  largest eigenvalues of

$$M^T M = \Sigma_X^{-1/2} \Sigma_{XY} \Sigma_Y^{-1} \Sigma_{YX} \Sigma_X^{-1/2}.$$

- Let  $f_1, \dots, f_p$  be the corresponding eigenvectors with unit norm.

## CCA: Main theorem ii

- Then the  $k$ -th pair of canonical variates is given by

$$U_k = e_k^T \Sigma_Y^{-1/2} \mathbf{Y}, \quad V_k = f_k^T \Sigma_X^{-1/2} \mathbf{X}.$$

- Moreover, we have

$$\rho_k = \text{Corr}(U_k, V_k) = \sqrt{\lambda_k}.$$

## Some vocabulary

1. **Canonical directions:**  $(e_k^T \Sigma_Y^{-1/2}, f_k^T \Sigma_X^{-1/2})$
2. **Canonical variates:**  $(U_k, V_k) = (e_k^T \Sigma_Y^{-1/2} \mathbf{Y}, f_k^T \Sigma_X^{-1/2} \mathbf{X})$
3. **Canonical correlations:**  $\rho_k = \sqrt{\lambda_k}$

## Example i

```
Sigma_Y <- matrix(c(1, 0.4, 0.4, 1), ncol = 2)
Sigma_X <- matrix(c(1, 0.2, 0.2, 1), ncol = 2)
Sigma_YX <- matrix(c(0.5, 0.3, 0.6, 0.4), ncol = 2)
Sigma_XY <- t(Sigma_YX)

rbind(cbind(Sigma_Y, Sigma_YX),
      cbind(Sigma_XY, Sigma_X))
```

## Example ii

##		[,1]	[,2]	[,3]	[,4]
##	[1,]	1.0	0.4	0.5	0.6
##	[2,]	0.4	1.0	0.3	0.4
##	[3,]	0.5	0.3	1.0	0.2
##	[4,]	0.6	0.4	0.2	1.0

## Example iii

```
library(expm)
sqrt_Y <- sqrtm(Sigma_Y)
sqrt_X <- sqrtm(Sigma_X)
M1 <- solve(sqrt_Y) %*% Sigma_YX %*% solve(Sigma_X)%*%
  Sigma_XY %*% solve(sqrt_Y)

(decomp1 <- eigen(M1))
```

## Example iv

```
## eigen() decomposition
## $values
## [1] 0.5457180317 0.0009089525
##
## $vectors
##           [,1]      [,2]
## [1,] -0.8946536  0.4467605
## [2,] -0.4467605 -0.8946536

decomp1$vectors[,1] %*% solve(sqrt_Y)
```



## Example v

```
##           [,1]      [,2]
## [1,] -0.8559647 -0.2777371

M2 <- solve(sqrt_X) %*% Sigma_XY %*% solve(Sigma_Y)%*%
      Sigma_YX %*% solve(sqrt_X)

decomp2 <- eigen(M2)
decomp2$vectors[,1] %*% solve(sqrt_X)

##           [,1]      [,2]
## [1,] 0.5448119 0.7366455
```

## Example vi

```
sqrt(decomp1$values)
```

```
## [1] 0.73872731 0.03014884
```

# Sample CCA

- Let  $\mathbf{Y}_1, \dots, \mathbf{Y}_n$  and  $\mathbf{X}_1, \dots, \mathbf{X}_n$  be random samples, and arrange them in  $n \times p$  and  $n \times q$  matrices  $\mathbb{Y}, \mathbb{X}$ , respectively.
  - Note that both sample sizes are equal.
  - Indeed, we assume that  $(\mathbf{Y}_i, \mathbf{X}_i)$  are sampled jointly, i.e. on the **same** experimental unit.
- Let  $\bar{\mathbf{Y}}$  and  $\bar{\mathbf{X}}$  be the sample means.
- Let  $S_Y$  and  $S_X$  be the sample covariances.
- Define

$$S_{YX} = \frac{1}{n-1} \sum_{i=1}^n (\mathbf{Y}_i - \bar{\mathbf{Y}}) (\mathbf{X}_i - \bar{\mathbf{X}})^T.$$

## Sample CCA: Main theorem i

- Let  $\hat{\lambda}_1 \geq \dots \geq \hat{\lambda}_p$  be the eigenvalues of  $S_Y^{-1/2} S_{YX} S_X^{-1} S_{XY} S_Y^{-1/2}$ .
  - Let  $\hat{e}_1, \dots, \hat{e}_p$  be the corresponding eigenvector with unit norm.
- Note that  $\hat{\lambda}_1 \geq \dots \geq \hat{\lambda}_p$  are also the  $p$  largest eigenvalues of

$$S_X^{-1/2} S_{XY} S_Y^{-1} S_{YX} S_X^{-1/2}.$$

- Let  $\hat{f}_1, \dots, \hat{f}_p$  be the corresponding eigenvectors with unit norm.

