

Correspondence Analysis

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Correspondence analysis (CA) is a generalized principal component analysis tailored for the analysis of qualitative data. Originally, CA was created to analyze contingency tables, but CA is so versatile that it is now often used with other data table types as long these tables contain only non-negative numbers.

CA transforms a data table into two sets of *factor scores*: One set for the rows and one set for the columns. The factor scores give the best representation of the similarity structure of the rows and the columns of the original data table. In addition, the factors scores can be plotted as maps, which display the essential information of the original table. In these maps, rows and columns are displayed as points whose coordinates are the factor scores and where the dimensions are called *factors*. Interestingly, in CA the factor scores of the rows and the columns have the same variance and, therefore, both rows and columns can be conveniently represented in one single map. The modern version of corre-

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spendence analysis and its geometric interpretation originated in France in the 1960s and is associated with the French school of “data analysis” (*analyse des données*).

As a technique, CA was often discovered (and, then, re-discovered) and so variations of correspondence analysis can be found under several names such as “*dual-scaling*,” “*optimal scaling*,” or “*reciprocal averaging*.” The multiple identities of correspondence analysis are a consequence of its versatility and large number of properties: It can be defined as the optimal solution for a lot of apparently different problems.

1 Notations

Matrices are denoted with upper case letters typeset in a boldface font, for example \mathbf{X} is a matrix. The elements of a matrix are denoted with a lower case italic font matching the matrix name with indices indicating the row and column positions of the element, for example $x_{i,j}$ is the element located at the i -th row and j -th-column of matrix \mathbf{X} . Vectors are denoted with lower case letters typeset in a boldface font, for example \mathbf{c} is a vector. The elements of a vector are denoted with a lower case italic font matching the vector name with an index indicating the position of the element in the vector, for example c_i is the i -th element of \mathbf{c} . The superscript T applied to a matrix or vector indicates that this matrix or vector is transposed.

2 An Example: How French Writers Punctuate

This example comes from Etienne Brunet who analyzed the way punctuation marks were used by six French writers: Rousseau, Chateaubriand, Hugo, Zola, Proust, and Giraudoux. In his paper, Brunet gave a table

Table 1: The punctuation marks of six French writers (from Brunet, 1989).

	Period	Comma	All the other marks
Rousseau	7836	13112	6026
Chateaubriand	53655	102383	42413
Hugo	115615	184541	59226
Zola	161926	340479	62754
Proust	38177	105101	12670
Giraudoux	46371	58367	14299

recording the number of times each of these writers used three punctuation marks: the period, the comma, and all the other marks (*i.e.*, interrogation mark, exclamation mark, colon, and semi-colon) grouped together. These data are reproduced in Table 1. From these data we can build the original data matrix which is denoted \mathbf{X} . It has $I = 6$ rows and $J = 3$ columns and is equal to

$$\mathbf{X} = \begin{bmatrix} 7836 & 13112 & 6026 \\ 53655 & 102383 & 42413 \\ 115615 & 184541 & 59226 \\ 161926 & 340479 & 62754 \\ 38177 & 105101 & 12670 \\ 46371 & 58367 & 14299 \end{bmatrix}. \quad (1)$$

In the matrix \mathbf{X} , the rows represent the authors and the columns represent the types of punctuation marks. At the intersection of a row and a column, we find the number of a given punctuation mark (represented by the column) used by a given author (represented by the row).

3 Analyzing the rows

Suppose that the focus is on the *authors*, and that we want to derive a map that reveals the similarities and differences in punctuation style between authors. In this map, the authors should be points and the distances between authors will reflect their stylistic proximity. So, on such a map, when two authors are close to each other these two authors punctuate in a similar way and when two authors are far away these two authors punctuate differently.

3.1 A first (bad) idea: doing PCA



Figure 1: PCA analysis of the Punctuation. Centered Data. Aloz is a supplementary element. Even though Aloz punctuates the same way as Zola, Aloz is further away from Zola than from any other author. The first dimension explains 98% of the variance. It reflects mainly the number of punctuation marks produced by the authors.

A first idea is to perform a principal component analysis (PCA) on X whose results are shown in Figure 1. The plot suggests that the data are quite unidimensional. And, in fact, the first component of this analysis explains 98% of the total inertia (a quantity akin to variance) of the data. How to interpret this component? It seems related to the *number* of punctuation marks produced by each author. This interpretation can be tested by creating a fictitious alias for Zola. To do so: Suppose that, unbeknown to most historians of French literature, Zola wrote a small novel under the (rather transparent) pseudonym of Aloz. In this novel,

he kept his usual way of punctuating, but because this is a short novel, he obviously produced a smaller number of punctuation marks than he did in his complete *œuvre*. Here is the (row) vector recording the number of occurrences of the punctuation marks for Aloz:

$$\left[2699 \ 5675 \ 1046 \right] . \quad (2)$$

For ease of comparison, Zola's row vector is reproduced here:

$$\left[161926 \ 340479 \ 62754 \right] . \quad (3)$$

So, Aloz and Zola have the same punctuation style and differ only in their prolixity. A good analysis should reveal such a similarity of style, but as Figure 1 shows, PCA fails. In this figure, we have projected Aloz (as a supplementary element) in the analysis of the authors and Aloz is, in fact, further away from Zola than from any other author. This example shows that using PCA to analyze the *style* of the authors is not a good idea because PCA is mainly sensitive to the total *number* of punctuation marks produced rather than to *how* punctuation is used. But, the style of the authors, is, in fact, expressed by the *relative frequencies* of their use of the punctuation marks. This suggests that the data matrix should be transformed such that each author is described by the *proportion* of the usage of the punctuation marks rather than by the *number* of punctuation marks used. With this transformation, each row of the data matrix is now called a *row profile*: The entries of a row profile are all non-negative and they sum to one. The transformed data matrix storing the row profiles is called (not-surprisingly) a *row profile matrix*. In order to obtain the row profiles, we divide each row by its sum. The matrix of row profiles

is denoted \mathbf{R} . It is computed as:

$$\mathbf{R} = \text{diag} \left\{ \mathbf{X} \mathbf{1}_{J \times 1} \right\}^{-1} \mathbf{X} = \begin{bmatrix} .2905 & .4861 & .2234 \\ .2704 & .5159 & .2137 \\ .3217 & .5135 & .1648 \\ .2865 & .6024 & .1110 \\ .2448 & .6739 & .0812 \\ .3896 & .4903 & .1201 \end{bmatrix} \quad (4)$$

(where the diag operator transforms a vector into a diagonal matrix with the elements of this vector on the diagonal, and where $\mathbf{1}_{J \times 1}$ is a J by 1 vector of ones).

