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# From Multiblock Partial Least Squares to Multiblock Redundancy Analysis. A Continuum Approach

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Received: October 2009; accepted: October 2010

**Abstract.** For the purpose of exploring and modelling the relationships between a dataset and several datasets, multiblock Partial Least Squares is a widely-used regression technique. It is designed as an extension of PLS which aims at linking two datasets. In the same vein, we propose an extension of Redundancy Analysis to the multiblock setting. We show that PLS and multiblock Redundancy Analysis aim at maximizing the same criterion but the constraints are different. From the solutions of both these approaches, it turns out that they are the two end points of a continuum approach that we propose to investigate.

**Keywords:** multiblock *PLS*, multiblock redundancy analysis, continuum approach, Ridge-type regularization, multicolinearity.

## 1. Introduction

This paper deals with the description and the prediction of multiblock data organized in (K+1) blocks consisting of K explanatory blocks  $(X_1,\ldots,X_K)$  and a Y dataset to be explained. The first issue is to describe the multiblock tables and sum up the relationships between the variables and between the datasets. For this purpose, we seek overall variables (latent variables) which highlight the relationships between the various datasets. The second issue is to predict Y from the K tables  $(X_1,\ldots,X_K)$ , determine which  $X_k$  blocks are best related to the Y variables and within these blocks which variables have an impact on Y.

Multiblock Partial Least Squares (Wold, 1984) is a regression technique that is widely used in the field of chemometrics, sensometrics and process monitoring for the purpose of exploring and modelling the relationships between several datasets to be predicted from several other datasets. Thus, not only a multiblock approach makes it possible to combine several sources of information, but it also highlights the importance of each block in the prediction of the response variables. In the case where only one block of variables is explained by several blocks of explanatory variables, (Westerhuis et al., 1998; Qin et al., 2001; Vivien, 2002) show that the solution obtained from the iterative algorithm of multiblock PLS (mbPLS) is equivalent to the solution obtained from a PLS regression of Y and X, where X is the merged dataset, namely  $X = [X_1|, \dots, |X_K|]$ . Redundancy Analysis, RA (Rao, 1964; Van Den Wollenberg, 1977), is yet another popular method for linking two datasets. In a previous paper, we compared the merits of RA and PLSregression (Bougeard et al., 2008). We propose to extend Redundancy Analysis to the multiblock setting and compare this approach to mbPLS. Redundancy Analysis, also called Principal Component Analysis with respect to Instrumental Variables, was introduced by Rao (1964) and further investigated by Van der Wollenberg (1997) and Sabatier (1984) among others. These authors gave several formulations of RA which clearly show how this method of analysis can be seen as a regression of Y upon linear combinations of the variables  $(x_1, \ldots, x_P)$  or as a principal component analysis of the Y variables where components are constrained to be linear combinations of  $(x_1, \ldots, x_P)$ . Similarly to PLSregression, the components thus obtained may be used for an exploratory purpose to investigate the relationships between  $(x_1, \ldots, x_P)$  and Y or to set up prediction models. This latter approach is called *reduced-rank regression* (Muller, 1981; Davies and Tso, 1982). For the purpose of exploring and modelling the relationships between a dataset Y and several datasets  $(X_1, \ldots, X_K)$ , we propose, in a first stage, a new method called multiblock Redundancy Analysis (mbRA), based on the same maximization criterion as mbPLS with different constraints on the components to be determined. In a second stage, we highlight the connection between multiblock Redundancy Analysis and multiblock PLS. It turns out that mbPLS and mbRA are the two end points of a continuum approach that we propose to investigate. As this continuum approach establishes a bridge between mbPLS and mbRA, we shall refer to it as "multiblock Continuum Redundancy PLS regression" (mbCR). We discuss how the proposed methods are related to other statistical techniques. Finally, the interest of the multiblock methods and the properties of the continuum are illustrated on the basis of a simulation study and on a real dataset in the field of veterinary epidemiology.

#### 2. Methods

#### 2.1. Notations

Consider the multiblock setting where we have (K+1) datasets: a dataset Y to be predicted from K datasets  $X_k$   $(k=1,\ldots,K)$ . The Y table contains Q variables and each

table  $X_k$  contains  $p_k$  variables. The merged dataset X is defined as  $[X_1|,\ldots,|X_K]$  and contains  $P=\sum_k p_k$  explanatory variables. All these quantitative variables are measured on the same N individuals and supposed to be column centred.

#### 2.2. Multiblock PLS Regression

Wold introduced multiblock Partial Least Squares as an alternative procedure based on the Non-linear Iterative PArtial Least Squares (NIPALS) algorithm (Wold, 1984; Struc and Pavesic, 2009). This algorithm was further investigated by Wangen and Kowalski (1988). The initial method aims at linking several tables  $X_k$  ( $k=1,\ldots,K$ ) to one (or more) data table(s) Y. Westerhuis *et al.* (1998), Qin *et al.* (2001), Vivien (2002) showed that the solution obtained from the iterative algorithm of mbPLS is equivalent to the solution obtained from a PLS regression of Y and X where X is the merged dataset. More precisely, Vivien (2002) proved that mbPLS seeks, in a first step, a component  $t^{(1)} = Xw^{(1)}$  which is highly related to a component  $u^{(1)} = Yv^{(1)}$  and which sums up partial components  $t_k^{(1)}$  respectively associated with the blocks  $X_k$ . More formally, mbPLS consists in maximizing criterion (1):

$$cov^{2}(u^{(1)}, t^{(1)}) \quad \text{with } t^{(1)} = \sum_{k=1}^{K} a_{k}^{(1)} t_{k}^{(1)}, \ u^{(1)} = Y v^{(1)}, \ t_{k}^{(1)} = X_{k} w_{k}^{(1)}, \\
\sum_{k=1}^{K} a_{k}^{(1)^{2}} = 1, \quad \|w_{k}^{(1)}\| = \|v^{(1)}\| = 1.$$
(1)