## Sample CCA: Main theorem ii

- Then the  $k$ -th pair of *sample* canonical variates is given by

$$\hat{U}_k = \mathbb{Y} S_Y^{-1/2} \hat{e}_k, \quad \hat{V}_k = \mathbb{X} S_X^{-1/2} \hat{f}_k.$$

- Moreover, we have that  $\hat{\rho}_k = \sqrt{\hat{\lambda}_k}$  is the sample correlation of  $\hat{U}_k$  and  $\hat{V}_k$ .

## Example (cont'd) i

*# Let's generate data*

```
library(mvtnorm)
```

```
Sigma <- rbind(cbind(Sigma_Y, Sigma_YX),  
               cbind(Sigma_XY, Sigma_X))
```

```
YX <- rmvnorm(100, sigma = Sigma)
```

```
Y <- YX[,1:2]
```

```
X <- YX[,3:4]
```

```
decomp <- cancortest(x = X, y = Y)
```

## Example (cont'd) ii

```
U <- Y %*% decomp$ycoef
```

```
V <- X %*% decomp$xcoef
```

```
diag(cor(U, V))
```

```
## [1] 0.66401984 0.02024187
```

```
decomp$cor
```

```
## [1] 0.66401984 0.02024187
```

## Example i

```
library(tidyverse)
library(dslabs)

X <- olive %>%
  select(-area, -region) %>%
  as.matrix

Y <- olive %>%
  select(region) %>%
  model.matrix(~ region - 1, data = .)
```



## Example ii

```
head(unname(Y))
```

##		[,1]	[,2]	[,3]
##	[1,]	0	0	1
##	[2,]	0	0	1
##	[3,]	0	0	1
##	[4,]	0	0	1
##	[5,]	0	0	1
##	[6,]	0	0	1

## Example iii

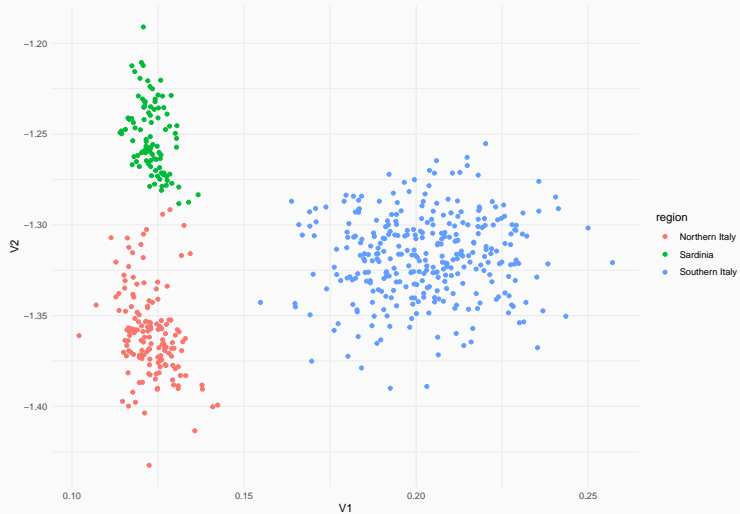
```
decomp <- cancel(X, Y)
```

```
V <- X %*% decomp$xccoef
```

## Example iv

```
data.frame(  
  V1 = V[,1],  
  V2 = V[,2],  
  region = olive$region  
) %>%  
  ggplot(aes(V1, V2, colour = region)) +  
  geom_point() +  
  theme_minimal()
```

# Example v



- The main difference between CCA and Multivariate Linear Regression is that CCA treats  $\mathbb{Y}$  and  $\mathbb{X}$  *symmetrically*.
- As with PCA, you can use CCA and the covariance matrix or the correlation matrix.
  - The latter is equivalent to performing CCA on the standardised variables.
- Note that sample CCA involves inverting the sample covariance matrices  $S_Y$  and  $S_X$ :
  - This means we need to assume  $p, q < n$ .
  - In general, this is what drives most of the performance (or lack thereof) of CCA.