A convenient way to evaluate the differences between writers is to compare these writers to the “average writer”—A writer who would use each punctuation mark according to its proportion in the sample. The profile of this average writer is called the (row) *barycenter* (also called *centroid*, *center of mass*, or *center of gravity*) of the data matrix. Here, the barycenter of \mathbf{R} is a vector with $J = 3$ elements, it is denoted \mathbf{c} , and computed as

$$\mathbf{c}^T = \underbrace{\left(\mathbf{1}_{1 \times I} \times \mathbf{X} \times \mathbf{1}_{J \times 1} \right)^{-1}}_{\text{Inverse of the total of X}} \times \underbrace{\mathbf{1}_{1 \times I} \mathbf{X}}_{\text{Total of the columns of X}} = \left[.2973 \ .5642 \ .1385 \right]. \quad (5)$$

If all authors punctuate the same way, they all punctuate like the average writer, and, therefore, in order to study the differences between authors, we need to analyze the matrix of *deviations* to the average writer. This matrix of deviations is denoted as \mathbf{Y} and it is computed as:

$$\mathbf{Y} = \mathbf{R} - \left(\mathbf{1}_{I \times 1} \times \mathbf{c}^T \right) = \begin{bmatrix} -.0068 & -.0781 & .0849 \\ -.0269 & -.0483 & .0752 \\ .0244 & -.0507 & .0263 \\ -.0107 & .0382 & -.0275 \\ -.0525 & .1097 & -.0573 \\ .0923 & -.0739 & -.0184 \end{bmatrix}. \quad (6)$$

3.2 Masses (rows) and weights (columns)

In correspondence analysis, we assign a *mass* to each row and a weight to each column. The mass of each row reflects its importance in the sample. In other words, the mass of each row is the proportion of this row in the total of the table. The masses of the rows are stored in a vector denoted \mathbf{m} , which is computed as

$$\mathbf{m} = \underbrace{\left(\mathbf{1}_{1 \times I} \times \mathbf{X} \times \mathbf{1}_{J \times 1} \right)^{-1}}_{\text{Inverse of the total of X}} \times \underbrace{\mathbf{X} \mathbf{1}_{J \times 1}}_{\text{Total of the rows of X}} = \left[.0189 \ .1393 \ .2522 \ .3966 \ .1094 \ .0835 \right]^T. \quad (7)$$

From the vector \mathbf{m} we define the diagonal matrix of masses as $\mathbf{D}_m = \text{diag} \{ \mathbf{m} \}$.

The weight of each column expresses its importance for *discriminating* between the authors. So, the weight of a column reflects the information this columns provides to the identification of a given row. Here, the idea is that the information provided by a column is inversely proportional to its frequency, which, itself, is equal to the value of this column component of the barycenter. Therefore the column weights are computed as the inverse of the values of the barycenter. Specifically, if we

denote by \mathbf{w} the J by 1 weight vector for the columns, we have:

$$\mathbf{w} = [w_j] = [c_j^{-1}] . \quad (8)$$

For our example, we obtain:

$$\mathbf{w} = [w_j] = [c_j^{-1}] = \begin{bmatrix} \frac{1}{.2973} \\ \frac{1}{.5642} \\ \frac{1}{.1385} \end{bmatrix} = \begin{bmatrix} 3.3641 \\ 1.7724 \\ 7.2190 \end{bmatrix} . \quad (9)$$

From vector \mathbf{w} , we define the matrix of column weights as

$$\mathbf{D}_c^{-1} = \text{diag} \{ \mathbf{w} \} = \text{diag} \left\{ [c_j^{-1}] \right\} = \begin{bmatrix} 3.3641 & 0 & 0 \\ 0 & 1.7724 & 0 \\ 0 & 0 & 7.2190 \end{bmatrix} . \quad (10)$$

3.3 Generalized Singular Value Decomposition of \mathbf{Y}

With all these notations defined, correspondence analysis boils down to a *generalized singular value decomposition* (GSVD) problem. Specifically, matrix \mathbf{Y} is decomposed using the GSVD under the constraints imposed by the matrices \mathbf{D}_m (masses for the rows) and \mathbf{D}_c^{-1} (weights for the columns):

$$\mathbf{Y} = \mathbf{P}\mathbf{\Delta}\mathbf{Q}^T \quad \text{with:} \quad \mathbf{P}^T\mathbf{D}_m\mathbf{P} = \mathbf{Q}^T\mathbf{D}_c^{-1}\mathbf{Q} = \mathbf{I} , \quad (11)$$

where \mathbf{P} is the matrix of the right singular vectors, \mathbf{Q} is the matrix of the left singular vectors, and $\mathbf{\Delta}$ is the diagonal matrix of the eigenvalues (in the GSVD framework, \mathbf{D}_m and \mathbf{D}_c^{-1} are also called *metric matrices*, or

simply *metrics*). From this GSVD, we get:

$$\mathbf{Y} = \underbrace{\begin{bmatrix} 1.7962 & 0.9919 \\ 1.4198 & 1.4340 \\ 0.7739 & -0.3978 \\ -0.6878 & 0.0223 \\ -1.6801 & 0.8450 \\ 0.3561 & -2.6275 \end{bmatrix}}_{\mathbf{P}} \times \underbrace{\begin{bmatrix} .1335 & 0 \\ 0 & .0747 \end{bmatrix}}_{\mathbf{\Delta}} \times \underbrace{\begin{bmatrix} 0.1090 & -0.4114 & 0.3024 \\ -0.4439 & 0.2769 & 0.1670 \end{bmatrix}}_{\mathbf{Q}^T}. \quad (12)$$

