

FINM 331: DATA ANALYSIS FOR FINANCE AND STATISTICS
FALL 2015
CORRESPONDENCE ANALYSIS

- we are going to look at variants of PCA that apply to frequency data (CA), network data (HITS), and text data (LSA)
- we are also going to look at a variant of PCA called MDS that uses dissimilarity/distance matrix in place of covariance/inner product matrix
- a recap of the matrix decompositions that we have been using and a new one that we will introduce:

SVD: for PCA, FA, HITS, LSI; $A \in \mathbb{R}^{n \times p}$,

$$A = U\Sigma V^T, \quad U^T U = I_n, \quad V^T V = I_p$$

EVD: for CCA, MDS; $A \in \mathbb{R}^{p \times p}$ symmetric,

$$X = Q\Lambda Q^T, \quad Q^T Q = I_p$$

GEVD: for LDA; $A \in \mathbb{R}^{p \times p}$ symmetric, $B \in \mathbb{R}^{p \times p}$ symmetric nonsingular,

$$B^{-1}A = Q\Lambda Q^T, \quad Q^T Q = I_p$$

GSVD: for CA; $A \in \mathbb{R}^{n \times p}$, $D_1 \in \mathbb{R}^{n \times n}$ diagonal nonsingular, $D_2 \in \mathbb{R}^{p \times p}$ diagonal nonsingular,

$$A = U\Sigma V^T, \quad U^T D_1^{-1} U = I_n, \quad V^T D_2^{-1} V = I_p$$

1. CORRESPONDENCE ANALYSIS

- in this case, the data matrix $X \in \mathbb{R}^{n \times p}$ is (part of) a *contingency table*
- a contingency table is essentially a table of *count* or *frequency* data
- an example is the Greenacre smokers data set

	none	light	medium	heavy	row total
senior managers	4	2	3	2	11
junior managers	4	3	7	4	18
senior employees	25	10	12	4	51
junior employees	18	24	33	13	88
secretaries	10	6	7	2	25
column total	61	54	62	25	193

- the bold faced row and column totals as well as the grand total can all be computed from the matrix

$$X = \begin{bmatrix} 4 & 2 & 3 & 2 \\ 4 & 3 & 7 & 4 \\ 25 & 10 & 12 & 4 \\ 18 & 24 & 33 & 13 \\ 10 & 6 & 7 & 2 \end{bmatrix} \in \mathbb{R}^{5 \times 4}$$

so this matrix in the middle forms the essence of a contingency table

- here $n = 5$ is the number of staff categories and $p = 4$ is the number of smoking categories — in a contingency table, the row and column categories are treated on equal footing
- more generally, for a data matrix $X = [x_{ij}] \in \mathbb{R}^{n \times p}$ that comes from a contingency table, x_{ij} 's are *frequencies*, i.e.,

x_{ij} = number of observations in a sample that falls into row category i and column category j

- alternatively, there are people who prefer to normalize by the grand total before forming the data matrix
- for the smokers data set, this gives

$$X = \begin{bmatrix} \frac{4}{193} & \frac{2}{193} & \frac{3}{193} & \frac{2}{193} \\ \frac{193}{25} & \frac{193}{10} & \frac{193}{12} & \frac{193}{4} \\ \frac{193}{18} & \frac{193}{24} & \frac{193}{33} & \frac{193}{13} \\ \frac{193}{10} & \frac{193}{6} & \frac{193}{7} & \frac{193}{2} \\ \frac{193}{193} & \frac{193}{193} & \frac{193}{193} & \frac{193}{193} \end{bmatrix} \in \mathbb{R}^{5 \times 4}$$

- here x_{ij} 's are *relative frequencies* instead of frequencies
- the only difference is that the result would differ by a constant so we would just stick to the frequency version in the following
- the most notations for the row, column, and grand totals are

$$\begin{aligned} x_{i\bullet} &= \sum_{j=1}^p x_{ij}, & i &= 1, \dots, n, \\ x_{\bullet j} &= \sum_{i=1}^n x_{ij}, & j &= 1, \dots, p, \\ x_{\bullet\bullet} &= \sum_{i=1}^n \sum_{j=1}^p x_{ij} = \sum_{i=1}^n x_{i\bullet} = \sum_{j=1}^p x_{\bullet j} = \mathbf{1}_n^\top X \mathbf{1}_p \end{aligned}$$