- There may be gains in efficiency by directly estimating the inverse covariance.
- When one of the two datasets  $\mathbb{Y}$  or  $\mathbb{X}$  represent indicators variables for a categorical variables (cf. the olive dataset), CCA is equivalent to **Linear Discriminant Analysis**.
  - To learn more about this method, see a course/textbook on Statistical Learning.

# Proportions of Explained Sample Variance i

- Just like in PCA, there is a notion of *proportion of explained variance* that may be helpful in determining the number of canonical variates to retain.
- Assume that  $\mathbf{Y}_1, \dots, \mathbf{Y}_n$  and  $\mathbf{X}_1, \dots, \mathbf{X}_n$  have been **standardized**. The matrices  $A$  and  $B$  of canonical directions have the following properties:
  - The **rows** are the canonical directions (by definition!)
  - The **columns** of the inverses  $A^{-1}, B^{-1}$  are the sample correlations between the canonical variates and the standardized variables.

## Proportions of Explained Sample Variance ii

- Moreover, we have
  - $\text{Corr}(\mathbb{Y}) = A^{-1}A^{-T}$
  - $\text{Corr}(\mathbb{X}) = B^{-1}B^{-T}$
- But recall that
  - $\text{tr}(\text{Corr}(\mathbb{Y})) = p$
  - $\text{tr}(\text{Corr}(\mathbb{X})) = q$



## Proportions of Explained Sample Variance iii

- Putting this all together, we have that
  - Proportion of total standardized sample variance in  $\mathbb{Y} = (\mathbb{Y}_1 \ \cdots \ \mathbb{Y}_p)$  explained by  $\hat{U}_1, \dots, \hat{U}_r$ :

$$R^2(\mathbf{Y} \mid \hat{U}_1, \dots, \hat{U}_r) = \frac{\sum_{i=1}^r \sum_{j=1}^p \text{Corr}(\hat{U}_i, \mathbb{Y}_k)^2}{p}$$

- Proportion of total standardized sample variance in  $\mathbb{X} = (\mathbb{X}_1 \ \cdots \ \mathbb{X}_q)$  explained by  $\hat{V}_1, \dots, \hat{V}_r$ :

$$R^2(\mathbf{X} \mid \hat{V}_1, \dots, \hat{V}_r) = \frac{\sum_{i=1}^r \sum_{j=1}^q \text{Corr}(\hat{V}_i, \mathbb{X}_k)^2}{q}$$

## Example i

```
# Olive data
X_sc <- scale(X)
Y_sc <- scale(Y)
decomp_sc <- cancort(X_sc, Y_sc)

V_sc <- X_sc %*% decomp_sc$xccoef
colnames(V_sc) <- paste0("CC", seq_len(ncol(V_sc)))

(prop_X <- rowMeans(cor(V_sc, X_sc)^2))
```

## Example ii

```
##    CC1    CC2    CC3    CC4    CC5    CC6    CC7    CC8
## 0.340 0.153 0.124 0.081 0.134 0.039 0.067 0.061
```

```
cumsum(prop_X)
```

```
##    CC1    CC2    CC3    CC4    CC5    CC6    CC7    CC8
## 0.34 0.49 0.62 0.70 0.83 0.87 0.94 1.00
```

## Example iii

*# But since we are dealing with correlations*  
*# We get the same with unstandardized variables*

```
decomp <- cancel(X, Y)
V <- X %*% decomp$xccoef
colnames(V) <- paste0("CC", seq_len(ncol(V)))

(prop_X <- rowMeans(cor(V, X)^2))
```

```
##   CC1   CC2   CC3   CC4   CC5   CC6   CC7   CC8
## 0.340 0.153 0.124 0.081 0.134 0.039 0.067 0.061
```

## Example iv

```
cumsum(prop_X)
```

```
##   CC1   CC2   CC3   CC4   CC5   CC6   CC7   CC8  
## 0.34 0.49 0.62 0.70 0.83 0.87 0.94 1.00
```

# Interpreting the population canonical variates i

- To help interpreting the canonical variates, let's go back to the population model.
- Define

$$A = \left( e_1^T \Sigma_Y^{-1/2} \quad \cdots \quad e_p^T \Sigma_Y^{-1/2} \right)^T,$$
$$B = \left( f_1^T \Sigma_X^{-1/2} \quad \cdots \quad f_p^T \Sigma_X^{-1/2} \right)^T.$$

- In other words, both  $A$  and  $B$  are  $p \times p$ , and their *rows* are the canonical directions.