The rows of matrix \mathbf{X} are now represented by their *factor scores* (which are the projections of the observations onto the singular vectors of \mathbf{X}). The row factor scores are stored in an $I = 3$ by $L = 2$ (L stands for the number of non-zero singular values) matrix denoted \mathbf{F} , which is obtained as

$$\mathbf{F} = \mathbf{P}\mathbf{\Delta} = \begin{bmatrix} 0.2398 & 0.0741 \\ 0.1895 & 0.1071 \\ 0.1033 & -0.0297 \\ -0.0918 & 0.0017 \\ -0.2243 & 0.0631 \\ 0.0475 & -0.1963 \end{bmatrix}. \quad (13)$$

The variance of the factor scores for a given dimension is equal to the squared singular value of this dimension (note that the variance of the observations is computed taking into account their masses). Or, equivalently, we say that the variance of the factor scores of one dimension is equal to the eigenvalue of this dimension (*i.e.*, the eigenvalue is the square of the singular value). This can be checked as follows:

$$\mathbf{F}^T \mathbf{D}_m \mathbf{F} = \mathbf{\Delta} \mathbf{P}^T \mathbf{D}_m \mathbf{P} \mathbf{\Delta} = \mathbf{\Delta}^2 = \mathbf{\Lambda} = \begin{bmatrix} 0.1335^2 & 0 \\ 0 & 0.0747^2 \end{bmatrix} = \begin{bmatrix} 0.0178 & 0 \\ 0 & 0.0056 \end{bmatrix}. \quad (14)$$

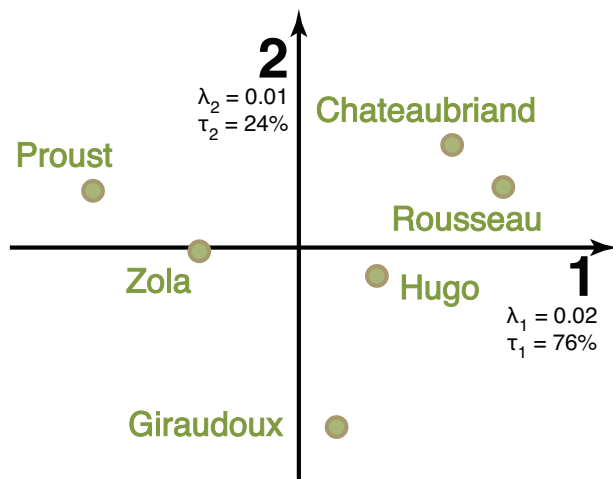


Figure 2: Plot of the correspondence analysis of the rows of matrix \mathbf{X} . The first two sets of factors of the analysis for the rows are plotted (i.e., this is the matrix \mathbf{F}). Each point represents an author. The variance of each set of factor scores is equal to its eigenvalue.

We can display the results by plotting the factor scores as a map where each point represents a row of the matrix \mathbf{X} (i.e., each point represents an author). This is done in Figure 2. On this map, the first dimension seems to be related to time (the rightmost authors are earlier authors, the leftmost authors are more recent), with the exception of Giraudoux who is a very recent author. The second dimension singularizes Giraudoux. These factors will be easier to understand after we have analyzed the columns. This can be done by analyzing the matrix \mathbf{X}^T . Equivalently this can be done by doing what is called the *dual analysis*.

4 Geometry of Correspondence Analysis

CA has a simple geometric interpretation. For example, when a row profile is interpreted as a vector, it can be represented as a point in a multidimensional space. This way, a row profile with values for J variables is represented as a point in a J -dimensional space but because the sum of a profile is equal to one, row profiles are, in fact points in a $(J - 1)$

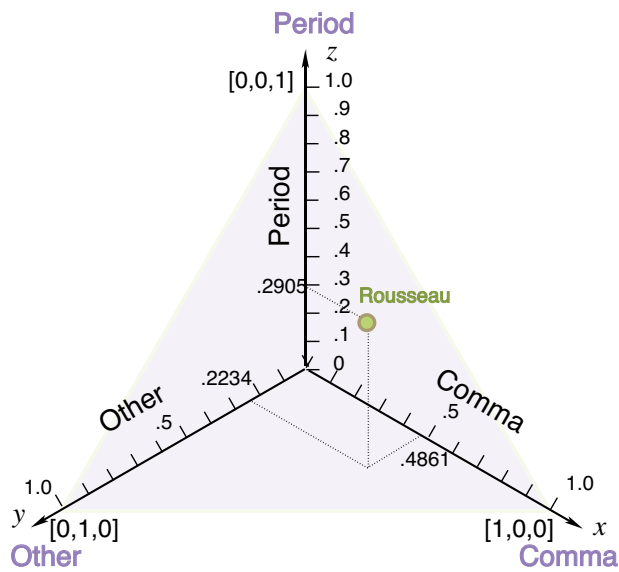


Figure 3: In three dimensions, the simplex is a 2-dimensional triangle whose vertices are the vectors $[1\ 0\ 0]$, $[0\ 1\ 0]$ and $[0\ 0\ 1]$. The point describing Rousseau (with coordinates $[\.2905\ .4861\ .2234]$) is also plotted.

dimensional space. Also, because the components of a row profile take value in the interval $[0\ 1]$, the points representing these row-profiles can only lay in the subspace whose “extreme points” (i.e., vertices) have one component equal to one and all other components equal to zero. This subspace is called a *simplex*. For example, Figure 3 shows the 2-dimensional simplex corresponding to the subspace of all possible row profiles with three components. As an illustration, the point describing Rousseau (with coordinates equal to $[\.2905\ .4861\ .2234]$) is also plotted. For this particular example, the simplex is an equilateral triangle and, so the three dimensional row profiles can conveniently be represented as points on this triangle as illustrated in Figure 4a which shows the simplex of Figure 3 in two dimensions. Figure 4b shows all six authors and their barycenter.