The optimal vector of loadings  $w^{(1)}$  is given by the eigenvector of the matrix (X'YY'X) associated with the largest eigenvalue  $\lambda_{mbPLS}^{(1)}$  (Westerhuis et~al., 1998). The vector  $v^{(1)}$  is given by the eigenvector of  $M_{mbPLS}=(Y'XX'Y)$  associated with the same eigenvalue. Thereafter the partial vectors of loadings  $w_k^{(1)}$  are given by  $w_k^{(1)}=w_k^{(1)*}/\|w_k^{(1)*}\|$  where  $w_k^{(1)*}$  are the block sub-vectors of  $w^{(1)}$ , namely  $w^{(1)}=[w_1^{(1)*}|,\ldots,|w_K^{(1)*}|]'$  (Qin et~al., 2001). It is clear that  $a_k^{(1)}=\|w_k^{(1)*}\|$  which indeed fulfills the constraint  $\sum_k a_k^{(1)^2}=1$ .

Thereafter, the same analysis is performed by replacing  $(X_1, \ldots, X_K)$  by their residual in the orthogonal projection onto the subspace spanned by the first global component  $t^{(1)}$  (Westerhuis and Smilde, 2001). This process is reiterated in order to determine subsequent components. It is worth noting that Wangen and Kowalski (1988) use a block score deflation, i.e., deflation of each block  $X_k$  with respect to its associated partial component  $t_k$ . This leads to a slightly different mbPLS strategy of analysis.

# 2.3. Proposition of a Multiblock Redundancy Analysis

For the purpose of exploring and modelling the relationships between two data tables  $X = (x_1, \ldots, x_P)$  and Y, it has been shown that Redundancy Analysis and PLS regression are based on the same criterion to maximize, namely  $cov^2(t, u)$  with t = Xw

and u=Yv, associated with different norm constraints imposed on the components to be determined (Burnham *et al.*, 1996; Bougeard *et al.*, 2008). More precisely, PLS regression imposes the constraints  $\|w\| = \|v\| = 1$  whereas Redundancy Analysis imposes the constraints  $\|t\| = \|v\| = 1$ . In the same vein as mbPLS, we propose an extension of Redundancy Analysis to the multiblock setting. This leads us to consider the following maximization problem (2).

$$cov2(u(1), t(1)) with t(1) = \sum_{k=1}^{K} a_k(1) t_k(1), u(1) = Y v(1), t_k(1) = X_k w_k(1), 
\sum_{k=1}^{K} a_k(1)2 = 1, ||t_k(1)|| = ||v(1)|| = 1.$$
(2)

As previously, the method derives a global component  $t^{(1)} = Xw^{(1)}$  oriented towards the explanation of Y, that sums up partial components  $t_k^{(1)}$  for  $k = (1, \ldots, K)$  respectively associated with the blocks  $X_k$ . In the case where there is only one block of explanatory variables (K = 1), it is clear that multiblock Redundancy Analysis (mbRA) amounts to RA.

Replacing the global component  $t^{(1)}$  by its expression as a linear combination of the partial components  $t_k^{(1)}$ , we are led to maximizing the criterion  $\cos^2(u^{(1)},t^{(1)})=[\sum_k a_k^{(1)} \cos(u^{(1)},t_k^{(1)})]^2$  under the constraints stated above. The optimal solutions are given by:

$$a_k^{(1)} = \frac{\operatorname{cov}(u^{(1)}, t_k^{(1)})}{\sqrt{\sum_{l=1}^K \operatorname{cov}^2(u^{(1)}, t_l^{(1)})}}.$$

Therefore the criterion to be maximized amounts to  $\sum_{k=1}^K \cos^2(u^{(1)}, t_k^{(1)})$ . The maximization problem becomes (3):

$$\sum_{k=1}^{K} \operatorname{cov}^{2}(u^{(1)}, t_{k}^{(1)}) \quad \text{with } t_{k}^{(1)} = X_{k} w_{k}^{(1)},$$

$$u^{(1)} = Y v^{(1)}, \ \|t_{k}^{(1)}\| = \|v^{(1)}\| = 1.$$
(3)

The criterion (3) highlights the optimal link between datasets Y and  $(X_1, \ldots, X_K)$ . It follows:

$$\sum_{k=1}^{K} \cot^{2} \left( u^{(1)}, t_{k}^{(1)} \right) = \sum_{k} \left[ w_{k}^{(1)'} X_{k}' u^{(1)} \right]^{2} = \sum_{k} \left[ b_{k}^{(1)'} (X_{k}' X_{k})^{-1/2} X_{k}' u^{(1)} \right]^{2}, (4)$$

where  $b_k^{(1)}$  is defined as  $b_k^{(1)}=(X_k'X_k)^{1/2}w_k^{(1)}$ . In the previous equations, we have dropped the term 1/N from the expression of the covariance, for simplicity sake. The constraint  $\|t_k^{(1)}\|=1$  can be expressed as  $\|b_k^{(1)}\|=1$ . The maximization of the criterion