## Interpreting the population canonical variates ii

- Using this notation, we can get all canonical variates using one linear transformation:

$$\mathbf{U} = \mathbf{A}\mathbf{Y}, \quad \mathbf{Y} = \mathbf{B}\mathbf{X}.$$

- We then have

$$\text{Cov}(\mathbf{U}, \mathbf{Y}) = \text{Cov}(\mathbf{A}\mathbf{Y}, \mathbf{Y}) = \mathbf{A}\Sigma_Y.$$

- Since  $\text{Cov}(\mathbf{U}) = \mathbf{I}_p$ , we have

$$\text{Corr}(U_k, Y_i) = \text{Cov}(U_k, \sigma_i^{-1}Y_i),$$

where  $\sigma_i^2$  is the variance of  $Y_i$ .

## Interpreting the population canonical variates iii

- If we let  $D_Y$  be the diagonal matrix whose  $i$ -th diagonal element is  $\sigma_i = \sqrt{\text{Var}(Y_i)}$ , we can write

$$\text{Corr}(\mathbf{U}, \mathbf{Y}) = A\Sigma_Y D_Y^{-1}.$$

- Using similar computations, we get

$$\begin{aligned}\text{Corr}(\mathbf{U}, \mathbf{Y}) &= A\Sigma_Y D_Y^{-1}, & \text{Corr}(\mathbf{V}, \mathbf{Y}) &= B\Sigma_{XY} D_Y^{-1}, \\ \text{Corr}(\mathbf{U}, \mathbf{X}) &= A\Sigma_{YX} D_X^{-1}, & \text{Corr}(\mathbf{V}, \mathbf{X}) &= B\Sigma_X D_X^{-1}.\end{aligned}$$

- **These quantities** (and their sample counterparts) **give us information about the contribution of the original variables to the canonical variates.**

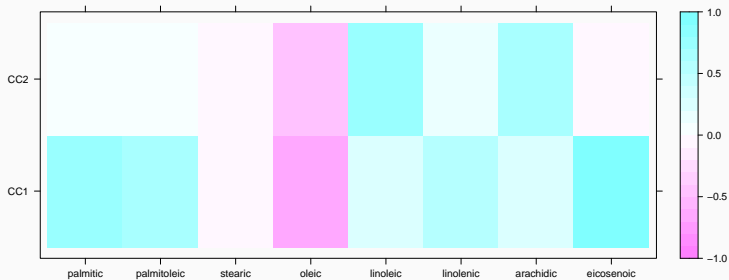


## Example i

```
# Let's go back to the olive data
decomp <- cancort(X, Y)
V <- X %*% decomp$xccoef
colnames(V) <- paste0("CC", seq_len(8))

library(lattice)
levelplot(cor(X, V[,1:2]),
          at = seq(-1, 1, by = 0.1),
          xlab = "", ylab = "")
```

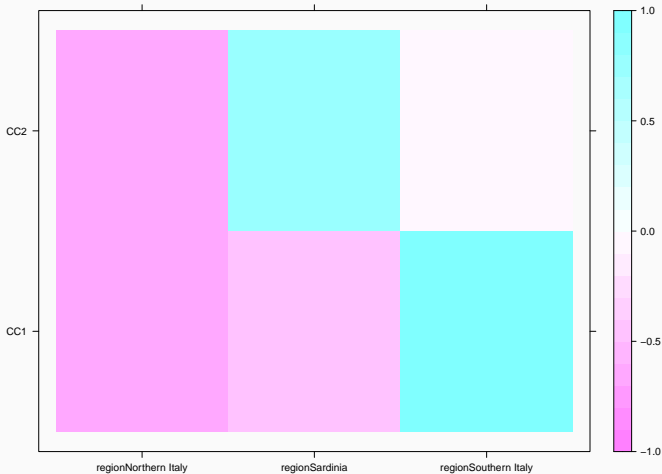
## Example ii



## Example iii

```
levelplot(cor(Y, V[,1:2]),  
          at = seq(-1, 1, by = 0.1),  
          xlab = "", ylab = "")
```

## Example iv



# Generalization of Correlation coefficients i

- The canonical correlations can be seen as a generalization of many notions of “correlation”.
- If both  $\mathbf{Y}, \mathbf{X}$  are one dimensional, then

$$\text{Corr}(a^T \mathbf{Y}, b^T \mathbf{X}) = \text{Corr}(\mathbf{Y}, \mathbf{X}), \quad \text{for all } a, b.$$

- In other words, the canonical correlation generalizes the **univariate correlation coefficient**.
- Then assume  $\mathbf{Y}$  is one-dimensional, but  $\mathbf{X}$  is  $q$ -dimensional. Then CCA is equivalent to (univariate) linear regression, and the first canonical correlation is equal to the **multiple correlation coefficient**.