The weights of the columns, which are used as constrains in the GSVD, have also a straightforward geometric interpretation. As illus-

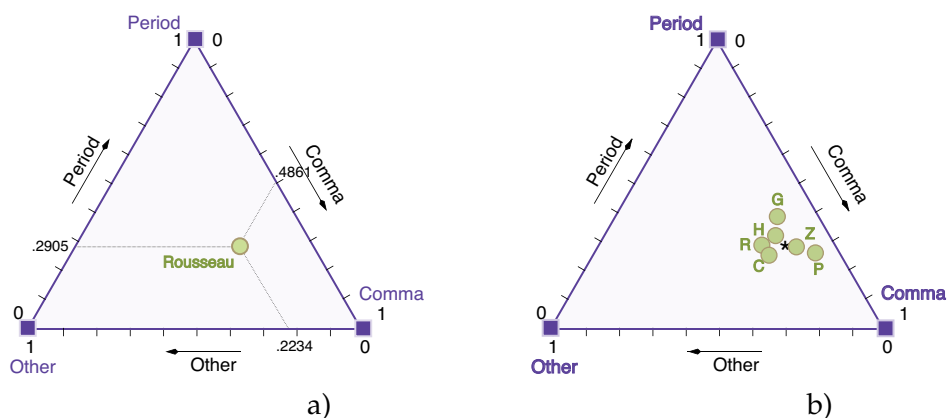


Figure 4: The simplex as a triangle. a) With Rousseau (compare with Figure 3) b) With all six authors and their barycenter.

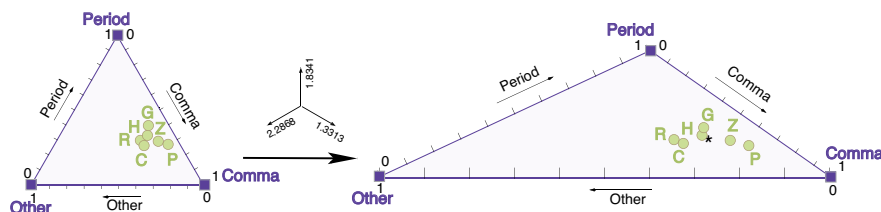


Figure 5: Geometric interpretation of the column weights. Each side of the simplex is stretched by a factor equal to the square root of the weights.

trated in Figure 5, each side of the simplex is stretched by a quantity equal to the square root of the dimension it represents (we use the square root because we are interested in *squared* distances but not in squared weights, so, using the square root of the weights ensures that the *squared* distances between authors will take into account the weights rather than the squared weights).

To find the factors, the masses of the rows are taken into account. Specifically, the first factor is computed such that it gives the maximum possible value of the sum of the masses times the squared projections of the authors points (*i.e.*, the projections have the largest possible variance). The second factor is constrained to be orthogonal (taking into

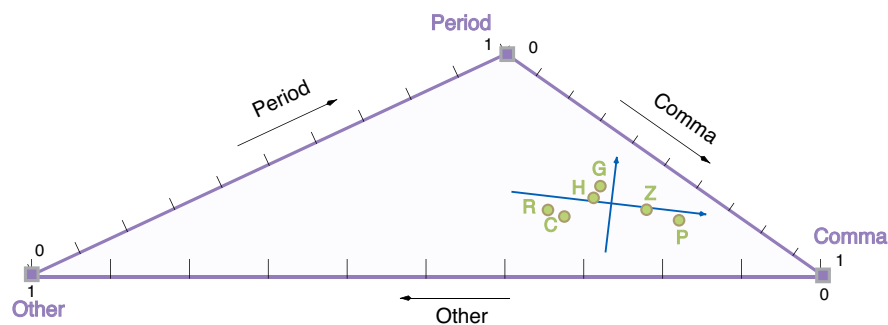


Figure 6: Correspondence analysis: The “stretched simplex” along with the factorial axes. The projections of the Authors’ points onto the factorial axes give the factor scores.

account the masses) to the first one and have the largest variance for the projections. The remaining factors are computed with similar constraints. Figure 6 shows the stretched simplex, the author points, and the two factors (note that the origin of the factors is the barycenter of the authors).

The “stretched simplex” shows the whole space of the possible profiles. Figure 6 shows that the authors occupy a small portion of this whole space—A pattern which reveals that the authors do not vary much in the way they punctuate. Also, the stretched simplex represents the columns as the vertices of the simplex: The columns are represented as row profiles with the column component being equal to one and all the other components being equal to zero. This representation is called an *asymmetric* representation because the rows always have a dispersion smaller than (or equal to) the columns.

5 Distance, Inertia, Chi-square, and CA

5.1 Chi-squared distances

In CA, the Euclidean distance in the “stretched simplex” is equivalent to a weighted distance in the original space. For reasons—that will be clear later—this distance is called the χ^2 -distance. The χ^2 -distance between two row profiles i and i' can be computed from the factor scores as

$$d_{i,i'}^2 = \sum_{\ell}^L (f_{i,\ell} - f_{i',\ell})^2 \quad (15)$$

or from the row-profiles as

$$d_{i,i'}^2 = \sum_j^J w_j (r_{i,j} - r_{i',j})^2. \quad (16)$$

5.2 Inertia

The variability of the row profiles relative to their barycenter is measured by a quantity—akin to a variance—called *inertia* and denoted \mathcal{I} . The inertia of the rows to their barycenter is computed as the weighed sum of the squared distances of the rows to their barycenter. We denote by $d_{c,i}^2$ the (squared) distance of the i -th row to the barycenter, it is computed as

$$d_{c,i}^2 = \sum_j^J w_j (r_{i,j} - c_j)^2 = \sum_{\ell}^L f_{i,\ell}^2 \quad (17)$$

where L is the number of factors extracted by the CA of the table, [this number is smaller or equal to $\min(I, J) - 1$]. The inertia of the rows to

their barycenter is then computed as

$$\mathcal{I} = \sum_i^I m_i d_{c,j}^2. \quad (18)$$

The inertia can also be expressed as the sum of the eigenvalues (see Equation 14):

$$\mathcal{I} = \sum_{\ell}^L \lambda_{\ell}. \quad (19)$$

This shows that in CA, each factor extracts a portion of the inertia, with the first factor extracting the largest portion, the second factor extracting the largest portion left of the inertia, etc.

