### Correspondence Analysis

ST 558: Multivariate Analytics

Module 9

Lecture 2

#### Correspondence Analysis: Overview

Correspondence analysis is another technique primarily used to graphically represent high-dimensional data; specifically, it is used to display *contingency table* data, most commonly in two dimensions.

The resulting representation gives an indication of

- Which rows are similar to each other
- Which columns are similar to each other
- Rows and columns that have higher than expected association with each other.

### Correspondence Analysis: Background

Recall that a <u>contingency table</u> is a way of representing the relationship between two (or more) different <u>categorical</u> variables:

		Row			
Variable 1	Level 1	Level 2	•••	Level c	Totals
Level 1	$n_{11}$	$n_{12}$	•••	$n_{1c}$	$n_1$ .
Level 2	$n_{21}$	$n_{22}$	•••	$n_{2c}$	$n_2$ .
:	:	:	٠.	••	
Level r	$n_{r1}$	$n_{r2}$	•••	$n_{rc}$	$n_r$ .
Column Totals	$n_{\cdot 1}$	n. <sub>2</sub>		$n_{\cdot c}$	n

# Correspondence Analysis: Background

We will let **P** be the matrix of *relative frequencies*.

$$\mathbf{P} = \begin{bmatrix} n_{11}/n & n_{12}/n & \cdots & n_{1c}/n \\ n_{21}/n & n_{22}/n & \cdots & n_{2c}/n \\ \vdots & \vdots & \ddots & \vdots \\ n_{r1}/n & n_{r2}/n & \cdots & n_{rc}/n \end{bmatrix}$$

Define

• 
$$\mathbf{r} = \frac{1}{n} [n_{1}, n_{2}, ..., n_{r}]^{T}$$
: the vector of row frequencies

• 
$$\mathbf{c} = \frac{1}{n}[n_1, n_2, \dots, n_c]^T$$
: the vector of *column frequencies*

### Correspondence Analysis: Background [n\_1]

r - 1	$\begin{vmatrix} n_1 \\ n_2 \end{vmatrix}$
$\frac{1-n}{n}$	
7.0	l n
	$\lfloor n_r \rfloor$

		Row			
Variable 1	Level 1	Level 2	•••	Level c	Totals
Level 1	$n_{11}$	$n_{12}$	•••	$n_{1c}$	$n_1$ .
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:	:	:	٠.	:	
Level r	$n_{r1}$	$n_{r2}$	•••	$n_{rc}$	$n_r$ .
Column Totals	n. <sub>1</sub>	n. <sub>2</sub>		$n_{\cdot c}$	n

$$\mathbf{c} = \frac{1}{n} \begin{bmatrix} n_{\cdot 1} \\ n_{\cdot 2} \\ \vdots \\ n_{\cdot c} \end{bmatrix}$$

**Example:** Suppose that we sample *n* = 200 OSU students and ask each sampled student

- What their favorite ice cream flavor is:
  - o Chocolate, Vanilla, or Strawberry
- What their favorite sport is:
  - o Baseball, Basketball, Football, Soccer, or Tennis

#### We obtain the following data:

	Baseball	Basketball	Football	Soccer	Tennis
Chocolate	11	20	17	11	11
Vanilla	6	17	19	7	17
Strawberry	18	8	8	17	13

Correspondence analysis proceeds via the following steps:

- 1. Calculate the *expected frequency matrix* (that is, the expected table if the rows and columns were *independent*)
- 2. Calculate the matrix of *deviations from independence*
- 3. Calculate the *generalized singular value decomposition* (generalized SVD) of the deviations from independence
- 4. Use the first several (typically 2) dimensions of the generalized SVD, **appropriately scaled**, as the coordinates for plotting the rows (columns, respectively) of the contingency table

- 1. Calculate the *expected frequency matrix* (that is, the expected table if the rows and columns were *independent*)
  - The expected frequency matrix if the row variable and column variable are independent of each other is given by

$$\widehat{\mathbf{P}} = \mathbf{r}\mathbf{c}^T$$

2. Calculate the matrix of *deviations from independence* 

$$\mathbf{P} - \widehat{\mathbf{P}} = \mathbf{P} - \mathbf{r}\mathbf{c}^T$$

- 1. Calculate the *expected frequency matrix* (that is, the expected table if the rows and columns
  - were independent)
  - The expected frequence column variable are in given by

This difference matrix tells us how extreme the individual cells in the contingency table are: elements with larger values in this deviations matrix are more unusual if the variables are independent.

