

Multivariate Data Analysis: The French Way

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Abstract: This paper presents exploratory techniques for multivariate data, many of them well known to French statisticians and ecologists, but few well understood in North American culture.

We present the general framework of duality diagrams which encompasses discriminant analysis, correspondence analysis and principal components and we show how this framework can be generalized to the regression of graphs on covariates.

1. Motivation

David Freedman is well known for his interest in multivariate projections [5] and his skepticism with regards to model-based multivariate inference, in particular in cases where the number of variables and observations are of the same order (see Freedman and Peters[12, 13]).

Brought up in a completely foreign culture, I would like to share an alien approach to some modern multivariate statistics that is not well known in North American statistical culture. I have written the paper ‘*the French way*’ with theorems and abstract formulation in the beginning and examples in the latter sections, Americans are welcome to skip ahead to the motivating examples.

Some French statisticians, fed Bourbakist mathematics and category theory in the 60’s and 70’s as all mathematicians were in France at the time, suffered from abstraction envy. Having completely rejected the probabilistic enterprise as useless for practical reasons, they composed their own abstract framework for talking about data in a geometrical context. I will explain the framework known as the duality diagram developed by Cazes, Caillès, Pages, Escoufier and their followers. I will try to show how aspects of the general framework are still useful today and how much every idea from Benzecri’s correspondence analysis to Escoufier’s conjoint analysis has been rediscovered many times. Section 2.1 sets out the abstract picture. Section 2.2-2.6 treat extensions of classical multivariate techniques; principal components analysis, instrument variables, canonical correlation analysis, discriminant analysis, correspondence analysis from this unified view. Section 3 shows how the methods apply to the analysis of network data.

2. The Duality Diagram

Established by the French school of “Analyse des Données” in the early 1970’s this approach was only published in a few texts, [1] and technical reports [9], none of

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which were translated into English. My PhD advisor, Yves Escoufier [8][10] publicized the method to biologists and ecologists presenting a formulation based on his RV-coefficient, that I will develop below. The first software implementation of duality based methods described here were done in LEAS (1984) a Pascal program written for Apple II computers. The most recent implementation is the **R** package `ade-4` (see A for a review of various implementations of the methods described here).

2.1. Notation

The data are p variables measures on n observations. They are recorded in a matrix X with n rows (the observations) and p columns (the variables). D_n is an $n \times n$ matrix of weights on the "observations", most often diagonal. We will also use a "neighborhood" relation (thought of as a metric on the observations) defined by taking a symmetric definite positive matrix Q . For example, to standardize the variables Q can be chosen as

$$Q = \begin{pmatrix} \frac{1}{\sigma_1^2} & 0 & 0 & 0 & \dots \\ 0 & \frac{1}{\sigma_2^2} & 0 & 0 & \dots \\ 0 & 0 & \frac{1}{\sigma_3^2} & 0 & \dots \\ \dots & \dots & \dots & 0 & \frac{1}{\sigma_p^2} \end{pmatrix}.$$

These three matrices form the essential "triplet" $(\mathbf{X}, \mathbf{Q}, \mathbf{D})$ defining a multivariate data analysis. As the approach here is geometrical, it is important to see that Q and D define geometries or inner products in \mathbb{R}^p and \mathbb{R}^n respectively through

$$\begin{aligned} x^t Q y &= \langle x, y \rangle_Q & x, y &\in \mathbb{R}^p \\ x^t D y &= \langle x, y \rangle_D & x, y &\in \mathbb{R}^n \end{aligned}$$

From these definitions we see there is a close relation between this approach and kernel based methods, for more details see [24]. Q can be seen as a linear function from \mathbb{R}^p to $\mathbb{R}^{p*} = \mathcal{L}(\mathbb{R}^p)$, the space of scalar linear functions on \mathbb{R}^p . D can be seen as a linear function from \mathbb{R}^n to $\mathbb{R}^{n*} = \mathcal{L}(\mathbb{R}^n)$. Escoufier[8] proposed to associate to a data set an operator from the space of observations \mathbb{R}^p into the dual of the space of variables \mathbb{R}^{n*} . This is summarized in the following diagram [1] which is made commutative by defining V and W as $X^t D X$ and $X Q X^t$ respectively.

$$\begin{array}{ccc} \mathbb{R}^{p*} & \xrightarrow{\quad X \quad} & \mathbb{R}^n \\ Q \uparrow & & \downarrow D \\ \mathbb{R}^p & \xleftarrow{\quad X^t \quad} & \mathbb{R}^{n*} \\ & & \uparrow W \\ & & \downarrow V \end{array}$$

This is known as the duality diagram because knowledge of the eigendecomposition of $X^t D X Q = V Q$ leads to that of the dual operator $X Q X^t D$. The main consequence is an easy transition between principal components and principal axes

as we will see in the next section. The terms duality diagram or triplet are often used interchangeably.