## Generalization of Correlation coefficients ii

- Now, let's go back to full-generality:  $\mathbf{Y} = (Y_1, \dots, Y_p)$ ,  $\mathbf{X} = (X_1, \dots, X_q)$ . Let  $a$  be all zero except for a one in position  $i$ , and let  $b$  be all zero except for a one in position  $j$ . We have

$$\begin{aligned} |\text{Corr}(Y_i, X_j)| &= |\text{Corr}(a^T \mathbf{Y}, b^T \mathbf{X})| \\ &\leq \max_{a,b} \text{Corr}(a^T \mathbf{Y}, b^T \mathbf{X}) \\ &= \rho_1. \end{aligned}$$

## Generalization of Correlation coefficients iii

- In other words, the **first canonical correlation is larger than any entry** (in absolute value) **in the matrix**  $\text{Corr}(\mathbf{Y}, \mathbf{X})$ .
- Finally, the  $k$ -th canonical correlation  $\rho_k$  can be interpreted as the **multiple correlation coefficient** of two different univariate linear regression model:
  - $U_k$  against  $\mathbf{X}$ ;
  - $V_k$  against  $\mathbf{Y}$ .

## Example (cont'd) i

```
# Canonical correlations
```

```
decomp$cor
```

```
## [1] 0.95 0.84
```

```
# Maximum value in correlation matrix
```

```
max(abs(cor(Y, X)))
```

```
## [1] 0.89
```



## Example (cont'd) ii

*# Multiple correlation coefficients*

```
sqrt(summary(lm(V[,1] ~ Y))$r.squared)
```

```
## [1] 0.95
```

```
sqrt(summary(lm(V[,2] ~ Y))$r.squared)
```

```
## [1] 0.84
```

# Geometric interpretation i

- Let's look at a geometric interpretation of CCA.
- First, some notation:
  - Let  $A$  be the matrix whose  $k$ -th row is the  $k$ -th canonical direction  $e_k^T \Sigma_Y^{-1/2}$ .
  - Let  $E$  be the matrix whose  $k$ -th *column* is the eigenvector  $e_k$ . Note that  $E^T E = I_p$ .
  - We thus have  $A = E^T \Sigma_Y^{-1/2}$ .
- We get all canonical variates  $U_k$  by transforming  $\mathbf{Y}$  using  $A$ :

$$\mathbf{U} = \mathbf{A}\mathbf{Y}.$$

## Geometric interpretation ii

- Now, using the spectral decomposition of  $\Sigma_Y$ , we can write

$$A = E^T \Sigma_Y^{-1/2} = E^T P_Y \Lambda_Y^{-1/2} P_Y^T,$$

where  $P_Y$  contains the eigenvectors of  $\Sigma_Y$  and  $\Lambda_Y$  is the diagonal matrix with its eigenvalues.