2. Calculate the matrix of *deviations from independence* 

 $\widehat{\mathbf{p}}$ 

$$\mathbf{P} - \widehat{\mathbf{P}} = \mathbf{P} - \mathbf{r}\mathbf{c}^T$$

- 3. Calculate the *generalized singular value decomposition* (generalized SVD) of the deviations from independence
  - $\circ$  The *generalized SVD* is a way of expressing the matrix  $\mathbf{P} \mathbf{rc}^T$  as a matrix product:

$$\mathbf{P} - \mathbf{r} \mathbf{c}^T = \mathbf{A} \mathbf{\Lambda} \mathbf{B}^T$$

where  $\Lambda$  is a diagonal matrix, and the columns of A and B are orthogonal to each other, and scaled so that the *norm* of each column is proportional to the sum of the corresponding row/column:

$$\mathbf{A}^T \mathbf{D}_r \mathbf{A} = \mathbf{B}^T \mathbf{D}_c \mathbf{B} = \mathbf{I}$$

3. Calculate the *generalized singular value decomposition* (generalized SVD) of the

$$\mathbf{D}_{r} = \operatorname{diag}(\mathbf{r})$$

$$= \begin{bmatrix} n_{1}./n & 0 & \cdots & 0 \\ 0 & n_{2}./n & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & n_{r}./n \end{bmatrix} \quad \mathbf{D}_{c} = \operatorname{diag}(\mathbf{c})$$

$$= \begin{bmatrix} n_{.1}/n & 0 & \cdots & 0 \\ 0 & n_{.2}/n & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & n_{.c}/n \end{bmatrix} \quad \mathbf{ix}$$

and **B** are orthogonal to each other, and scaled so that the *norm* of each column is proportional to the sum of the corresponding row/column:

$$\mathbf{A}^T \mathbf{D}_r \mathbf{A} = \mathbf{B}^T \mathbf{D}_c \mathbf{B} = \mathbf{I}$$

The generalized SVD tells us the predominant patterns in the rows and columns of a matrix.

3. Calcula decomp deviation of The ger

Like the eigendecomposition/PCA, the first few components (columns of and A, B, with corresponding diagonal elements of  $\Lambda$ ) of this matrix product can give us a good approximation to the matrix.

 $\mathbf{P} - \mathbf{r} \mathbf{c}^T$  as a matrix product:

$$\mathbf{P} - \mathbf{r}\mathbf{c}^T = \mathbf{A}\mathbf{\Lambda}\mathbf{B}^T$$

where  $\Lambda$  is a diagonal matrix, and the columns of A and B are orthogonal to each other, and scaled so that the *norm* of each column is proportional to the sum of the corresponding row/column:

$$\mathbf{A}^T \mathbf{D}_r \mathbf{A} = \mathbf{B}^T \mathbf{D}_c \mathbf{B} = \mathbf{I}$$

4. Use the first several (typically 2) dimensions of the generalized SVD, appropriately scaled, as the coordinates for plotting the rows (columns, respectively)

$$\mathbf{X} = \mathbf{D}_r^{-1} \mathbf{A} \mathbf{\Lambda}$$

$$\mathbf{Y} = \mathbf{D}_c^{-1} \mathbf{B} \mathbf{\Lambda}$$

4. Use the first several (typically 2) dimensions of the generalized SVD, appropriately scaled, as the coordinates for plotting the rows (columns, respectively)

Columns of **X** give coordinates for plotting the <u>row</u> deviations.

$$X = D_r^{-1} A \Lambda$$

$$\mathbf{Y} = \mathbf{D}_c^{-1} \mathbf{B} \mathbf{\Lambda}$$

Columns of **Y** give coordinates for plotting the <u>column</u> deviations.