Notes:

1. Suppose we have data and inner products defined by Q and D :

$$(x, y) \in \mathbb{R}^p \times \mathbb{R}^p \mapsto x^t Q y = \langle x, y \rangle_Q \in \mathbb{R}$$

$$(x, y) \in \mathbb{R}^n \times \mathbb{R}^n \mapsto x^t D x = \langle x, y \rangle_D \in \mathbb{R}$$

We say an operator O is B -symmetric if $\langle x, O y \rangle_B = \langle O x, y \rangle_B$, or equivalently $BO = O^t B$. The duality diagram is equivalent to three matrices $(\mathbf{X}, \mathbf{Q}, \mathbf{D})$ such that X is $n \times p$ and Q and D are symmetric matrices of the right size (Q is $p \times p$ and D is $n \times n$). The operators defined as $X Q^t X D = W D$ and $X^t D X Q = V Q$ are called the characteristic operators of the diagram [8], in particular $V Q$ is Q -symmetric and $W D$ is D -symmetric.

2. $V = X^t D X$ will be the variance-covariance matrix if X is centered with regards to $(D, X^t D \mathbf{1}_n = 0)$.
3. There is an important symmetry between the rows and columns of X in the diagram, and one can imagine situations where the role of observation or variable is not uniquely defined. For instance in microarray studies the genes can be considered either as variables or observations. This makes sense in many contemporary situations which evade the more classical notion of n observations seen as a random sample of a population. It is certainly not the case that the 30,000 probes are a sample of genes since these probes try to be an exhaustive set.

2.1.1. Properties of the Diagram

Here are some of the properties that prove useful in various settings:

- Rank of the diagram $X, X^t, V Q$ and $W D$ all have same rank which will usually be smaller than both n and p .
- For Q and D symmetric matrices $V Q$ and $W D$ are diagonalisable and have the same eigenvalues. We note them in decreasing order

$$\lambda_1 \geq \lambda_2 \geq \lambda_3 \geq \dots \geq \lambda_r \geq 0 \geq \dots \geq 0.$$

- Eigendecomposition of the Diagram:
 $V Q$ is Q symmetric, thus we can find Z such that

$$V Q Z = Z \Lambda, Z^t Q Z = \mathbb{I}_p, \text{ where } \Lambda = \text{diag}(\lambda_1, \lambda_2, \dots, \lambda_p) \quad (2.1)$$

In practical computations, we start by finding the Cholesky decompositions of Q and D , which exist as long as these matrices are symmetric and positive definite, call these $H^t H = Q$ and $K^t K = D$. Here H and K are triangular. Then we use the singular value decomposition of $K X H$:

$$K X H = U S T^t, \quad \text{with } T^t T = \mathbb{I}_p, U^t U = \mathbb{I}_n, S \text{ diagonal}$$

. Then $Z = (H^{-1})^t T$ satisfies (2.1) with $\Lambda = S^2$. The renormalized columns of $Z, A = S Z$ are called the principal axes and satisfy:

$$A^t Q A = \Lambda$$

Similarly, we can define $L = K^{-1}U$ that satisfies

$$WDL = LA, L^t DL = \mathbb{I}_n, \text{ where } \Lambda = \text{diag}(\lambda_1, \lambda_2, \dots, \lambda_r, 0, \dots, 0). \quad (2.2)$$

$C = LS$ is usually called the matrix of principal components. It is normed so that

$$C^t DC = \Lambda.$$

- Transition Formulæ: Of the four matrices Z, A, L and C we only have to compute one, all others are obtained by the transition formulæ provided by the duality property of the diagram.

$$XQZ = LS = C \quad X^t DL = ZS = A$$

- The $\text{Trace}(VQ) = \text{Trace}(WD)$ is often called the inertia of the diagram, (inertia in the sense of Huyghens inertia formula for instance). The inertia with regards to a point A of a cloud of p_i -weighted points being $\sum_{i=1}^n p_i d^2(x_i, a)$. When we look at ordinary PCA with $Q = \mathbb{I}_p$, $D = \frac{1}{n} \mathbb{I}_n$, and the variables are centered, this is the sum of the variances of all the variables, if the variables are standardized Q is the diagonal matrix of inverse variances, the inertia is the number of variables p .