- Therefore, we can see that

$$\mathbf{U} = A\mathbf{Y} = E^T P_Y \Lambda_Y^{-1/2} P_Y^T \mathbf{Y}.$$

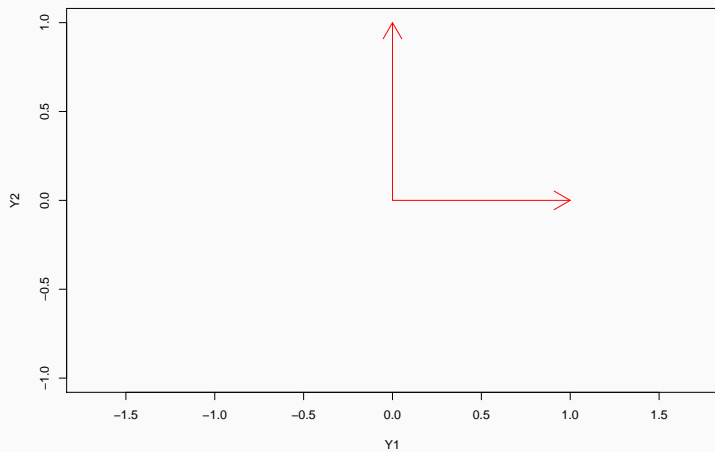
## Geometric interpretation iii

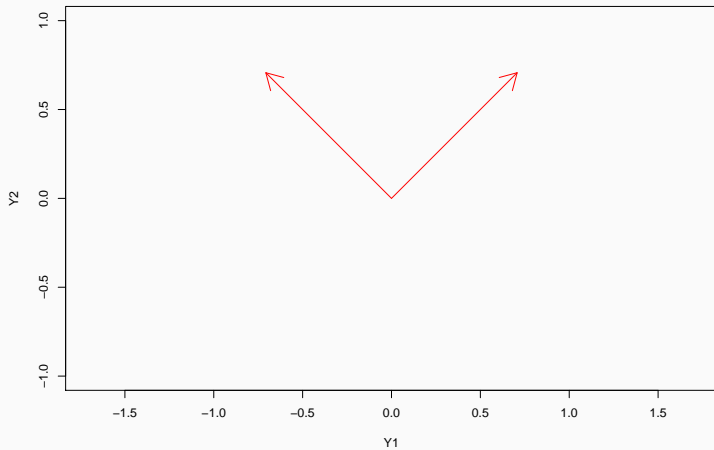
- Let's look at this expression in stages:
  - $P_Y^T \mathbf{Y}$ : This is the matrix of **principal components** of  $\mathbf{Y}$ .
  - $\Lambda_Y^{-1/2} (P_Y^T \mathbf{Y})$ : We standardize the principal components to have unit variance.
  - $P_Y (\Lambda_Y^{-1/2} P_Y^T \mathbf{Y})$ : We rotate the standardized PCs using a transformation that **only involves**  $\Sigma_Y$ .
  - $E^T (P_Y \Lambda_Y^{-1/2} P_Y^T \mathbf{Y})$ : We rotate the result using a transformation that **involves the whole covariance matrix**  $\Sigma$ .

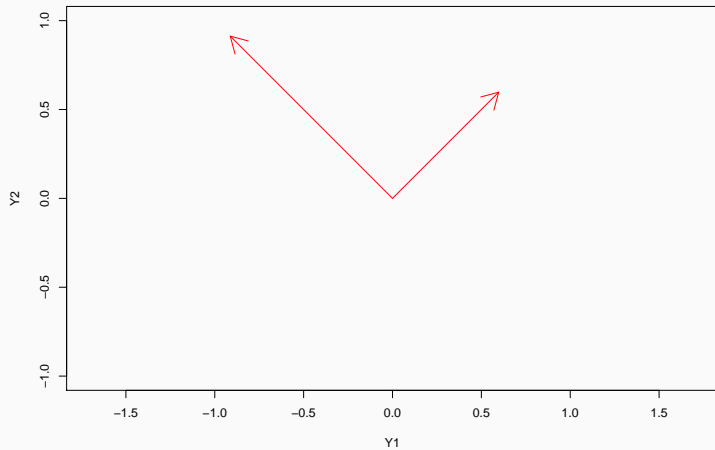
## Example i

- Let's go back to the covariance matrix at the beginning of this slide deck:

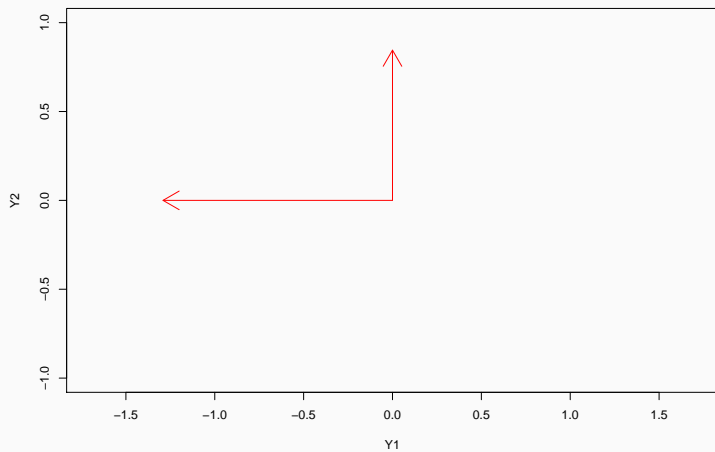
$$\Sigma = \begin{pmatrix} 1.0 & 0.4 & 0.5 & 0.6 \\ 0.4 & 1.0 & 0.3 & 0.4 \\ 0.5 & 0.3 & 1.0 & 0.2 \\ 0.6 & 0.4 & 0.2 & 1.0 \end{pmatrix}.$$