(4) leads to  $b_k^{(1)} = (X_k' X_k)^{-1/2} X_k' u^{(1)} / \|(X_k' X_k)^{-1/2} X_k' u^{(1)}\|$ . Including this expression in criterion (4) and replacing  $u^{(1)}$  by  $Yv^{(1)}$ , we are led to:

$$\sum_{k=1}^{K} \operatorname{cov}^{2}(u^{(1)}, t_{k}^{(1)}) = \sum_{k} v^{(1)'} Y' X_{k} (X_{k}' X_{k})^{-1} X_{k}' Y v^{(1)}.$$
 (5)

It directly follows that the solution is given by  $v^{(1)}$  the normalized eigenvector of the matrix  $M_{mbRA} = \sum_k Y' X_k (X_k' X_k)^{-1} X_k' Y$  associated with the largest eigenvalue  $\lambda_{mbRA}^{(1)}$ . The partial components  $(t_1^{(1)},\ldots,t_K^{(1)})$  are therefore given by  $t_k^{(1)} = X_k w_k^{(1)} = X_k (X_k' X_k)^{-1/2} b_k^{(1)} = P_{X_k} u^{(1)} / \|P_{X_k} u^{(1)}\|$ , where  $P_{X_k} = X_k (X_k' X_k)^{-1} X_k'$  is the projector onto the subspace spanned by the  $X_k$  variables. The partial components  $(t_1^{(1)},\ldots,t_K^{(1)})$  are given by the projection of  $u^{(1)}$  on each subspace respectively spanned by  $X_1,\ldots$  and  $X_K$ . The coefficients  $a_k^{(1)}$  can also be given by  $a_k^{(1)} = \cos(u^{(1)},t_k^{(1)})/\sqrt{\sum_l \cos^2(u^{(1)},t_l^{(1)})} = \|P_{X_k} u^{(1)}\|/\sqrt{\sum_l \|P_{X_l} u^{(1)}\|^2}$ . These coefficients reflect the link between the Y and the  $X_k$  datasets for  $k=(1,\ldots,K)$ . This implies that the global component  $t^{(1)} = \sum_k a_k^{(1)} t_k^{(1)} = \sum_k P_{X_k} u^{(1)}/\sqrt{\sum_l \|P_{X_l} u^{(1)}\|^2}$ . We recall that the optimal solution of the maximization of the criterion (2) is based

We recall that the optimal solution of the maximization of the criterion (2) is based on the eigenvector of the matrix  $M_{mbRA} = \sum_k Y' X_k (X_k' X_k)^{-1} X_k' Y$ . Because projectors are symmetric and idempotent  $(P_{X_k}^2 = P_{X_k})$ , it follows that  $M_{mbRA} = \sum_k (P_{X_k} Y)' (P_{X_k} Y)$ . From this standpoint, mbRA appears as a principal component analysis of the table obtained by the vertical concatenation of the projection of Y onto each subspace spanned by the  $X_k$  blocks. Moreover, criterion (5) can also be written as:

$$(5) = v^{(1)'}Y' \sum_{k=1}^{K} P_{X_k} Y v^{(1)} = u^{(1)'} \sum_{k} P_{X_k} u^{(1)} = \sum_{k} \operatorname{var} (P_{X_k} u^{(1)}).$$
 (6)

It follows that mbRA consists in maximizing the sum of the variance of the projections of  $u^{(1)} = Yv^{(1)}$  onto the subspace spanned by the  $X_k$  variables.

As a summing up, the various components in mbRA can be determined as follows:

- 1. Compute  $P_{X_k} = X_k (X_k' X_k)^{-1} X_k'$  and  $M_{mbRA} = \sum_k (P_{X_k} Y)' (P_{X_k} Y)$ .
- 2. Compute  $v^{(1)}$ , the normalized eigenvector of  $M_{mbRA}$  associated with the largest eigenvalue and set  $u^{(1)} = Yv^{(1)}$ .
- 3. Set  $t_k^{(1)} = P_{X_k} u^{(1)} / \|P_{X_k} u^{(1)}\|$ .
- 4. Set  $t^{(1)} = \sum_{k} P_{X_k} u^{(1)} / \sqrt{\sum_{k} \|P_{X_l} u^{(1)}\|^2}$ .

In order to obtain second order solutions, i.e., a global component  $t^{(2)}$  and partial components  $(t_1^{(2)},\ldots,t_K^{(2)})$  and  $u^{(2)}$ , we propose to follow the same strategy as mbPLS. This consists in deflating the  $(X_1,\ldots,X_K)$  datasets by projection onto  $t^{(1)}$  and considering the residuals. Subsequent components can be found by reiterating this process.

### 2.4. Continuum Between Multiblock PLS and Multiblock Redundancy Analysis

It turns out that mbRA and mbPLS regression are respectively based on the eigenstructure of matrices  $M_{mbRA} = \sum_k Y' X_k (X_k' X_k)^{-1} X_k' Y$  and  $M_{mbPLS} = Y' X X' Y =$  $\sum_{k} Y' X'_{k} X_{k} Y$ . Thus, it appears that mbPLS corresponds to a shrinkage of matrices  $(X_k'X_k)^{-1}$  towards the identity matrices  $I_{p_k}$  for  $k=(1,\ldots,K)$ . From this standpoint, we can adopt a gradual shrinkage of the matrices  $(X'_k X_k)^{-1}$  towards  $I_{p_k}$  by considering a convex combination of these matrices (Saudargiene, 1999). More precisely for a scalar  $\gamma$  comprised between 0 and 1, the various components of the continuum approach can be determined as follows:

- 1. Compute  $P_{X_k,\gamma} = X'_k [(1-\gamma)(X'_k X_k) + \gamma I_{p_k}]^{-1} X_k$  and  $M_{\gamma} = \sum_{k} (P_{X_{k},\gamma}Y)'(P_{X_{k},\gamma}Y).$
- 2. Compute  $v_{\gamma}^{(1)}$  the normalized eigenvector of  $M_{\gamma}$  associated with the largest eigen-
- 2. Compute  $v_{\gamma}$  ' the normalized eigenvector of  $M_{\gamma}$  associated with the largest eigenvalue  $\lambda_{\gamma}^{(1)}$  and set  $u_{\gamma}^{(1)} = Y v_{\gamma}^{(1)}$ .