- the objective in *correspondence analysis* (CA) is to find a *row weight vector* and a *column weight vector*,

$$\mathbf{r} = \begin{bmatrix} r_1 \\ \vdots \\ r_n \end{bmatrix} \in \mathbb{R}^n \quad \text{and} \quad \mathbf{c} = \begin{bmatrix} c_1 \\ \vdots \\ c_p \end{bmatrix} \in \mathbb{R}^p$$

such that

$$\begin{aligned} r_i &\propto \sum_{j=1}^p c_j \frac{x_{ij}}{x_{i\bullet}}, & i &= 1, \dots, n, \\ c_j &\propto \sum_{i=1}^n r_i \frac{x_{ij}}{x_{\bullet j}}, & j &= 1, \dots, p, \end{aligned}$$

hold simultaneously

- here ' $x \propto y$ ' means ' x is proportional to y ,' i.e., $x = \lambda y$ for some constant $\lambda \in \mathbb{R}$
- now let

$$D_r = \begin{bmatrix} x_{1\bullet} & & \\ & \ddots & \\ & & x_{n\bullet} \end{bmatrix} \in \mathbb{R}^{n \times n}, \quad D_c = \begin{bmatrix} x_{\bullet 1} & & \\ & \ddots & \\ & & x_{\bullet p} \end{bmatrix} \in \mathbb{R}^{p \times p}$$

and rewrite the above equations in terms of vectors, we get

$$\mathbf{r} \propto D_r^{-1} X \mathbf{c}, \quad \mathbf{c} \propto D_c^{-1} X^\top \mathbf{r} \quad (1.1)$$

- substituting one into the other, we get

$$\mathbf{r} \propto D_r^{-1} X D_c^{-1} X^\top \mathbf{r}, \quad \mathbf{c} \propto D_c^{-1} X^\top D_r^{-1} X \mathbf{c} \quad (1.2)$$

- so $\mathbf{r} \in \mathbb{R}^n$ is an eigenvector of $D_r^{-1} X D_c^{-1} X^\top \in \mathbb{R}^{n \times n}$ and $\mathbf{c} \in \mathbb{R}^p$ is an eigenvector of $D_c^{-1} X^\top D_r^{-1} X \in \mathbb{R}^{p \times p}$
- note that the nonzero eigenvalues of $D_r^{-1} X D_c^{-1} X^\top = D_r^{-1} X (X D_c^{-1})^\top$ and $D_c^{-1} X^\top D_r^{-1} X = (X D_c^{-1})^\top D_r^{-1} X$ are the same since the nonzero eigenvalues of AB always equals the nonzero eigenvalues of BA for any $A \in \mathbb{R}^{n \times p}$ and $B \in \mathbb{R}^{p \times n}$
- we are only interested in nonzero eigenvalues and so (1.2) becomes

$$D_r^{-1} X D_c^{-1} X^\top \mathbf{r} = \lambda \mathbf{r}, \quad D_c^{-1} X^\top D_r^{-1} X \mathbf{c} = \lambda \mathbf{c}, \quad (1.3)$$

where $0 \neq \lambda \in \mathbb{R}$

- if $D_r = I_n$ and $D_c = I_p$, i.e., the $n \times n$ and $p \times p$ identity matrices, then (1.3) reduces to

$$X X^\top \mathbf{r} = \mathbf{r}, \quad X^\top X \mathbf{c} = \mathbf{c},$$

i.e., \mathbf{r} and \mathbf{c} are left and right singular vectors of X

2. GENERALIZED SINGULAR VALUE DECOMPOSITION

- but $D_r \neq I_n$ and $D_c \neq I_p$ in general and that's why we need the *generalized singular value decomposition* or GSVD

$$X = U \Sigma V^\top, \quad U^\top D_r^{-1} U = I_n, \quad V^\top D_c^{-1} V = I_p, \quad (2.1)$$

which can be shown to exist for any symmetric positive definite matrices $D_1 \in \mathbb{R}^{n \times n}$ and $D_2 \in \mathbb{R}^{p \times p}$ and any $X \in \mathbb{R}^{n \times p}$