2.2. Comparing Two Diagrams: the RV coefficient

Many problems can be rephrased in terms of comparison of two "duality diagrams" or put more simply, two characterizing operators, built from two "triplets", usually with one of the triplets being a response or having constraints imposed on it. Most often what is done is to compare two such diagrams and try to get one to match the other in some optimal way.

To compare two symmetric operators there is either a vector covariance $\text{cov}V(O_1, O_2) = \text{Tr}(O_1^t O_2)$ or their vector correlation[8]

$$RV(O_1, O_2) = \frac{\text{Tr}(O_1 O_2)}{\sqrt{\text{Tr}(O_1^t O_1) \text{Tr}(O_2^t O_2)}}$$

If we were to compare the two triplets $(X_{n \times 1}, 1, \frac{1}{n} I_n)$ and $(Y_{n \times 1}, 1, \frac{1}{n} I_n)$ we would have $RV = \rho^2$.

PCA can be seen as finding the matrix Y which maximizes of the RV coefficient between characterizing operators that is between $(X_{n \times p}, Q, D)$ and $(Y_{n \times q}, I, D)$, under the constraint that Y be of rank $q < p$.

$$RV(XQX^t D, YY^t D) = \frac{\text{Tr}(XQX^t D Y Y^t D)}{\sqrt{\text{Tr}(XQX^t D)^2 \text{Tr}(Y Y^t D)^2}}$$

This maximum is attained where Y is chosen as the first q eigenvectors of $XQX^t D$ normed so that $Y^t D Y = \Lambda_q$. The maximum RV is

$$RVmax = \sqrt{\frac{\sum_{i=1}^q \lambda_i^2}{\sum_{i=1}^p \lambda_i^2}}.$$

Of course, classical PCA has $D = \text{identity}$, $Q = \frac{1}{n} \text{identity}$ but the extra flexibility is often useful. We define the distance between triplets (X, Q, D) and (Z, Q, M)

where Z is also $n \times p$ as the distance deduced from the RV inner product between operators XQX^tD and ZMZ^tD . In fact, the reason the French like this scheme so much is that most linear methods can be reframed in these terms. We will give a few examples such as Principal Component Analysis (ACP in French), Correspondence Analysis (AFC in French), Discriminant Analysis (AFD in French), PCA with regards to instrumental variable (ACPVI in French) and Canonical Correlation Analysis (AC).

2.3. Explaining one diagram by another

Principal Component Analysis with respect to Instrumental Variables was a technique developed by C.R. Rao [25] to find the best set of coefficients in multivariate regression setting where the response is multivariate, given by a matrix Y . In terms of diagrams and RV coefficients this problem can be rephrased as that of finding M to associate to X so that $(\mathbf{X}, \mathbf{M}, \mathbf{D})$ is *as close as possible* to $(\mathbf{Y}, \mathbf{Q}, \mathbf{D})$ in the RV sense.

The answer is provided by defining M such that

$$YQY^tD = \lambda XMX^tD.$$

If this is possible then the two eigendecompositions of the triplet give the same answers. We simplify notation by the following abbreviations:

$$X^tDX = S_{xx} \quad Y^tDY = S_{yy} \quad X^tDY = S_{xy} \quad \text{and} \quad R = S_{xx}^{-1}S_{xy}QS_{yx}S_{xx}^{-1}.$$

Then

$$\|YQY^tD - XMX^tD\|^2 = \|YQY^tD - XRX^tD\|^2 + \|XRX^tD - XMX^tD\|^2$$

The first term on the right hand side does not depend on M , and the second term will be zero for the choice $M = R$.

In order to have rank $q < \min(\text{rank}(X), \text{rank}(Y))$ the optimal choice of a positive definite matrix M of $M = RBB^tR$ where the columns of B are the eigenvectors of X^tDXR with 1:

$$B = \left(\frac{1}{\sqrt{\lambda_1}}\beta_1, \dots, \frac{1}{\sqrt{\lambda_q}}\beta_q \right) \left\{ \begin{array}{l} X^tDXR\beta_k = \lambda_k\beta_k \\ \beta_k^tR\beta_k = \lambda_k, k = 1 \dots q \\ \lambda_1 > \lambda_2 > \dots > \lambda_q \end{array} \right.$$