  3. Set  $w_{k,\gamma}^{(1)} = [(1-\gamma)(X_k'X_k) + \gamma I_{p_k}]^{-1} X_k' u_{\gamma}^{(1)} / \|X_k[(1-\gamma)(X_k'X_k) + \gamma I_{p_k}]^{-1/2}$ .  $X_k' u_{\gamma}^{(1)} \|$  and then the partial components  $t_{k,\gamma}^{(1)} = X_k w_{k,\gamma}^{(1)}$ .

  4. Set the coefficients  $a_{k,\gamma}^{(1)} = \text{cov}(u_{\gamma}^{(1)}, t_{k,\gamma}^{(1)}) / \sqrt{\sum_{l} \text{cov}^2(u_{\gamma}^{(1)}, t_{l,\gamma}^{(1)})}$  and then set the
- global component  $t_{\gamma}^{(1)} = \sum_{k} a_{k,\gamma}^{(1)} t_{k,\gamma}^{(1)}$  or set directly  $t_{\gamma}^{(1)} = \sum_{k} P_{X_k, \gamma} u_{\gamma}^{(1)} / \sqrt{\sum_{k} \|P_{X_l, \gamma} u_{\gamma}^{(1)}\|^2}.$

It is clear that the case ( $\gamma = 0$ ) corresponds to mbRA applied to the datasets  $(Y, X_1, \ldots, X_K)$  whereas the case  $(\gamma = 1)$  corresponds to mbPLS. We shall refer to this strategy of analysis as multiblock Continuum Redundancy PLS regression (mbCR). As previously, subsequent components can be obtained by a stagewise procedure by deflating the  $X_k$  datasets with respect to the global components obtained in earlier stages.

The introduction of parameter  $\gamma$  is intended to prevent the instability of the prediction models in case of multicolinearity among the variables in  $X_k$ . Indeed, the sensitivity to multicolinearity can be reflected by the condition index (Belsley et al., 1980). The condition index  $\eta_k$  of matrix  $(X'_k X_k)$  is the ratio of its largest eigenvalue  $\lambda_k^{(1)}$  to its smallest eigenvalue  $\lambda_k^{(p_k)}$  of matrix  $(X_k'X_k)$ . A large value of  $\eta_k$  flags the presence of multicolinearity among  $X_k$  which is likely to lead to an unstable model. The condition index of each matrix  $[(1-\gamma)X'_kX_k + \gamma I_{p_k}]$  is given by:

$$\eta_{k,\gamma} = \frac{[(1-\gamma)\lambda_k^{(1)} + \gamma]}{[(1-\gamma)\lambda_k^{(p_k)} + \gamma]} \quad \text{for } k = (1, \dots, K).$$

It is easy to prove, by considering its derivative with respect to  $\gamma$ , that each  $\eta_{k,\gamma}$  decreases when  $\gamma$  increases. Within mbCR, mbPLS corresponds to the smallest values of  $\eta_{k,\gamma}$  whereas mbRA corresponds to the largest ones. Thus, parameter  $\gamma$  stands as a regularization parameter as it improves the conditioning of each matrix  $(X'_k X_k)$ .

## 2.5. Prediction of Y from $(X_1, \ldots, X_K)$

For all the methods previously described, e.g., mbPLS, mbRA and mbCR, the prediction of the Y variables can be obtained by regressing the Y variables onto the global components  $(t^{(1)},\ldots,t^{(h)})$ . These components being orthogonal by construction, the Y table is split up into:  $Y=t^{(1)}c^{(1)'}+\cdots+t^{(h)}c^{(h)'}+Y^{(h)},Y^{(h)}$  being the matrix of residuals. Moreover, the global components can be expressed as linear combinations of X:  $t^{(1)}=Xw^{*(1)},\ldots,t^{(h)}=Xw^{*(h)}$ . The vectors of loadings  $w^*$  and c are defined as in PLS regression. This leads to the model (7):

$$Y = X \left[ w^{*(1)} c^{(1)'} + \dots + w^{*(h)} c^{(h)'} \right] + Y^{(h)}.$$
(7)

From a practical point of view, the final model may be obtained by selecting the optimal number h of components to be introduced in the model and the  $\gamma$  parameter, by a validation technique such as cross-validation (Stone, 1974). This consists in splitting the whole dataset into two sets, namely a calibration set and a validation set. The calibration set is used to select the parameters of the model and the root mean square error of calibration  $(RMSE_C)$  which reflects the fitting ability of the model. The validation set is used to compute the root mean square error of validation  $(RMSE_V)$  which reflects the prediction ability of the model under consideration.

$$RMSE^{(h)} = ||Y - \hat{Y}^{(h)}|| / \sqrt{Q},$$
 (8)

where  $\hat{Y}^{(h)}$  is the matrix of predicted values from a model with h components. Thereafter, this procedure is repeated several times. For each number h of components to be introduced in the model, the optimal value of  $\gamma$  is determined by minimizing  $RMSE_V$ . Among all these models corresponding to the various values of h, a compromised model with a correct fitting ability and a good prediction ability is retained.