- for our purpose we just need $D_r = \text{diag}(x_{1\bullet}, \dots, x_{n\bullet})$ and $D_c = \text{diag}(x_{\bullet 1}, \dots, x_{\bullet p})$, obviously symmetric positive definite since $x_{i\bullet} > 0$ and $x_{\bullet j} > 0$ for all $i = 1, \dots, n$ and $j = 1, \dots, p$
- we will call $\mathbf{u}_k \in \mathbb{R}^n$, $\mathbf{v}_k \in \mathbb{R}^p$ the left and right *generalized singular vectors* and $\sigma_k \in \mathbb{R}$ *generalized singular values* of $X \in \mathbb{R}^{n \times p}$ with respect to *weight matrices* D_r and D_c
- so you know how to compute GSVD or use a software that has a GSVD function, then you could compute the row and column weight vectors as generalized left and right singular vectors, i.e.,

$$\mathbf{r}_k = \mathbf{u}_k, \quad \mathbf{c}_k = \mathbf{v}_k \quad (2.2)$$

- however GSVD is not as common as SVD, for instance it's not in R unless you load some additional packages
- in the following we will see how to do CA with the usual SVD

3. CORRESPONDENCE ANALYSIS WITH SVD

- we need first deal with an issue: check that

$$D_r^{-1} X D_c^{-1} X^\top \mathbf{1}_n = \mathbf{1}_n, \quad D_c^{-1} X^\top D_r^{-1} X \mathbf{1}_p = \mathbf{1}_p,$$

i.e., these matrices have a trivial eigenvector, a vector of all ones, corresponding to the eigenvalue 1, that we want to exclude from our possibilities for \mathbf{r} and \mathbf{c}

- in other words, we want to exclude the left generalized singular vector $\mathbf{1}_n$ and right generalized singular vector $\mathbf{1}_p$ corresponding to the generalized singular value 1 from the GSVD of X

- how do you ‘exclude’ a pair of left/right singular vectors $\mathbf{u}_i, \mathbf{v}_i$, corresponding to a singular value σ_i ? the answer is *deflation*, i.e., just subtract the rank-1 matrix $\sigma_i \mathbf{u}_i \mathbf{v}_i^\top$ from X :

$$X - \sigma_i \mathbf{u}_i \mathbf{v}_i^\top$$

- why does this work? remember that the SVD of X can be written as

$$X = \sigma_1 \mathbf{u}_1 \mathbf{v}_1^\top + \cdots + \sigma_i \mathbf{u}_i \mathbf{v}_i^\top + \cdots + \sigma_r \mathbf{u}_r \mathbf{v}_r^\top$$

and if we subtract $\sigma_i \mathbf{u}_i \mathbf{v}_i^\top$, we get

$$X - \sigma_i \mathbf{u}_i \mathbf{v}_i^\top = \sigma_1 \mathbf{u}_1 \mathbf{v}_1^\top + \cdots + \sigma_{i-1} \mathbf{u}_{i-1} \mathbf{v}_{i-1}^\top + \sigma_{i+1} \mathbf{u}_{i+1} \mathbf{v}_{i+1}^\top + \cdots + \sigma_r \mathbf{u}_r \mathbf{v}_r^\top,$$

i.e., the term $\sigma_i \mathbf{u}_i \mathbf{v}_i^\top$ no longer appears in the SVD of $X - \sigma_i \mathbf{u}_i \mathbf{v}_i^\top$

- this also works for GSVD, but we have to scale the rank-1 term accordingly
- doing deflation gives us the matrix

$$X - \frac{\mathbf{a} \mathbf{b}^\top}{x_{\bullet\bullet}} \in \mathbb{R}^{n \times p} \quad (3.1)$$

where

$$\mathbf{a} = D_r \mathbf{1}_n \in \mathbb{R}^n, \quad \mathbf{b} = D_c \mathbf{1}_p \in \mathbb{R}^p$$