The PCA with regards to instrumental variables of rank q is equivalent to the PCA of rank q of the triplet (X, R, D)

$$R = S_{xx}^{-1}S_{xy}QS_{yx}S_{xx}^{-1}.$$

2.4. One Diagram to replace Two Diagrams

Canonical correlation analysis was introduced by Hotelling[18] to find the common structure in two sets of variables X_1 and X_2 measured on the same observations. This is equivalent to merging the two matrices columnwise to form a large matrix with n rows and $p_1 + p_2$ columns and taking as the weighting of the variables the matrix defined by the two diagonal blocks $(X_1^tDX_1)^{-1}$ and $(X_2^tDX_2)^{-1}$

$$Q = \left(\begin{array}{c|c} (X_1^tDX_1)^{-1} & 0 \\ \hline 0 & (X_2^tDX_2)^{-1} \end{array} \right)$$

$$\begin{array}{ccccc}
\mathbb{R}^{p_1^*} & \xrightarrow{X_1} & \mathbb{R}^n & & \mathbb{R}^{p_2^*} & \xrightarrow{X_2} & \mathbb{R}^n \\
\mathbb{I}_{p_1} \uparrow & \downarrow V_1 & D \downarrow & \uparrow W_1 & \mathbb{I}_{p_2} \uparrow & \downarrow V_2 & D \downarrow & \uparrow W_2 \\
\mathbb{R}^{p_1} & \xleftarrow{X_1^t} & \mathbb{R}^{n^*} & & \mathbb{R}^{p_2} & \xleftarrow{X_2^t} & \mathbb{R}^{n^*} \\
& & \mathbb{R}^{p_1+p_2^*} & \xrightarrow{[X_1; X_2]} & \mathbb{R}^n & & \\
Q \uparrow & \downarrow V & D \downarrow & \uparrow W & & & \\
& & \mathbb{R}^{p_1+p_2} & \xleftarrow{[[X_1; X_2]^t]} & \mathbb{R}^{n^*} & &
\end{array}$$

This analysis gives the same eigenvectors as the analysis of the triple $(X_2^t D X_1, (X_1^t D X_1)^{-1}, (X_2^t D X_2)^{-1})$ also known as the canonical correlation analysis of X_1 and X_2 .

2.5. Discriminant Analysis

If we want to find linear combinations of the original variables $X_{n \times p}$ that characterize the best the group structure of the points given by a zero/one group coding matrix Y , with as many columns as groups, we can phrase the problem as a duality diagram. Suppose that the observations are given individual weights in the diagonal matrix D , and that the variables are centered with regards to these weights.

Let A be the $q \times p$ matrix of group means in each of the p variables. This satisfies

$$Y^t D X = \Delta_Y \mathbf{1}_q A \quad \text{where } \Delta_Y \text{ is the diagonal matrix of group weights.}$$

The variance covariance matrix will be denoted $T = X^t D X$, with elements

$$t_{jk} = \text{cov}(x_j, x_k) = \sum_{i=1}^n d_i (x_{ij} - \bar{x}_j)(x_{ik} - \bar{x}_k)$$

. The between group variance-covariance is

$$B = A^t \Delta_Y A.$$

The duality diagram for linear discriminant analysis is

$$\begin{array}{ccccc}
\mathbb{R}^{p^*} & \xrightarrow{X} & \mathbb{R}^n & & \\
T^{-1} \uparrow & \downarrow B & \Delta_Y \downarrow & \uparrow AT^{-1}A^t & \\
\mathbb{R}^p & \xleftarrow{X^t} & \mathbb{R}^{n^*} & &
\end{array}$$

Corresponding to the triplet $(\mathbf{A}, \mathbf{T}^{-1}, \Delta_Y)$, because

$$(X^t D Y) \Delta_Y^{-1} (Y^t D X) = A^t \Delta_Y A$$

this gives equivalent results to the triplet $(\mathbf{Y}^t \mathbf{D} \mathbf{X}, \mathbf{T}^{-1}, \Delta_Y^{-1})$.

The discriminating variables are the eigenvectors of the operator

$$A^t \Delta_Y A T^{-1}$$

They can also be seen as the PCA with regards to instrumental variables of (Y, Δ^{-1}, D) with regards to (X, M, D) .

2.6. Correspondence Analysis

Correspondence analysis can be used to analyse several types of multivariate data. All involve some categorical variables. Here are some examples of the type of data that can be decomposed using this method:

- Contingency Tables (cross-tabulation of two categorical variables)
- Multiple Contingency Tables (cross-tabulation of several categorical variables).
- Binary tables obtained by cutting continuous variables into classes and then recoding both these variables and any extra categorical variables into 0/1 tables, 1 indicating presence in that class. So for instance a continuous variable cut into three classes will provide three new binary variables of which only one can take the value one for any given observation.