5.3 Inertia and the Chi-squared test

Interestingly, the inertia in CA is closely related to the chi-square test which is traditionally performed on a contingency table in order to test the independence of the rows and the columns of the table. Under independence, the frequency of each cell of the table is proportional to the product of its row and column marginal probabilities. So, if we denote by $x_{+,+}$ the grand total of matrix \mathbf{X} , the expected frequency of the cell at the i -th row and j -th column is denoted $E_{i,j}$ and computed as:

$$E_{i,j} = m_i c_j x_{+,+}. \quad (20)$$

The chi-squared test statistic, denoted χ^2 is computed as the sum of the squared difference between the actual values and the corresponding expected values weighted by the expected values:

$$\chi^2 = \sum_{i,j} \frac{(x_{i,j} - E_{i,j})^2}{E_{i,j}}. \quad (21)$$

When rows and columns are independent, χ^2 follows a chi-squared distribution with $(I - 1)(J - 1)$ degrees of freedom. Therefore, χ^2 can be used to evaluate the likelihood of the row and columns independence hypothesis. The statistics χ^2 can be rewritten to show its close relationship with the inertia of CA, namely:

$$\chi^2 = \mathcal{I} \times x_{+,+} = \varphi^2 \times x_{+,+}, \quad (22)$$

with $\varphi^2 = \mathcal{I}$ being a coefficient of effect size associated to chi^2 (this index takes values between 0 and $\min(I, J) - 1$). Equation 22 shows that CA decomposes—in orthogonal components—the pattern of deviations to independence.

6 Dual Analysis: the Column Space

In a contingency table, the rows and the columns of the table play a similar role, and therefore the analysis that was performed on the rows can also be performed on the columns by exchanging the role of the rows and the columns. This is illustrated by the analysis of the columns of matrix \mathbf{X} , or equivalently by the rows of the transposed matrix \mathbf{X}^T . The matrix of column profiles for \mathbf{X}^T is called \mathbf{O} (like cOolumn), and is computed as

$$\mathbf{O} = \text{diag} \left\{ \mathbf{X}^T \mathbf{1}_{I \times 1} \right\}^{-1} \mathbf{X}^T \quad (23)$$

The matrix of the column deviations to their barycenter is called \mathbf{Z} and it is computed as:

$$\mathbf{Z} = \mathbf{O} - \left(\mathbf{1}_{I \times 1} \times \mathbf{m}^T \right) = \begin{bmatrix} -.0004 & -.0126 & .0207 & -.0143 & -.0193 & .0259 \\ -.0026 & -.0119 & -.0227 & .0269 & .0213 & -.0109 \\ .0116 & .0756 & .0478 & -.0787 & -.0453 & -.0111 \end{bmatrix}. \quad (24)$$

Weights and masses of the column analysis are the inverse of their equivalent for the row analysis. This implies that the punctuation marks factor scores are obtained from the GSVD with the constraints imposed by the two metric matrices \mathbf{D}_c (masses for the columns) and \mathbf{D}_m^{-1} (weights for the rows, compare with Equation 11):

$$\mathbf{Z} = \mathbf{U}\mathbf{\Delta}\mathbf{V}^T \quad \text{with:} \quad \mathbf{U}^T \mathbf{D}_c \mathbf{U} = \mathbf{V}^T \mathbf{D}_m^{-1} \mathbf{V} = \mathbf{I}. \quad (25)$$

This gives:

$$\mathbf{Z} = \underbrace{\begin{bmatrix} 0.3666 & -1.4932 \\ -0.7291 & 0.4907 \\ 2.1830 & 1.2056 \end{bmatrix}}_{\mathbf{U}} \times \underbrace{\begin{bmatrix} .1335 & 0 \\ 0 & .0747 \end{bmatrix}}_{\mathbf{\Delta}} \times \underbrace{\begin{bmatrix} 0.0340 & 0.1977 & 0.1952 & -0.2728 & -0.1839 & 0.0298 \\ 0.0188 & 0.1997 & -0.1003 & 0.0089 & 0.0925 & -0.2195 \end{bmatrix}}_{\mathbf{V}^T}. \quad (26)$$

The factor scores for the punctuation marks are stored in a $J = 3 \times L = 2$ matrix denoted \mathbf{G} which is computed in the same way \mathbf{F} was

computed (see Equation 13). Specifically, \mathbf{G} is computed as:

$$\mathbf{G} = \mathbf{U}\mathbf{\Delta} = \begin{bmatrix} 0.0489 & -0.1115 \\ -0.0973 & 0.0367 \\ 0.2914 & 0.0901 \end{bmatrix}. \quad (27)$$

6.1 Transition formula: from the rows to the columns and back

A Comparison of Equation 26 and Equation 12, shows that the singular values are the same for both the row and the column analyses. This means that the inertia extracted by each factor (*i.e.*, the eigenvalue associated to this factor, which is also the square of the singular value) is the same for both analyses. Because the variance extracted by the factors can be added, to obtain the total inertia of the data table, this also means that each analysis is decomposing the same inertia which, here, is equal to:

$$\mathcal{I} = .1335^2 + .747^2 = .0178 + .0056 = 0.0234. \quad (28)$$

Also, the generalized singular decomposition of one set (say the columns) can be obtained from the other set (say the rows). For example the generalized singular vectors of the analysis of the columns can be computed directly from the analysis from the rows as

$$\mathbf{U} = \mathbf{D}_c^{-1}\mathbf{Q}. \quad (29)$$

Combining Equations 29 and 27 shows that the factors for the rows of \mathbf{Z} (*i.e.*, the punctuation marks) can be obtained directly from the singular value decomposition of the authors matrix (*i.e.*, matrix \mathbf{Y}) as

$$\mathbf{G} = \mathbf{U}\mathbf{\Delta} = \mathbf{D}_c^{-1}\mathbf{Q}\mathbf{\Delta}. \quad (30)$$

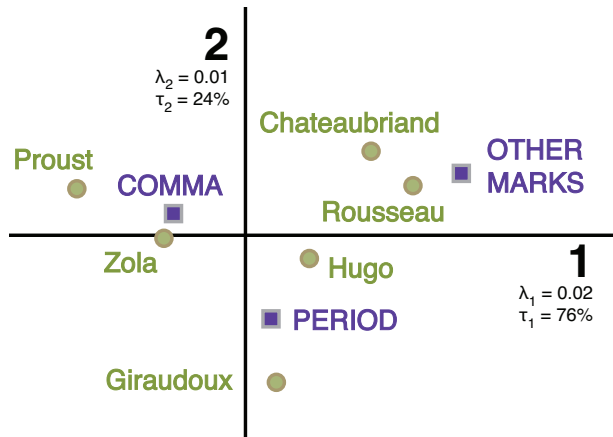


Figure 7: Correspondence analysis of six authors (Rousseau, Chateaubriand, Hugo, Zola, Proust, and Giraudoux) described by the way they used three punctuation marks (Comma, Period, and Others).