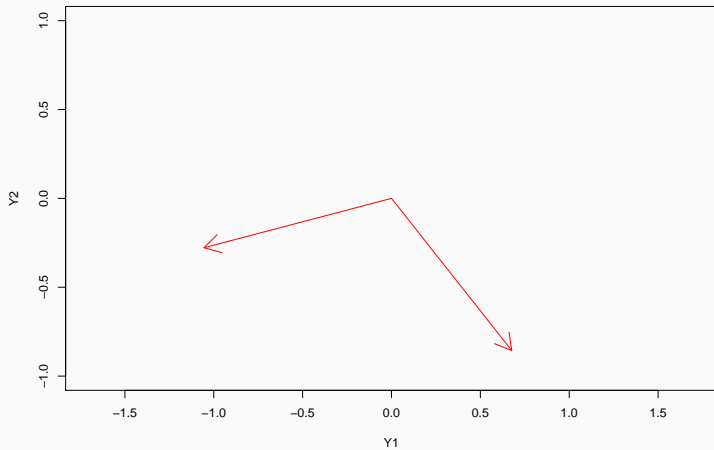












# Large sample inference

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# Test of independence i

- Recall what we said at the outset: CCA tries to explain the covariance  $\text{Cov}(\mathbf{Y}, \mathbf{X})$ .
- If there is no correlation between  $\mathbf{Y}, \mathbf{X}$ , then  $\Sigma_{YX} = 0$ .
  - In particular,  $a^T \Sigma_{YX} b = 0$  for any choice of  $a \in \mathbb{R}^p, b \in \mathbb{R}^q$ , and therefore all canonical correlations are equal to 0.
- To test for independence between  $\mathbf{Y}$  and  $\mathbf{X}$ , we will use a **likelihood ratio test**.

## LRT for $\Sigma_{YX} = 0$ i

Let  $(\mathbf{Y}_i, \mathbf{X}_i)$ ,  $i = 1, \dots, n$ , be a random sample from a normal distribution  $N_{p+q}(\mu, \Sigma)$ , with

$$\Sigma = \begin{pmatrix} \Sigma_Y & \Sigma_{YX} \\ \Sigma_{XY} & \Sigma_X \end{pmatrix}.$$

Let  $S_Y, S_X$  be the sample covariances of  $\mathbf{Y}_1, \dots, \mathbf{Y}_n$ , respectively, and let  $S_n$  be the  $p + q$ -dimensional sample covariance of  $(\mathbf{Y}_i, \mathbf{X}_i)$ .

## LRT for $\Sigma_{YX} = 0$ ii

Then the likelihood ratio test for  $H_0 : \Sigma_{YX} = 0$  rejects  $H_0$  for large values of

$$-2 \log \Lambda = n \log \left( \frac{|S_Y| |S_X|}{|S_n|} \right) = -n \log \prod_{i=1}^p (1 - \hat{\rho}_i^2),$$

where  $\hat{\rho}_1, \dots, \hat{\rho}_p$  are the sample canonical correlations.

# Null distribution

1. For large  $n$ , the statistic  $-2 \log \Lambda$  is approximately chi-square with degrees of freedom equal to

$$\left( \frac{(p+q)(p+q+1)}{2} \right) - \left( \frac{p(p+1)}{2} + \frac{q(q+1)}{2} \right) = pq.$$

2. Bartlett's correction uses a different statistic (but the same null distribution):

$$- \left( n - 1 - \frac{1}{2}(p+q+1) \right) \log \prod_{i=1}^p (1 - \hat{\rho}_i^2).$$

## Example i

- We will look at a different example, this time from the field of vegetation ecology.
- We have two datasets:
  - varechem: 14 chemical measurements from the soil.
  - varespec: 44 estimated cover values for lichen species.
- The data has 24 observations.
- For more details, see Väre, H., Ohtonen, R. and Oksanen, J. (1995) *Effects of reindeer grazing on understorey vegetation in dry Pinus sylvestris forests*. Journal of Vegetation Science 6, 523–530.