- now we could perform GSVD to (3.1) but we want to avoid GSVD and use only SVD, so we consider the matrix

$$Y := \sqrt{x_{\bullet\bullet}} D_r^{-1/2} \left(X - \frac{\mathbf{a} \mathbf{b}^\top}{x_{\bullet\bullet}} \right) D_c^{-1/2} \in \mathbb{R}^{n \times p}$$

instead

- if we work out the entries of Y , we would see that

$$y_{ij} = \frac{x_{ij} - e_{ij}}{\sqrt{e_{ij}}}$$

where

$$e_{ij} = \frac{x_{i\bullet} x_{\bullet j}}{x_{\bullet\bullet}}$$

for $i = 1, \dots, n$ and $j = 1, \dots, p$

- the square of the Frobenius norm of Y is then

$$\|Y\|_F^2 = \sum_{i=1}^n \sum_{j=1}^p y_{ij}^2 = \sum_{i=1}^n \sum_{j=1}^p \frac{(x_{ij} - e_{ij})^2}{e_{ij}} \quad (3.2)$$

- for those who know some statistics the expression on the right of (3.2) is a χ^2 -test statistics and e_{ij} may be interpreted as the *estimated expected value*
- now we may perform a usual SVD to Y to obtain

$$Y = U \Sigma V^\top, \quad U^\top U = I_n, \quad V^\top V = I_p$$

where $U = [\mathbf{u}_1, \dots, \mathbf{u}_n]$ and $V = [\mathbf{v}_1, \dots, \mathbf{v}_p]$

- in particular

$$Y \mathbf{v}_k = \sigma_k \mathbf{u}_k, \quad Y^\top \mathbf{u}_k = \sigma_k \mathbf{v}_k, \quad k = 1, \dots, r$$

where $r = \text{rank}(Y)$

- we claim that defining $\mathbf{r}_k := \sigma_k D_r^{-1/2} \mathbf{u}_k$ and $\mathbf{c}_k := \sigma_k D_c^{-1/2} \mathbf{v}_k$ gives us a solution to (1.1)

- to show this, note that

$$\begin{aligned}\mathbf{r}_k &= \sigma_k D_r^{-1/2} \mathbf{u}_k = D_r^{-1/2} (\sigma_k \mathbf{u}_k) = D_r^{-1/2} Y \mathbf{v}_k \\ &= D_r^{-1/2} Y \left(\frac{1}{\sigma_k} D_c^{1/2} \mathbf{c}_k \right) = \frac{1}{\sigma_k} D_r^{-1/2} Y D_c^{1/2} \mathbf{c}_k,\end{aligned}$$

and likewise

$$\mathbf{c}_k = \frac{1}{\sigma_k} D_c^{-1/2} Y^\top D_r^{1/2} \mathbf{r}_k$$

- since $\mathbf{a}^\top \mathbf{r}_k = 0$ and $\mathbf{b}^\top \mathbf{c}_k = 0$ (see appendix below), we get

$$\mathbf{r}_k = \frac{\sqrt{x_{\bullet\bullet}}}{\sigma_k} D_r^{-1} X \mathbf{c}_k, \quad \mathbf{c}_k = \frac{\sqrt{x_{\bullet\bullet}}}{\sigma_k} D_c^{-1} X^\top \mathbf{r}_k \quad (3.3)$$

as required

- note that

$$\mathbf{r}_i^\top D_r \mathbf{r}_j = \sigma_i \sigma_j \mathbf{u}_i^\top D_r^{-1/2} D_r D_r^{-1/2} \mathbf{u}_j = \sigma_i \sigma_j \mathbf{u}_i^\top \mathbf{u}_j = \begin{cases} \sigma_i^2 & \text{if } i = j, \\ 0 & \text{if } i \neq j, \end{cases}$$

and

$$\mathbf{c}_i^\top D_c \mathbf{c}_j = \sigma_i \sigma_j \mathbf{v}_i^\top D_c^{-1/2} D_c D_c^{-1/2} \mathbf{v}_j = \sigma_i \sigma_j \mathbf{v}_i^\top \mathbf{v}_j = \begin{cases} \sigma_i^2 & \text{if } i = j, \\ 0 & \text{if } i \neq j, \end{cases}$$