As a consequence, we can, in fact, find directly the factor scores of the columns from their profile matrix (*i.e.*, the matrix \mathbf{O}), and from the factor scores of the rows. Specifically, the equation which gives the values of \mathbf{G} from \mathbf{F} is

$$\mathbf{G} = \mathbf{O}\mathbf{F}\mathbf{\Delta}^{-1}, \quad (31)$$

and conversely \mathbf{F} could be obtained from \mathbf{G} as

$$\mathbf{F} = \mathbf{R}\mathbf{G}\mathbf{\Delta}^{-1}. \quad (32)$$

These equations are called “*transition formulas from the rows to the columns*” (and vice versa) or simply the *transition formulas*.

6.2 One single GSVD for CA

Because the factor scores obtained for the rows and the columns have the same variance (*i.e.*, they have the same “scale”), it is possible to plot them in the same space. This is illustrated in Figure 7. The symmetry of the rows and the columns in CA is revealed by the possibility of *directly*

obtaining the factors scores from one single GSVD. Specifically, let \mathbf{N} denote the matrix \mathbf{X} divided by the sum of all its elements. This matrix—sometimes called a *correspondence* matrix—has all elements larger than or equal to zero and their sum equals to one. The factors scores for the rows and the columns are obtained from the following GSVD:

$$\left(\mathbf{N} - \mathbf{m}\mathbf{c}^T\right) = \mathbf{S}\mathbf{\Delta}\mathbf{T}^T \quad \text{with} \quad \mathbf{S}^T\mathbf{D}_m^{-1}\mathbf{S} = \mathbf{T}^T\mathbf{D}_c^{-1}\mathbf{T} = \mathbf{I}. \quad (33)$$

With the decomposition from Equation 33, the factor scores for the rows (\mathbf{F}) and the columns (\mathbf{G}) are obtained respectively as

$$\mathbf{F} = \mathbf{D}_m^{-1}\mathbf{S}\mathbf{\Delta} \quad \text{and} \quad \mathbf{G} = \mathbf{D}_c^{-1}\mathbf{T}\mathbf{\Delta}. \quad (34)$$

7 Supplementary elements

Often in CA we want to know the position in the analysis of rows or columns that were not analyzed. These rows or columns are called illustrative or supplementary rows or columns (or supplementary observations or variables). The appellation “out of sample” observations or variables is also sometimes used. By contrast with these supplementary elements (which are *not* used to compute the factors) the *active* elements are those used to actually compute the factors. Table 2 shows the punctuation data table with four additional columns giving the detail of the “other punctuation marks” (*i.e.*, the exclamation mark, the interrogation mark, the semi-colon, and the colon). These punctuation marks were not analyzed for two reasons: first, these marks are too rare and therefore they would distort the factor space and, second, the “Other” marks comprises all these other marks and therefore to analyze them with “Other” would be redundant. There is also a new author in Table 2: We counted the marks used by a different author, namely Hervé Abdi in the first chapter of

	Active Elements				Supplementary Elements					
	Period	Comma	Other Marks	x_{i+}	$\frac{x_{i+}}{x_{++}}$	Exclamation	Question	Semicolon	Colon	
Rousseau	7836	13112	6026	26974	.0189	413	1240	3401	972	
Chateaubriand	53655	102383	42413	198451	.1393	4669	4595	19354	13795	
Hugo	115615	184541	59226	359382	.2522	19513	9876	22585	7252	
Zola	161926	340479	62754	565159	.3966	24025	10665	18391	9673	
Proust	38117	105101	12670	155948	.1094	2756	2448	3850	3616	
Giraudoux	46371	58367	14229	119037	.0835	5893	5042	1946	1418	
x_{+j}	423580	803983	197388	1424951						
$w^T = \frac{x_{++}}{x_{+j}}$	3.3641	1.7724	7.2190	x_{++}						
$c^T = \frac{x_{+j}}{x_{+++}}$.2973	.5642	.1385							
Abdi (Chapter 1)	216	139	26							

Table 2: Number of punctuation marks used by six major French authors (from Brunet, 1989). The exclamation point, question mark, semicolon, and colon are supplementary columns. Abdi (1994) Chapter 1 is a supplementary row. Notations x_{i+} : sum of the i -th row; x_{+j} : sum of the j -th column; x_{++} : grand total.

Table 3: Factor scores, contributions, and cosines for the Rotas. Negative contributions are shown in *italic*. Abdi (1994) Chapter 1 is a supplementary row.

Axis	λ	%	Rousseau	Chateaubriand	Hugo	Zola	Proust	Giraudoux	Abdi (Chapter 1)
Factor Scores									
1	.0178	76	0.2398	0.1895	0.1033	-0.0918	-0.2243	0.0475	-0.0908
2	.0056	24	0.0741	0.1071	-0.0297	0.0017	0.0631	-0.1963	0.5852
Contributions									
1			<i>0.0611</i>	<i>0.2807</i>	<i>0.1511</i>	<i>0.1876</i>	<i>0.3089</i>	0.0106	-
2			<i>0.0186</i>	<i>0.2864</i>	<i>0.0399</i>	<i>0.0002</i>	<i>0.0781</i>	<i>0.5767</i>	-
Cosines									
1			0.9128	0.7579	0.9236	0.9997	0.9266	0.0554	0.0235
2			0.0872	0.2421	0.0764	0.0003	0.0734	0.9446	0.9765
Squared Distances to Grand Barycenter									
-	-	-	0.0630	0.0474	0.0116	0.0084	0.0543	0.0408	0.3508