# 2.6. Alternative Methods

It is worth mentioning that several methods are proposed in the literature in order to investigate the relationships among datasets. Among these methods, we can distinguish strategies of analysis which fit into the framework of generalized canonical analysis (Horst, 1961; Carroll, 1968). We refer to Kissita (2003) for a review of such methods. Another family of methods pertains to PLS regression and its extensions. We refer to Vivien (2002) for a detailed discussion of these methods. PLS path modelling, PLS-PM (Wold, 1982; Markauskaite, 2001) and more generally structural equation modelling are also worth mentioning in this context. The Generalized Structured Component Analysis, GSCA (Hwang and Takane, 2004), as an alternative method to PLS-PM may also be mentioned. However, this method pertaining to the field of structural equation modelling follows a specific pattern of analysis based on conceptual models which should be set up by the user beforehand. Among all these techniques of analysis, we single out those methods which are based on the same maximization criterion as mbPLS and mbRA. Generalized canonical analysis with a reference table, GCART (Kissita, 2003) fits within the

framework of generalized canonical analysis, whereas generalized concordance analysis, CONCORg (Lafosse and Ten Berge, 2006) and Orthogonal Multiple Co-Inertia Analysis, OMCIA (Vivien et al., 2005) fit within the framework of PLS regression. A main difference of these methods with mbPLS on the one hand and mbRA on the other hand, lies in the fact that these methods focus on the partial components rather than the global components. This is in particular reflected by the adopted deflation procedure which consists in deflating with respect to the vectors of loadings within each dataset (CONCORg) or the partial components (OMCIA and ACGTR). Therefore, within each dataset, the vector of loadings or the partial components are orthogonal, but not the global components. We believe that the global components give more insight into the problem under study and give valuable tools both for the prediction and the investigation of the relationships among datasets as will be illustrated in the next section (Westerhuis and Smilde, 2001).

#### 3. Applications

For the purpose of comparing the performances of the methods, we apply multiblock PLS regression, multiblock Redundancy Analysis and the continuum approach to a simulation study and to a real dataset pertaining to the field of veterinary epidemiology.

#### 3.1. Simulation Study

A simulation study is conducted in order to investigate the performance of the three methods under study, e.g., mbPLS, mbRA and mbCR. A simplified model is specified which involves three datasets  $X_1, X_2$  and Y with two variables per dataset. The conditions considered in this simulation study are the size of multicolinearity among the variables in  $X_k$  and the sample size. The multicolinearity within  $X_1$  and  $X_2$  is set to be identical and varied at three levels (low, Cor = 0.1; medium, Cor = 0.5; high, Cor = 0.9). The average correlation between variables in  $X_k$  and Y is set to 0.3. Furthermore, five different sample size are considered (N = 15, 25, 50, 100, 200). At each level of the experimental conditions, i.e., the three levels of multicolinearity times the five levels of the sample sizes, one hundred samples are randomly generated. The methods mbPLS, mbRA and mbCR are applied to each sample. The performance of these three methods is evaluated on the basis of a cross-validation procedure, described in Section 2.5, repeated one hundred times. Fitting ability  $(RMSE_C)$  and prediction ability  $(RMSE_V)$  are computed using respectively the calibration set and the validation set. As these measures express a lack of fit, the smaller they are, the better the method of analysis is. Moreover, for mbCR, the optimal value of the tuning parameter  $\gamma$  is automatically selected in each sample in accordance with the procedure described in Section 2.5. The average value of  $\gamma$  is given for each level of the experimental conditions. Results for all methods under the different conditions, for a model based on the first global component, are given in Table 1.

Fitting ability ( $R_c = RMSE_C$ ) and prediction ability ( $R_v = RMSE_V$ ) obtained from multiblock PLS (mbPLS), multiblock Re