## Example ii

```
library(vegan)

data(varespec)
data(varechem)

# There are too many variables in varespec
# Let's pick first 10
Y <- varespec %>%
  select(Callvulg:Diphcomp) %>%
  as.matrix
```

## Example iii

```
# The help page in `vegan` suggests a better  
# chemical model
```

```
X <- varechem %>%  
  model.matrix( ~ A1 + P*(K + Baresoil) - 1,  
               data = .)
```

```
decomp <- cancort(x = X, y = Y)
```

```
n <- nrow(X)  
(LRT <- -n*log(prod(1 - decomp$cor^2)))
```

## Example iv

```
## [1] 156
```

```
p <- min(ncol(X), ncol(Y))  
q <- max(ncol(X), ncol(Y))  
LRT > qchisq(0.95, df = p*q)
```

```
## [1] TRUE
```

## Example v

```
LRT_bart <- -(n - 1 - 0.5*(p + q + 1)) *  
  log(prod(1 - decomp$cor^2))
```

```
c("Large Sample" = LRT,  
  "Bartlett" = LRT_bart)
```

```
## Large Sample      Bartlett  
##                156          94
```

```
LRT_bart > qchisq(0.95, df = p*q)
```

```
## [1] TRUE
```

## Sequential inference i

- The LRT above was for independence, i.e.  $\Sigma_{YX} = 0$ .
- Given our description of CCA above, this test is equivalent to having all canonical correlations being equal to 0.

$$\Sigma_{YX} = 0 \iff \rho_1 = \cdots = \rho_p = 0.$$

- If we reject the null hypothesis, it is natural to ask how many canonical correlations are nonzero.
- Recall that by design  $\rho_1 \geq \cdots \geq \rho_p$ . We thus get a sequence of null hypotheses:

$$H_0^k : \rho_1 \neq 0, \dots, \rho_k \neq 0, \rho_{k+1} = \cdots = \rho_p = 0.$$

## Sequential inference ii

- We can test the  $k$ -th hypothesis using a *truncated* version of the likelihood ratio test statistic:

$$LRT_k = - \left( n - 1 - \frac{1}{2}(p + q + 1) \right) \log \prod_{i=k+1}^p (1 - \hat{\rho}_i^2),$$

where its null distribution is approximately chi-square on  $(p - k)(q - k)$  degrees of freedom.

## Example (cont'd) i

*# We can get the truncated LRTs in one go*

```
(log_ccs <- rev(log(cumprod(1 - rev(decomp$cor)^2))))
```

```
## [1] -6.513 -4.002 -2.259 -1.011 -0.262 -0.073
```

```
(LRTs <- -(n - 1 - 0.5*(p + q + 1)) * log_ccs)
```

```
## [1] 94.4 58.0 32.7 14.7 3.8 1.1
```

## Example (cont'd) ii

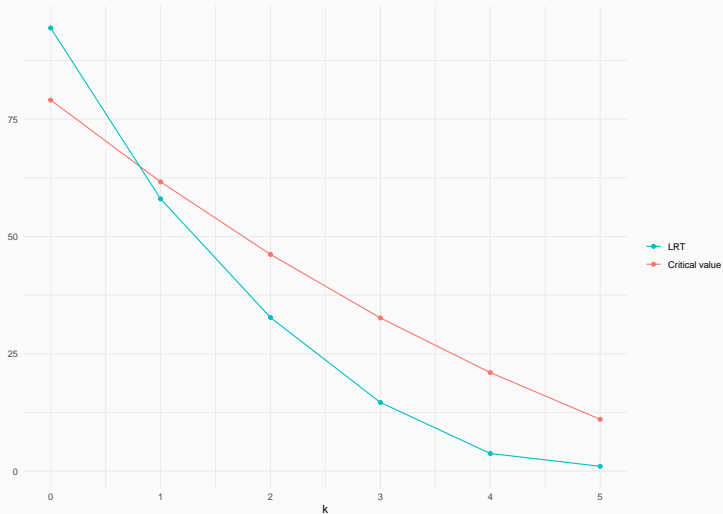
```
k_seq <- seq(0, p - 1)
LRTs > qchisq(0.95,
              df = (p - k_seq)*(q - k_seq))
```

```
## [1] TRUE FALSE FALSE FALSE FALSE FALSE
```

```
# We only reject the first null hypothesis
# of independence
```



## Example (cont'd) iii



# Summary

- CCA is a dimension reduction method like PCA
  - But we are reducing the dimension of two datasets **jointly**.
  - Instead of maximising variance, we maximise **correlation**.
- The goal is to explain the association between **Y** and **X**.
  - Unlike MLR, both datasets are treated equally.
- All visualization methods we discussed in the context of PCA (e.g. component plots, loading plots, biplots) are available for CCA.
  - See the R package `vegan`.
- **Limitation:** CCA performs poorly when  $p$  and/or  $q$  are close to  $n$ .