Axis	λ	%	Period	Comma	Other Marks	Exclamation	Question	Semicolon	Colon
Factor Scores									
1	.0178	76	-0.0489	0.0973	-0.2914	-0.0596	-0.1991	-0.4695	-0.4008
2	.0056	24	0.1115	-0.0367	-0.0901	0.2318	0.2082	-0.2976	-0.4740
Contributions									
1			<i>0.0399</i>	<i>0.2999</i>	<i>0.6601</i>	-	-	-	-
2			<i>0.6628</i>	<i>0.1359</i>	<i>0.2014</i>	-	-	-	-
Cosines									
1			0.1614	0.8758	0.9128	0.0621	0.4776	0.7133	0.4170
2			0.8386	0.1242	0.0872	0.9379	0.5224	0.2867	0.5830
Squared Distances to Grand Barycenter									
-	-		0.0148	0.0108	0.0930	0.0573	0.0830	0.3090	0.3853

Table 4: Factor scores, contributions, and cosines for the columns. Negative contributions are shown in *italic*. Exclamation mark, question mark, semicolon, and colon are supplementary columns.

his 1994 book called “*Les réseaux de neurones.*” This author was not analyzed because data are available for only one chapter (not his complete work) and also because this author (despite all his stylistic qualities) is not, strictly speaking, a literary author.

The values of the projections on the factors for the supplementary elements are computed from the *transition formula*. Specifically, a supplementary row is projected into the space defined using the transition formula for the active rows (*cf.* Equation 32) and replacing the active row profiles by the supplementary row profiles. So, if we denote by \mathbf{R}_{sup} the matrix of the supplementary row profiles, then \mathbf{F}_{sup} —the matrix of the supplementary row factor scores—is computed as:

$$\mathbf{F}_{\text{sup}} = \mathbf{R}_{\text{sup}} \times \mathbf{G} \times \Delta^{-1} . \quad (35)$$

Table 3 lists the factor scores for the active and supplementary rows. For example, the factor scores of the author Abdi are computed as

$$\mathbf{F}_{\text{sup}} = \mathbf{R}_{\text{sup}} \mathbf{G} \Delta^{-1} = \begin{bmatrix} 0.0908 & -0.5852 \end{bmatrix} . \quad (36)$$

Supplementary columns are projected into the factor space using the transition formula from the active rows (*cf.* Equation 31) and replacing the active column profiles by the supplementary column profiles. So, if we denote by \mathbf{O}_{sup} the supplementary column profile matrix, then \mathbf{G}_{sup} , the matrix of the supplementary column factor scores, is computed as:

$$\mathbf{G}_{\text{sup}} = \mathbf{O}_{\text{sup}} \mathbf{F} \Delta^{-1} . \quad (37)$$

Table 4 lists the factor scores for the active and supplementary elements and Figure 8 displays the supplementary columns (*i.e.*, the “Other” punctuation marks) and the supplementary author (Abdi).

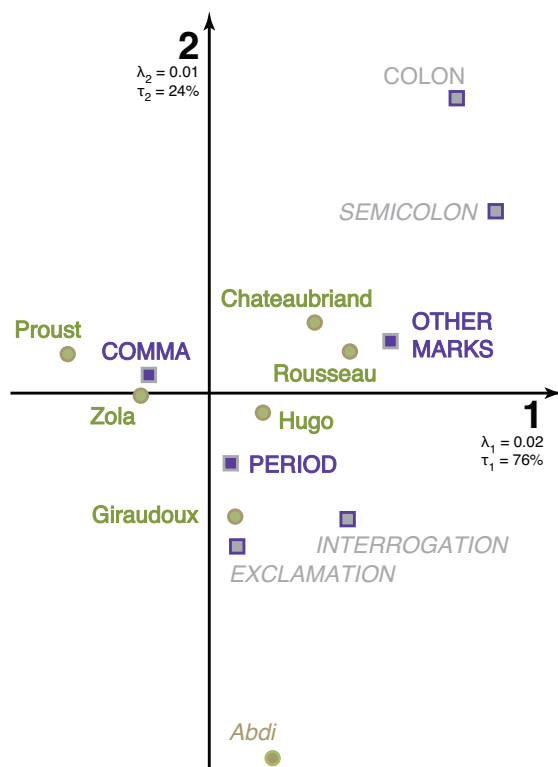


Figure 8: Correspondence analysis of the punctuation of six authors, Comma, Period, and Others are active columns; Rousseau, Chateaubriand, Hugo, Zola, Proust, and Giraudoux are active rows; Colon, Semicolon, Interrogation and Exclamation are supplementary columns; Abdi is a supplementary row.

Figure 8 reveals that the “Other” punctuation mark category comprises two distinct groups: 1) the first group includes “Colon” and “Semi-Colon” which are punctuation marks used more than average by the early authors; whereas 2) the second group includes “interrogation” and “exclamation” marks which are punctuation marks used more than average by Giraudoux (who, as a writer of plays, frequently uses spoken dialogues rich in “interrogation” and “exclamation” marks). Note that because the four supplementary punctuation marks add up to the “Other” punctuation mark, the “Other” mark is the barycenter of the four “Other” marks.

For the supplementary author, Abdi—whose linguistic corpus originates from scientific writing—is plotted far from the origin on the negative side of Dimension 2 because he wrote short sentences (a hallmark of scientific writing) and therefore used more periods and fewer commas and other punctuation marks than the average writer.

8 Little Helpers: Contributions and cosines

Contributions and cosines are coefficients whose goal is to facilitate the interpretation. The contributions identify the important elements for a given factor, whereas the (squared) cosines identify the factors important for a given element. These coefficients express importance as the proportion of something into a total. The contribution is the ratio of the weighted squared projection of an element on a factor to the sum of the weighted projections of all the elements for this factor (which happens to be the eigenvalue of this factor). The squared cosine is the ratio of the squared projection of an element onto a factor to the sum of the projections of this element on all the factors (which happens to be the squared distance from this point to the barycenter). Contributions and squared cosines, being proportions, take values between 0 and 1.