To first approximation, correspondence analysis can be understood as an extension of principal components analysis (PCA) where the variance in PCA is replaced by an *inertia* proportional to the χ^2 distance of the table from independence. CA decomposes this measure of departure from independence along axes that are orthogonal according to a χ^2 inner product. If we are comparing two categorical variables, the simplest possible model is that of independence in which case the counts in the table would obey approximately the margin products identity. For an $m \times p$ contingency table \mathbf{N} with $n = \sum_{i=1}^m \sum_{j=1}^p n_{ij} = n_{..}$ observations and associated to the frequency matrix

$$\mathbf{F} = \frac{\mathbf{N}}{n}$$

Under independence the approximation

$$n_{ij} \doteq \frac{n_{i.} n_{.j}}{n} n$$

can also be written:

$$\mathbf{N} \doteq \mathbf{c} \mathbf{r}^t \mathbf{n}$$

where

$$r = \frac{1}{n} N \mathbf{1}_p \text{ is the vector of row sums of } F \text{ and } c^t = \frac{1}{n} N^t \mathbf{1}_m \text{ are the column sums}$$

The departure from independence is measured by the χ^2 statistic

$$\chi^2 = \sum_{i,j} \left[\frac{(n_{ij} - \frac{n_{i.} n_{.j}}{n})^2}{\frac{n_{i.} n_{.j}}{n^2} n} \right]$$

Under the usual validity assumptions that the cell counts n_{ij} are not too small, this statistic is distributed as a χ^2 with $(m-1)(p-1)$ degrees of freedom if the data are independent. If we do not reject independence, there is no more to be said about the table, no interaction of interest to analyse. There is in fact no ‘multivariate’ effect.

On the contrary if this statistic is large, we decompose it into one dimensional components.

Correspondence analysis is equivalent to the eigendecomposition of the triplet $(\mathbf{X}, \mathbf{Q}, \mathbf{D})$ with

$$\mathbf{X} = \mathbf{D}_r^{-1} \mathbf{F} \mathbf{D}_c^{-1} - \mathbf{1}^t \mathbf{1}, \mathbf{Q} = \mathbf{D}_c, \mathbf{D} = \mathbf{D}_r$$

$$\mathbf{D}_c = \text{diag}(\mathbf{c}), \mathbf{D}_r = \text{diag}(\mathbf{r})$$

$X'D_r \mathbf{1}_m = \mathbf{1}_p$, the average of each column is one.

Notes:

1. Consider the matrix $\mathbf{D}_r^{-1} \mathbf{F} \mathbf{D}_c^{-1}$ and take the principal components with regards to the weights \mathbf{D}_r for the rows and \mathbf{D}_c for the columns. The recentered matrix $\mathbf{D}_r^{-1} \mathbf{F} \mathbf{D}_c^{-1} - \mathbf{1}'_m \mathbf{1}_p$ has a generalized singular value decomposition

$$\mathbf{D}_r^{-1} \mathbf{F} \mathbf{D}_c^{-1} - \mathbf{1}'_m \mathbf{1}_p = \mathbf{U} \mathbf{S} \mathbf{V}', \text{ with } \mathbf{U}' \mathbf{D}_r \mathbf{U} = \mathbf{I}_m, \mathbf{V}' \mathbf{D}_c \mathbf{V} = \mathbf{I}_p$$

having total inertia:

$$\mathbf{D}_r (\mathbf{D}_r^{-1} \mathbf{F} \mathbf{D}_c^{-1} - \mathbf{1}'_m \mathbf{1}_p)' \mathbf{D}_c (\mathbf{D}_r^{-1} \mathbf{F} \mathbf{D}_c^{-1} - \mathbf{1}'_m \mathbf{1}_p) = \frac{\chi^2}{\mathbf{n}}$$

2. PCA of the row profiles $F \mathbf{D}_r^{-1}$, taken with weight matrix D_c with the metric $Q = \mathbf{D}_c^{-1}$.
3. Notice that

$$\sum_i f_i \left(\frac{f_{ij}}{f_i \cdot f_{.j}} - 1 \right) = 0$$

the row and columns profiles are centered

$$\sum_j f_j \left(\frac{f_{ij}}{f_i \cdot f_{.j}} - 1 \right) = 0$$

This method has been rediscovered many times, the most recently by Jon Kleinberg's in his method for analysing Hubs and Authorities [19], see Fouss, Saerens and Renders, 2004[11] for a detailed comparison.