4. Use the first several (typically 2) dimensions of the generalized SVD, appropriately scaled, as the coordinates for plotting the rows (columns, respectively)

Typically we plot the first two columns of X, with each point representing the corresponding row of  $P - rc^T$ 

$$\mathbf{X} = \mathbf{D}_r^{-1} \mathbf{A} \mathbf{\Lambda}$$

$$\mathbf{Y} = \mathbf{D}_c^{-1} \mathbf{B} \mathbf{\Lambda}$$

Typically we plot the first two columns of **Y**, with each point representing the corresponding column of **P** – **rc**<sup>T</sup>

**Example:** Suppose that we sample n = 200 OSU students and ask each sampled student

- What their favorite ice cream flavor is:
  - o Chocolate, Vanilla, or Strawberry
- What their favorite sport is:
  - o Baseball, Basketball, Football, Soccer, or Tennis

	Baseball	Basketball	Football	Soccer	Tennis	Row Totals
Chocolate	11	20	17	11	11	70
Vanilla	6	17	19	7	17	66
Strawberry	18	8	8	17	13	64
Column Totals	35	45	44	35	41	200

**Example:** Suppose that we sample *n* = 200 OSU students and ask each sampled student

- What their favorite ice cream flavor is
- What their favorite sport is

The relative frequency matrix is therefore:

	Baseball	Basketball	Football	Soccer	Tennis	Row Totals
Chocolate	0.055	0.100	0.085	0.055	0.055	0.350
Vanilla	0.030	0.085	0.095	0.035	0.085	0.330
Strawberry	0.090	0.040	0.040	0.085	0.065	0.320
Column Totals	0.175	0.225	0.22	0.175	0.205	1.000

**Example:** Suppose that we sample *n* = 200 OSU students and ask each sampled student

- What their favorite ice cream flavor is
- What their favorite sport is

The *expected* relative frequency matrix is therefore:

	Baseball	Basketball	Football	Soccer	Tennis	Row Totals
Chocolate	0.061	0.079	0.077	0.061	0.072	0.350
Vanilla	0.058	0.074	0.073	0.058	0.068	0.330
Strawberry	0.056	0.072	0.070	0.056	0.066	0.320
Column Totals	0.175	0.225	0.22	0.175	0.205	1.000

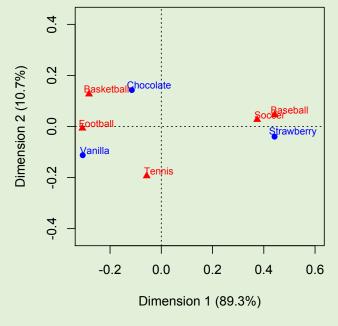
**Example:** Suppose that we sample *n* = 200 OSU students and ask each sampled student

- What their favorite ice cream flavor is
- What their favorite sport is

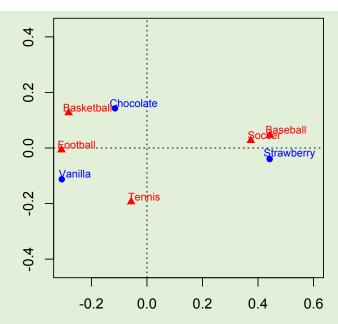
The matrix of *deviations from independence* is therefore:

	Baseball	Basketball	Football	Soccer	Tennis	Row Totals
Chocolate	-0.006	0.021	0.008	-0.006	-0.017	0
Vanilla	-0.028	0.011	0.022	-0.023	0.017	0
Strawberry	0.034	-0.032	-0.030	0.029	-0.001	0
Column Totals	0	0	0	0	0	0

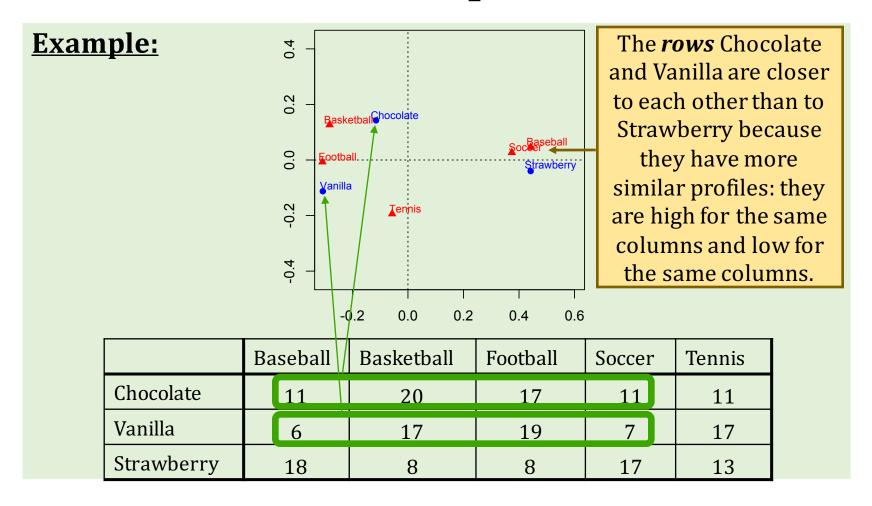
**Example:** Suppose that we sample n = 200 OSU students and ask each sampled student what their favorite ice cream flavor is and what their favorite sport is. The resulting correspondence analysis plot is:

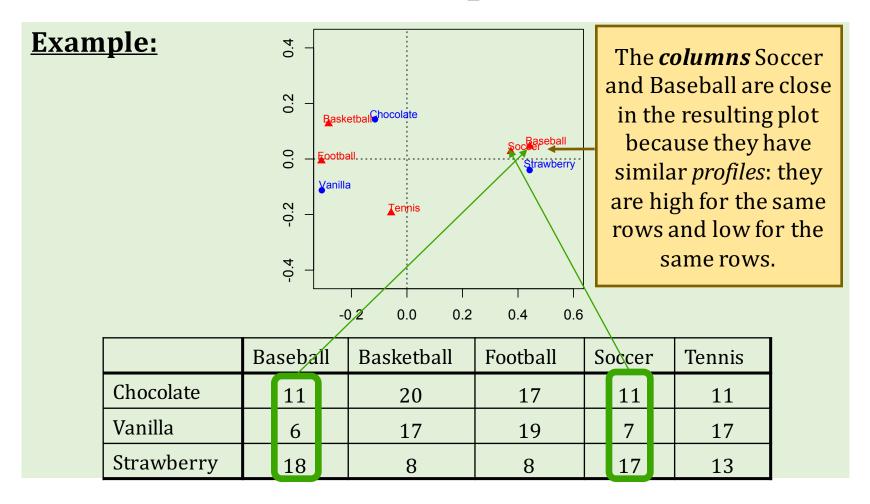


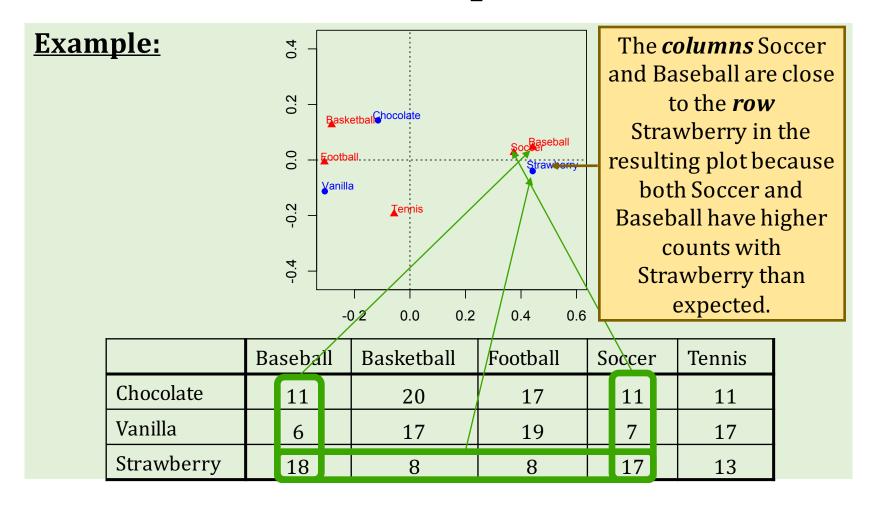




	Baseball	Basketball	Football	Soccer	Tennis
Chocolate	11	20	17	11	11
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### Correspondence Analysis: Summary

Correspondence analysis is a *descriptive* technique used to visualize and summarize relationships between categorical variables.

We have considered only two-way tables: that is, tables comparing two different categorical variables. Correspondence analysis may be extended to an arbitrary number of categorical variables.