The squared *cosines*, denoted h , between row i and factor ℓ (respectively, between column j and factor ℓ) are obtained as:

$$h_{i,\ell} = \frac{f_{i,\ell}^2}{\sum_{\ell} f_{i,\ell}^2} = \frac{f_{i,\ell}^2}{d_{c,i}^2} \quad \text{and} \quad h_{j,\ell} = \frac{g_{j,\ell}^2}{\sum_{\ell} f_{j,\ell}^2} = \frac{g_{j,\ell}^2}{d_{r,j}^2}. \quad (38)$$

Squared cosines help locating the factors important for a given observation. The *contributions*, denoted b , of row i to factor ℓ and of column j to

factor ℓ are obtained respectively as:

$$b_{i,\ell} = \frac{m_i f_{i,\ell}^2}{\sum_i m_i f_{i,\ell}^2} = \frac{m_i f_{i,\ell}^2}{\lambda_\ell} \quad \text{and} \quad b_{j,\ell} = \frac{c_j g_{j,\ell}^2}{\sum_j c_j g_{j,\ell}^2} = \frac{c_j g_{j,\ell}^2}{\lambda_\ell}. \quad (39)$$

Contributions help locating the observations important for a given factor. An often used rule of thumb is to consider that the important contributions are those larger than the average contribution, which is equal to one divided by the number of elements (*i.e.*, $\frac{1}{I}$ for the rows and $\frac{1}{J}$ for the columns). A dimension is then interpreted by opposing the positive elements with large contributions to the negative elements with large contributions. Cosines and contributions for the punctuation example are given in Tables 3 and 4.

9 Inferences for Correspondence Analysis

CA was first developed as a *descriptive* multivariate method, but recently it also started to incorporate some inferential aspects. When CA analyzes a contingency table, the inertia decomposed by CA is proportional to χ^2 (see Equation 22) and, therefore an independence χ^2 statistic with $(I - 1)(J - 1)$ degrees of freedom implements an *omnibus* test for CA. For our example, the value of the independence χ^2 is equal to

$$\begin{aligned} \chi^2 &= \mathcal{I} \times x_{+,+} = 1,424,951 \times (.0178 + .0056) \\ &= 1,424,951 \times 0.0234 = 33,340.15. \end{aligned} \quad (40)$$

When compared to a χ^2 distribution with $(6 - 1)(3 - 1) = 10$ degrees of freedom, the value of the χ^2 statistic equal to 33,240.15 indicates that the result is highly significant and that we can confidently conclude that the authors *do* differ in how they punctuate (even though the differences

are quite small as indicated by a value of $\varphi^2 = 0.0234$ whose maximum possible value would be equal to 2).

Edmond Malinvaud and Gilbert Saporta extended this χ^2 based statistical approach to test the significativity of the dimensions extracted by CA. To do so, they defined a χ^2 -like statistics denoted Q'_ℓ [with ℓ taking values between 0 and $L = \min(I, J) - 2$] computed as

$$Q'_\ell = x_{+,+} \left(\sum_{k=\ell+1}^L \lambda_k \right). \quad (41)$$

Under the same statistical assumptions as the independence χ^2 , the statistics Q'_ℓ follows a χ^2 distribution with $\nu = (I - \ell - 1)(J - \ell - 1)$. When $\ell = 0$, $Q'_0 = \chi^2 = 33,340.15$ with a number of degrees of freedom equal to $(I - 1)(J - 1) = 10$ (see Equation 22) and is identical to the χ^2 statistics from Equation 40. The significativity of Dimension 1, is tested by computing Q'_1 as:

$$\begin{aligned} Q'_1 &= x_{+,+} \left(\sum_{k=1+1}^2 \lambda_k \right) = x_{+,+} \left(\sum_{k=2}^2 \lambda_k \right) \\ &= x_{+,+} \times \lambda_2 = 1,424,951 \times .0056 = 7,949.57 \end{aligned} \quad (42)$$

a value distributed as χ^2 with $\nu = (I - \ell - 1)(J - \ell - 1) = 4 \times 1 = 4$ degrees of freedom. Here again, the large value of χ^2 indicates that Dimension 1 is highly significant (even though it describes a small effect as indicated by the eigenvalue of $\lambda_1 = .0178$). Note that Q'_ℓ is not defined for the last dimension of CA.

When the data matrix to be analyzed is not a true contingency table, the Q'_ℓ statistic is not distributed as χ^2 and therefore this distribution cannot be derived analytically; So, here, appropriate permutation

or Monte-Carlo cross-validation approaches need to be implemented to derive the sampling distribution of Q'_ℓ .

10 Multiple correspondence analysis

CA works with a contingency table which is equivalent to the analysis of two nominal variables (*i.e.*, one for the rows and one for the columns). Multiple correspondence analysis (MCA) is an extension of CA which allows the analysis of the pattern of relationship among several nominal variables. MCA is used to analyze a set of observations described by a set of nominal variables. Each nominal variable comprises several levels, and each of these levels is coded as a binary variable. For example gender (F *vs.* M) is a nominal variable with two levels. The pattern for a male respondent will be coded as [0 1] and as [1 0] for a female. The complete data table is composed of binary columns with one and only one column taking the value "1" per nominal variable.

MCA can also accommodate quantitative variables by recoding them as "bins." For example, a score with a range of -5 to $+5$ could be recoded as a nominal variable with three levels: less than 0, equal to 0, or more than 0. With this schema, a value of 3 will be expressed by the pattern [0 0 1]. The coding schema of MCA implies that each row has the same total, which for CA implies that each row has the same *mass*.

Essentially, MCA is computed by using a CA program on the data table. It can be shown that the binary coding scheme used in MCA creates artificial factors and therefore artificially reduces the inertia explained by the first factors of the analysis. A solution to this problem is to correct the eigenvalues obtained from the CA program; note that recent statistical packages are likely to automatically implement this correction.

Hervé Abdi and Lynne J. Williams

See also

Barycentric discriminant analysis, Canonical correlation analysis, categorical variables, Chi-square test, Data mining, Predictive Discriminant analysis, Exploratory data analysis, Exploratory factor analysis, Guttman scaling, Matrix algebra, Principal component analysis.

Further readings

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