D II	undancy Analys or $mbCR$ , the s	sis (mbKA) and average optimal	dundancy Analysis ( $mbKA$ ) and multiblock Continuum Kedundancy . For $mbCR$ , the average optimal tuning parameter ( $\gamma_{\text{opt}}$ ) is also given	ntinuum Kedunger (Yopt) is also	dancy PLS reg	ression (mbC/R	) under different	dundancy Analysis ( $mbKA$ ) and multiblock Continuum Kedundancy $PLS$ regression ( $mbCR$ ) under different simulation conditions. For $mbCR$ , the average optimal tuning parameter ( $\gamma_{\mathrm{opt}}$ ) is also given	tions.
	$mbRA \ (\gamma = 0)$	0)		$mbPLS \ (\gamma = 1)$	= 1)		Continuum $mbCR$ ( $\gamma_{\mathrm{opt}}$ )	$CR\left(\gamma_{ m opt} ight)$	
	Cor = 0.1	Cor = 0.5	Cor = 0.9	Cor = 0.1	Cor = 0.5	Cor = 0.9	Cor = 0.1	Cor = 0.5	Cor = 0.9
N = 15	$R_c = 0.66$	$R_c = 0.72$	$R_c = 0.73$	$R_c = 0.67$	$R_c = 0.71$	$R_c = 0.75$	$R_c = 0.67$	$R_c = 0.71$	$R_c = 0.74$
	$R_v = 0.79$	$R_v = 0.84$	$R_v = 0.88$	$R_v = 0.77$	$R_v = 0.81$	$R_v = 0.83$	$R_v=0.75$	$R_v = 0.79$	$R_v=0.81$
							$\gamma_{\rm opt} = 0.65$	$\gamma_{ m opt} = 0.71$	$\gamma_{\rm opt} = 0.61$
N = 25	$R_c = 0.71$	$R_c = 0.76$	$R_c = 0.78$	$R_c = 0.71$	$R_c = 0.76$	$R_c = 0.79$	$R_c = 0.71$	$R_c = 0.76$	$R_c = 0.79$
	$R_v = 0.80$	$R_v = 0.86$	$R_v = 0.88$	$R_v = 0.79$	$R_v = 0.83$	$R_v = 0.85$	$R_v = 0.78$	$R_v = 0.83$	$R_v = 0.84$
							$\gamma_{ m opt} = 0.65$	$\gamma_{\rm opt} = 0.74$	$\gamma_{ m opt}=0.64$
N = 50	$R_c = 0.75$	$R_c = 0.81$	$R_c = 0.83$	$R_c = 0.76$	$R_c = 0.81$	$R_c = 0.83$	$R_c = 0.75$	$R_c = 0.81$	$R_c = 0.83$
	$R_v = 0.81$	$R_v = 0.87$	$R_v = 0.89$	$R_v = 0.81$	$R_v = 0.86$	$R_v = 0.88$	$R_v = 0.80$	$R_v = 0.85$	$R_v = 0.87$
							$\gamma_{\rm opt} = 0.67$	$\gamma_{\rm opt} = 0.75$	$\gamma_{\rm opt}=0.70$
N = 100	$R_c = 0.77$	$R_c = 0.83$	$R_c = 0.86$	$R_c = 0.78$	$R_c = 0.83$	$R_c = 0.86$	$R_c = 0.78$	$R_c = 0.83$	$R_c = 0.86$
	$R_v = 0.81$	$R_v = 0.86$	$R_v = 0.90$	$R_v = 0.81$	$R_v = 0.86$	$R_v = 0.89$	$R_v = 0.81$	$R_v = 0.86$	$R_v = 0.89$
							$\gamma_{\rm opt} = 0.69$	$\gamma_{\rm opt}=0.80$	$\gamma_{\rm opt} = 0.71$
N = 200	$R_c = 0.79$	$R_c = 0.84$	$R_c = 0.87$	$R_c = 0.79$	$R_c = 0.84$	$R_c = 0.87$	$R_c = 0.79$	$R_c = 0.84$	$R_c = 0.87$
	$R_v = 0.81$	$R_v = 0.87$	$R_v = 0.90$	$R_v = 0.81$	$R_v = 0.86$	$R_v = 0.89$	$R_v = 0.81$	$R_v = 0.86$	$R_v = 0.89$
							$\gamma_{\rm opt} = 0.69$	$\gamma_{\rm opt} = 0.76$	$\gamma_{\mathrm{opt}} = 0.75$

Table 2

Comparison of methods with respect to their fitting and prediction ability under different simulation conditions: proportion of times that the methods, i.e., multiblock Redundancy Analysis (mbRA), multiblock PLS (mbPLS) or multiblock Continuum Redundancy PLS regression (mbCR), outperform each other

Compared N methods	Cor =	0.1 ab. Prediction ab.	Cor =	0.5 ab. Prediction ab.	Cor =	0.9 ab. Prediction ab.
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mbRA > mbPLS 15	83%	29%	81%	8%	90%	11%
25	78%	27%	75%	4%	91%	13%
50	84%	34%	71%	5%	84%	11%
100	83%	31%	75%	15%	89%	3%
200	73%	42%	73%	24%	87%	7%
mbCR > mbPLS 15	80%	100%	67%	100%	84%	100%
25	77%	100%	66%	100%	84%	100%
50	79%	100%	54%	100%	73%	100%
100	68%	100%	70%	100%	76%	100%
200	60%	100%	63%	100%	75%	100%
mbCR > mbRA 15	22%	100%	20%	100%	9%	100%
25	27%	100%	20%	100%	8%	100%
50	20%	100%	29%	100%	15%	100%
100	16%	100%	27%	100%	9%	100%
200	34%	100%	26%	100%	13%	100%

When the level of multicolinearity increases, regardless of the sample size and the method,  $RMSE_C$  and  $RMSE_V$  become also larger. In presence of high multicolinearity within the  $X_k$  datasets, the performance of mbRA, mbPLS and mbCR decreases. For mbCR, the average  $\gamma$  value is higher for a medium level of multicolinearity within  $X_k$ (Cor = 0.5) than for a high or a low level (Cor = 0.1) or Cor = 0.9. As expected, on the one hand, multiblock Redundancy Analysis has a better fitting ability than multiblock PLS especially when the level of multicolinearity is low. On the other hand, multiblock PLS has a better prediction ability for medium and high level of multicolinearity. The multiblock Continuum Redundancy PLS regression has a good and comparable fitting ability to mbRA. Moreover, whatever the size of the sample size and the level of multicolinearity, mbCR has a better fitting ability than mbPLS and mbRA. Results can also be viewed from a more general perspective by computing the number of times that the methods outperform each other (Table 2). It can be seen that, overall, mbRA has a better fitting ability than mbPLS and mbCR and, contrariwise, it is less effective insofar as the prediction is concerned. mbCR outperforms the other two methods in terms of prediction ability.

## 3.2. Case Study

The dataset consists in the measurements of several variables on 404 chicken flocks that were studied during rearing, transport and at slaughterhouse (Lupo *et al.*, 2008).