- this is not surprising since the method using GSVD (2.2) and the method using SVD give the same solution (3.3) up to a constant
- the sample means of \mathbf{r}_k and \mathbf{c}_k are both zero since

$$\frac{1}{x_{\bullet\bullet}} \mathbf{a}^\top \mathbf{r}_k = 0, \quad \frac{1}{x_{\bullet\bullet}} \mathbf{b}^\top \mathbf{c}_k = 0$$

- the sample variances of \mathbf{r}_k and \mathbf{c}_k are both $\sigma_k^2/x_{\bullet\bullet}$ since

$$\frac{1}{x_{\bullet\bullet}} \sum_{i=1}^n x_{i\bullet} r_{ki}^2 = \frac{\mathbf{r}_k^\top D_r \mathbf{r}_k}{x_{\bullet\bullet}} = \frac{\sigma_k^2}{x_{\bullet\bullet}}$$

and

$$\frac{1}{x_{\bullet\bullet}} \sum_{j=1}^p x_{\bullet j} c_{kj}^2 = \frac{\mathbf{c}_k^\top D_c \mathbf{c}_k}{x_{\bullet\bullet}} = \frac{\sigma_k^2}{x_{\bullet\bullet}}$$

- we use CA the way we use PCA — scree plots, scatter plots, biplots — but with \mathbf{r}_k and \mathbf{c}_k playing the roles of \mathbf{u}_k and \mathbf{v}_k in PCA

4. APPENDIX

- we verify that $\mathbf{a}^\top \mathbf{r}_k = 0$ and $\mathbf{b}^\top \mathbf{c}_k = 0$
- for your convenience

$$\begin{aligned}\mathbf{r}_k &= \sigma_k D_r^{-1/2} \mathbf{u}_k, & \mathbf{a} &= D_r \mathbf{1}_n, & \mathbf{b} &= D_c \mathbf{1}_p, \\ Y &= \sqrt{x_{\bullet\bullet}} D_r^{-1/2} \left(X - \frac{\mathbf{a} \mathbf{b}^\top}{x_{\bullet\bullet}} \right) D_c^{-1/2}, & Y \mathbf{v}_k &= \sigma_k \mathbf{u}_k\end{aligned}$$

- so

$$\begin{aligned}\mathbf{a}^\top \mathbf{r}_k &= \mathbf{1}_n^\top D_r^{1/2} (\sigma_k \mathbf{u}_k) = \mathbf{1}_n^\top D_r^{1/2} Y \mathbf{v}_k \\ &= \sqrt{x_{\bullet\bullet}} \mathbf{1}_n^\top \left(X - \frac{\mathbf{a} \mathbf{b}^\top}{x_{\bullet\bullet}} \right) D_c^{-1/2} \mathbf{v}_k\end{aligned}$$

- we claim that

$$\left(X - \frac{\mathbf{a}\mathbf{b}^\top}{x_{\bullet\bullet}}\right)^\top \mathbf{1}_n = \mathbf{0}_p$$

- this follows from

$$X^\top \mathbf{1}_n = \begin{bmatrix} x_{\bullet 1} \\ x_{\bullet 2} \\ \vdots \\ x_{\bullet p} \end{bmatrix}$$

and

$$\mathbf{b}\mathbf{a}^\top \mathbf{1}_n = (\mathbf{1}_n^\top D_r \mathbf{1}_n) \mathbf{b} = (x_{1\bullet} + \cdots + x_{n\bullet}) \mathbf{b} = x_{\bullet\bullet} \mathbf{b}$$

and so

$$\left(\frac{\mathbf{a}\mathbf{b}^\top}{x_{\bullet\bullet}}\right)^\top \mathbf{1}_n = \mathbf{b} = D_c \mathbf{1}_p = \begin{bmatrix} x_{\bullet 1} \\ x_{\bullet 2} \\ \vdots \\ x_{\bullet p} \end{bmatrix}$$