In statistics the most commonplace use of Correspondence Analysis is in ordination or seriation, that is , the search for a hidden gradient in contingency tables. As an example we take data analysed by Cox and Brandwood [4] and [6] who wanted to seriate Plato's works using the proportion of sentence endings in a given book, with a given stress pattern. We propose the use of correspondence analysis on the table of frequencies of sentence endings, for a detailed analysis see Charnomordic and Holmes[2].

The first 10 profiles (as percentages) look as follows:

	Rep	Laws	Crit	Phil	Pol	Soph	Tim
UUUUU	1.1	2.4	3.3	2.5	1.7	2.8	2.4
-UUUU	1.6	3.8	2.0	2.8	2.5	3.6	3.9
U-UUU	1.7	1.9	2.0	2.1	3.1	3.4	6.0
UU-UU	1.9	2.6	1.3	2.6	2.6	2.6	1.8
UUU-U	2.1	3.0	6.7	4.0	3.3	2.4	3.4
UUUU-	2.0	3.8	4.0	4.8	2.9	2.5	3.5
--UUU	2.1	2.7	3.3	4.3	3.3	3.3	3.4
-U-UU	2.2	1.8	2.0	1.5	2.3	4.0	3.4
-UU-U	2.8	0.6	1.3	0.7	0.4	2.1	1.7
-UUU-	4.6	8.8	6.0	6.5	4.0	2.3	3.3
.....etc (there are 32 rows in all)							

The eigenvalue decomposition (called the scree plot) of the chisquare distance matrix (see [2]) shows that two axes out of a possible 6 (the matrix is of rank 6) will provide a summary of 85% of the departure from independence, this suggests that a planar representation will provide a good visual summary of the data.

	Eigenvalue	inertia %	cumulative %
1	0.09170	68.96	68.96
2	0.02120	15.94	84.90
3	0.00911	6.86	91.76
4	0.00603	4.53	96.29
5	0.00276	2.07	98.36
6	0.00217	1.64	100.00

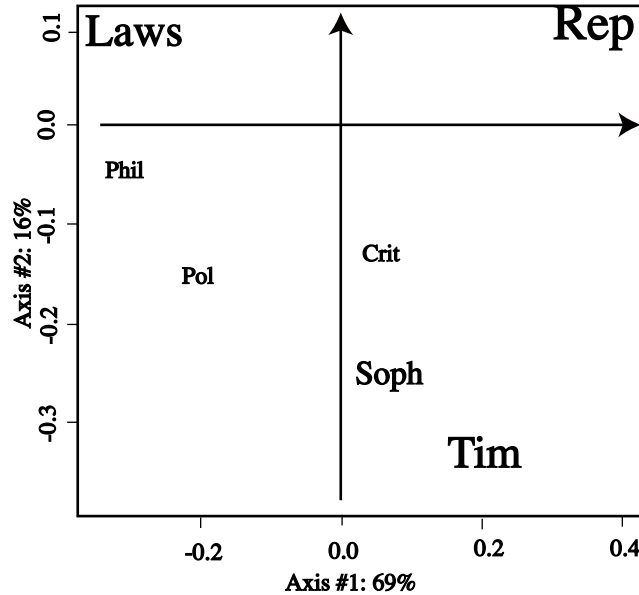


Figure 1: Correspondence Analysis of Plato's Works

We can see from the plot that there is a seriation that as in most cases follows a parabola or arch [16] from Laws on one extreme being the latest work and Republica being the earliest among those studied.

3. From Discriminant Analysis to Networks

Consider a graph with vertices the members of a group and edges if two members interact. We suppose each vertex comes with an observation vector x_i , and that each has the same weight $\frac{1}{n}$. In the extreme case of discriminant analysis, the graph is supposed to connect all the points of a group in a complete graph, and be disconnected from the other observations. Discriminant Analysis is just the explanation of this particular graph by linear combinations of variables, what we propose here is to extend this to more general graphs in a similar way. We will suppose all the observations are the nodes of the graph and each has the same weight $\frac{1}{n}$. The basic