The Y table to be explained contains two quantitative variables which reflect the official reasons for condemnation at slaughterhouse, i.e., rate of infectious (INFECT) or traumatic (TRAUMA) origin. The explanatory table is organized in three blocks. Table  $X_1$  contains 26 variables pertaining to the rearing features. Table  $X_2$  contains 11 variables which refer to the transport conditions between farm and slaughterhouse. Table  $X_3$  contains 4 variables pertaining to the slaughtering conditions. The condition index computed for each explanatory table flags the presence of multicolinearity in  $X_1$ . Indicator (dummy) variables are considered for the categorical variables. Variables are column centred and scaled to unit variance.

The relationships between  $(X_1, X_2, X_3)$  and Y can be investigated using the global components  $(t^{(1)},\ldots,t^{(h)})$ . The graphical displays in Fig. 1 depict the loadings associated with the first two components  $t^{(1)}$  and  $t^{(2)}$  for mbRA ( $\gamma = 0$ ) and mbPLS regression  $(\gamma = 1)$ . It highlights the relationships among the explanatory variables from  $X_1, X_2$  and  $X_3$ , and makes it possible to identify some risk factors associated with the condemnation reasons (Y table). The graphical displays associated with mbRA and mbPLS show that the Y variables are strongly related to the first two components. For simplicity sake, we will only interpret the graphical display associated with mbRA. The condemnation rate at slaughterhouse for infectious reason (INFECT) is in particular associated with the age of the poultry house (ebanage in  $X_1$ ), the frequency of visits of the farmer to the poultry house during the starting period (EPassage in  $X_1$ ) and the standard chicken type (eilotyp5 in  $X_1$ ), among others. The condemnation rate at slaughterhouse for traumatic reason (TRAUMA) is in particular linked to the presence of an operator at the evisceration line (ievinbr in  $X_3$ ) and the average lairage time at the slaughterhouse (dattentemoyin  $X_2$ ), among others. This means that particular care with respect to these variables should be taken in order to reduce the number of carcasses which are condemned at slaughterhouse.

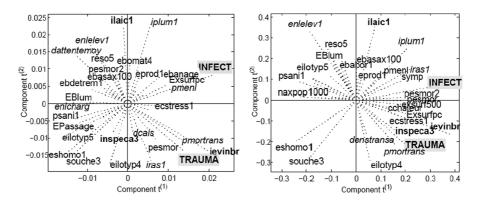


Fig. 1. Plots of the variable loadings associated with the first two components, for mbRA and mbPLS. 13 (resp. 14) variables that were not deemed important for the interpretation of the mbRA (resp. mbPLS) graphical display were discarded from the plot (although these variables were included in the analysis). Y variables are bold with a grey background,  $X_1$  variables are normal,  $X_2$  variables are slanted and  $X_3$  variables are bold.



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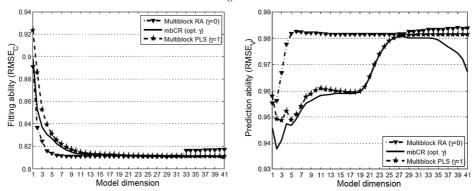


Fig. 2. Fitting and prediction ability as functions of the number of components introduced in the model. Comparison of mbCR (optimal parameter), mbRA ( $\gamma = 0$ ), mbPLS ( $\gamma = 1$ ).

The choice of the optimal model (i.e., optimal number of components and  $\gamma$  parameter) is a compromise achieved by both minimizing the root mean square error of calibration  $(RMSE_C)$  and validation  $(RMSE_V)$ , which respectively reflect the fitting ability and the prediction ability of the model under consideration. The cross-validation procedure is repeated (m = 200) times by setting one third of the individuals out and by varying the  $\gamma$  value from 0 to 1 with an increment of 0.01. We undertake a comparison of mbCR, mbRA and mbPLS on the basis of  $RMSE_C$  and  $RMSE_V$  criteria. Figure 2 shows  $RMSE_C$  and  $RMSE_V$  criteria as functions of the number h of components  $(t^{(1)}, \ldots, t^{(h)})$  introduced in the model. It can be seen in Fig. 2 that mbCR, mbRAand mbPLS have comparable fitting abilities, although mbRA slightly outperforms the other methods. Insofar as the prediction is concerned, the continuum approach mbCRoutperforms mbPLS and mbRA especially for the first four dimensions. We can notice that the performances of the methods under study, especially the fitting ability, depend on the number of components introduced in the model. It leads to the usual dilemma between performance (i.e., keep a large number of components) and parsimony (i.e., reduce the number of components). A compromise between a correct fitting and a good prediction ability makes the choice easier and leads to a model with two components for all the methods considered herein. The best method to predict Y from  $(X_1, X_2, X_3)$  is obtained by mbCR with two components. For a two-dimensional model, the median value from (m=200) cross-validations of the optimal  $\gamma$  parameter is 0.98. The median value is given because the optimal parameter distribution is bimodal: 17% of the  $\gamma$  value are comprised between 0 and 0.19 and 83% between 0.77 and 1. This kind of optimal parameter value, close to one of the bound of the continuum, is also found in other continuum methods (Gonzalez et al., 2008; Hwang, 2009).