decomposition of the variance is written

$$\begin{aligned} \mathbf{cov}(x_j, x_k) = t_{jk} &= \frac{1}{n} \sum_{i=1}^n (x_{ij} - \bar{x}_j)(x_{ik} - \bar{x}_k) \\ \text{Call the group means } \bar{x}_{gj} &= \frac{1}{n_g} \sum_{i \in G_g} x_{ij}, g = 1 \dots q \\ \sum_{i \in G_g} (x_{ij} - \bar{x}_{gj})(\bar{x}_{gj} - \bar{x}_k) &= (\bar{x}_{gj} - \bar{x}_k) \sum_{i \in G_g} (x_{ij} - \bar{x}_{gj}) = 0 \\ \text{Huyghens formula} \quad t_{jk} &= w_{jk} + b_{jk} \\ \text{Where } w_{jk} &= \sum_{g=1}^q \sum_{i \in G_g} (x_{ij} - \bar{x}_{gj})(x_{ik} - \bar{x}_{gk}) \\ \text{and } b_{jk} &= \sum_{g=1}^q \frac{n_g}{n} (\bar{x}_{gj} - \bar{x}_j)(\bar{x}_{gk} - \bar{x}_k) \\ T &= W + B \end{aligned}$$

As we showed above, linear discriminant analysis finds the linear combinations a such that $\frac{a^t B a}{a^t T a}$ is maximized. This is equivalent to maximizing the quadratic form $a^t B a$ in a , subject to the constraint $a^t T a = 1$. As we saw above, this is solved by the solution to the eigenvalue problem

$$B a = \lambda T a \text{ or } T^{-1} B a = \lambda a \text{ if } T^{-1} \text{ exists}$$

then $a^t B a = \lambda a^t T a = \lambda$. We are going to extend this to graphs by relaxing the group definition to partition the variation into local and global components

3.1. Decomposing the Variance into Local and Global Components

Lebart was a pioneer in adapting the eigenvector decompositions to cater to spatial structure in the data (see [20, 21, 22]).

$$\begin{aligned} \mathbf{cov}(x_j, x_k) &= \frac{1}{n} \sum_{i=1}^n (x_{ij} - \bar{x}_j)(x_{ik} - \bar{x}_k) \\ &= \frac{1}{2n^2} \sum_{i=1}^n \sum_{i'=1}^n (x_{ij} - x_{i'j})(x_{ik} - x_{i'k}) \\ \mathbf{var}(x_j) &= \frac{1}{2n^2} \left\{ \sum_{(i,i') \in E} (x_{ij} - x_{i'j})^2 + \sum_{(i,i') \notin E} (x_{ij} - x_{i'j})^2 \right\} \end{aligned}$$

If we call M is the incidence matrix of the directed graph: $m_{ij} = 1$ if i points to j . Suppose for the time being that M is symmetrical (the graph is undirected). The degree of vertex i is $m_i = \sum_{i'=1}^n m_{ii'}$. We take the convention that there are no self

loops. Then another way of writing the variance formula is

$$\begin{aligned}\mathbf{var}(x_j) &= \frac{1}{2n^2} \left\{ \sum_{i=1}^n \sum_{i'=1}^n m_{ii'} (x_{ij} - x_{i'j})^2 + \sum_{(i,i') \notin E} (x_{ij} - x_{i'j})^2 \right\} \\ \mathbf{var}_{loc}(x_j) &= \frac{1}{2m} \sum_{i=1}^n \sum_{i'=1}^n m_{ii'} (x_{ij} - x_{i'j})^2 \\ \text{where } m &= \sum_{i=1}^n \sum_{i'=1}^n m_{ii'}\end{aligned}$$

The total variance is the variance of the complete graph. Geary's ratio [14] is used to see whether the variable x_j can be considered as independent of the graph structure. If the neighboring values of x_j seem correlated then the local variance will only be an underestimate of the variance:

$$G = c(x_j) = \frac{\mathbf{var}_{loc}(x_j)}{\mathbf{var}(x_j)}$$

Call D the diagonal matrix with the total degrees of each node in the diagonal $D = \text{diag}(m_i)$.

For all variables taken together, $j = 1, \dots, p$ note the local covariance matrix $\mathbf{V} = \frac{1}{2m} X^t (D - M) X$, if the graph is just made of disjoint groups of the same size, this is proportional to the W within class variance-covariance matrix. The proportionality can be accomplished by taking the average of the sum of squares to the average of the neighboring nodes ([23]). We can generalize the Geary index to account for irregular graphs coherently. In this case we weight each node by its degree. Then we can write the Geary ratio for any n -vector x as

$$c(x) = \frac{x^t (D - M) x}{x^t D x}, \quad D = \begin{pmatrix} m_1 & 0 & 0 & 0 \\ & m_2 & 0 & 0 \\ \vdots & & \ddots & \vdots \\ \dots & 0 & 0 & m_n \end{pmatrix}$$

We can ask for the coordinate(s) that are the most correlated to the graph structure then if we want to minimize the Geary ratio, choose x such that $c(x)$ is minimal, this is equivalent to minimizing $x^t (D - M) x$ under the constraint $x^t D x = 1$. This can be solved by finding the smallest eigenvalue μ with eigenvector x such that:

$$\begin{aligned}(D - M)x &= \mu D x \\ D^{-1}(D - M)x &= \mu x \\ (1 - \mu)x &= D^{-1} M x\end{aligned}$$

This is exactly the defining equation of the correspondence analysis of the matrix M . This can be extended to as many coordinates as we like, in particular we can take the first 2 largest eigenvectors and provide the best planar representation of the graph in this way.

3.2. Regression of graphs on node covariates

The covariables measured on the nodes can be essential to understanding the fine structure of graphs. We call \mathbf{X} the $n \times p$ matrix of measurements at the vertices of the

graph; they may be a combination of both categorical variables (gene families, GO classes) and continuous measurements (expression scores). We can use the PCAIV method defined in section 2 to the eigenvectors of the graph defined above. This provides a method that uses the covariates in X to explain the graph. To be more precise, given a graph (V, E) with adjacency matrix M , define the Laplacian

$$L = D^{-1}(M - I) \quad D = \text{diag}(d_1, d_2, \dots, d_n) \text{ diagonal matrix of degrees}$$

Using the eigenanalysis of the graph, we can summarize the graph with a few variables, the first few relevant eigenvectors of L , these can then be regressed on the covariates using Principal Components with respect to Instrumental Variables [25] as defined above to find the linear combination of node covariates that explain the graph variables the best.

Appendix A: Resources

A.1. Reading

There are few references in English explaining the duality/operator point of view, apart from the already cited references of Escoufier [8, 10]. Frédérique Glaçon's PhD thesis [15] (in French) clearly lays out the duality principle before going on to explain its application to the conjoint analysis of several matrices, or data cubes. The interested reader fluent in French could also consult any one of several Masters level textbooks on the subject for many details and examples:

Brigitte Escoufier and Jérôme Pagès [7] have a textbook with many examples, although their approach is geometric, they do not delve into the Duality Diagram, more than explaining on page 100 its use in transition formula between eigenbases of the different spaces.

[22] is one of the broader books on multivariate analyses, making connections between modern uses of eigendecomposition techniques, clustering and segmentation. This book is unique in its chapter on stability and validation of results (without going as far as speaking of inference).

Caillez and Pagès [1] is hard to find, but was the first textbook completely based on the diagram approach, as was the case in the earlier literature they use transposed matrices.

A.2. Software

The methods described in this article are all available in the form of R packages which I recommend. The most complete package is `ade4` [3] which covers almost all the problems I mention except that of regressing graphs on covariates, however a complete understanding of the duality diagram terminology and philosophy is necessary as these provide the building blocks for all the functions in the form a class called `dudi` (this actually stands for duality diagram). One of the most important features in all the '`dudi.*`' functions is that when the argument `scanff` is at its default value `TRUE` the first step imposed on the user is the perusal of the screeplot of eigenvalues. This can be very important, as choosing to retain 2 values by default before consulting the eigenvalues can lead to the main mistake that can be made when using these techniques: the separation of two close eigenvalues. When two eigenvalues are close the plane will be stable, but not each individual axis or

principal component resulting in erroneous results if for instance the 2nd and 3rd eigenvalues were very close and the user chose to take 2 axes[17].

Another useful addition also comes from the ecological community and is called `vegan`. Here is a list of suggested functions from several packages:

- Principal Components Analysis (PCA) is available in `prcomp` and `princomp` in the standard package `stats` as `pca` in `vegan` and as `dudi.pca` in `ade4`.
- Two versions of PCAIV are available, one is called Redundancy Analysis (RDA) and is available as `rda` in `vegan` and `pcaiv` in `ade4`.
- Correspondence Analysis (CA) is available in `cca` in `vegan` and as `dudi.coa` in `ade4`.
- Discriminant analysis is available as `lda` in `stats`, as `discrimin` in `ade4`
- Canonical Correlation Analysis is available in `cancor` in `stats` (Beware `cca` in `ade4` is Canonical Correspondence Analysis).
- STATIS (Conjoint analysis of several tables) is available in `ade4`.

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