The importance of each block  $X_k$  for explaining Y is reflected by the coefficients  $a_k^{(h)^2}$  for a given dimension h. For h components introduced in the model, the importance of the block  $X_k$  is given by the average value  $\overline{a_k}^{(1-h)^2}$  of the coefficients  $(a_k^{(1)^2},\ldots,a_k^{(h)^2})$  associated with dimensions  $(1,\ldots,h)$ . Table 3 gives the importance of the rearing features  $(X_1)$ , the transport conditions  $(X_2)$  and the slaughtering conditions  $(X_3)$  in the explanation of the official reasons for condemnation at slaughterhouse (Y).

Table 3

Importance of  $(X_1,X_2,X_3)$  in the Y explanation for the optimal model with (h=2) components. Comparison of the block weight of multiblock Redundancy Analysis  $(\gamma=0)$ , continuum mbCR  $(\gamma_{\rm opt.}=0.98)$  and multiblock PLS  $(\gamma=1)$ 

	$mbRA  (\gamma = 0)$	$mbCR  (\gamma_{\rm opt.} = 0.98)$	$mbPLS \ (\gamma = 1)$
$\sqrt[\infty]{a_1}^{(1-2)2}$	51%	51%	48%
$\%\overline{a_2}^{(1-2)2}$	39%	39.5%	44%
$\%\overline{a_3}^{(1-2)2}$	10%	9.5%	8%
Total	100%	100%	100%

As discussed above, a prediction model can be set up by regressing the Y variables on the basis of the first two global components. Table 4 gives a comparison of the regression coefficients obtained by mbRA, mbCR and mbPLS. We use the results of the (m=200) cross-validated regression coefficients in order to compute the standard deviations for the various coefficients. Each variable from X is considered to be significantly linked with each variable from Y when the 95% confidence interval associated with the regression coefficient does not contain zero. It turns out that 19 (46%) explanatory variables are interpreted in a same way whatever the method used. For example, the variable eprod1 (i.e., presence of other animal productions on the farm, in  $X_1$ ) is a risk factor both for the infectious and the traumatic reasons for all the methods under study. We can notice that 15 (37%) explanatory variables have a different interpretation when using mbPLS instead of mbRA or mbCR. For example, the variable denstransa (i.e., chicken density in crates, in  $X_2$ ) is highlighted as a risk factor for infectious reason only by mbPLS (positive regression coefficients). The X variables which most influence Y, especially the infectious reason (INFECT), are in particular the frequency of visits of the farmer to the poultry house during the starting period (EPassage, in  $X_1$ ), the area of the poultry house (exsurf 500, in  $X_1$ ) and factors related to whether the production is made with standard chicken (eilotyp5, in  $X_1$ ) or the presence of the farmer during bird crating (enlelev1, in  $X_2$ ). This means that particular care with respect to these variables should be taken in order to reduce the number of carcasses which are condemned at slaughterhouse.

# 4. Concluding Remarks

For the purpose of exploring and modelling the relationships between one block of variables Y and several blocks of explanatory variables  $(X_1, \ldots, X_K)$ , we propose an extension of Redundancy Analysis in order to improve the fitting ability of multiblock PLS regression. As mbPLS and mbRA are based on the same criterion to be maximized associated with different norm constraints, we also investigate a continuum approach, called multiblock Continuum Redundancy PLS regression (mbCR). The key feature of this approach is the shrinkage of the variance-covariance matrices  $(X'_k X_k)$  towards the identity

Table 4

Comparison of the regression coefficients of  $X=[X_1|X_2|X_3]$  on Y using two global components. X variables which have a regression coefficient with an asterisk have a significant link with Y

Block	Variable	$mbRA (\gamma = 0)$		$mbCR  (\gamma = 0.98)$		$mbPLS \ (\gamma = 1)$	
		INFECT	TRAUMA	INFECT	TRAUMA	INFECT	TRAUMA
$\overline{X_1}$	ebasax100	0.02	-0.05	0.03	-0.05	0.01	-0.04*
	cchaleur	0.04	0.01	0.04	0.01	0.04	0.04
	ebabor1	-0.04	-0.07*	-0.03	-0.08*	-0.01	-0.06*
	ebanage	0.10*	0.01	0.09*	0.01	0.01	-0.01
	ebchauf1	-0.04	-0.01	-0.05	-0.01	-0.02	-0.02
	ebdetrem1	-0.07*	-0.08*	-0.05	-0.08*	-0.02	-0.06*
	EBlum	-0.07*	-0.05*	-0.06*	-0.05	-0.02	-0.06*
	ebomat4	0.03	-0.07*	0.03	-0.07*	-0.01	-0.06*
	ecstress1	0.03	0.04	0.04	0.04	0.05*	0.05*
	edesins2	-0.08*	-0.01	-0.08*	-0.01	-0.03	-0.03
	eilotyp4	-0.08	0.06	-0.03	0.06	0.04	0.04
	eilotyp5	-0.19*	0.10*	-0.14*	0.10*	0.00	0.09*
	EPassage	-0.26*	0.00	-0.19*	-0.02	-0.05*	-0.07*
	eprod1	-0.13*	-0.05*	-0.12*	-0.06*	-0.05*	-0.07*
	ESatpres1	0.07	0.01	0.05	0.01	-0.02	0.01
	eshomo1	0.07	-0.03	0.07*	-0.04	0.01	-0.02
	estri1	-0.06	-0.02	-0.05	-0.02	0.02	-0.01
	exsurf500	-0.25*	-0.01	-0.24*	-0.01	-0.09*	-0.01
	exsurfpc	0.06	0.04	0.06	0.04	0.03*	0.02
	frac	-0.02	-0.01	-0.01	0.00	0.02	-0.01
	pesmor2	0.09	0.02	0.09*	0.02	0.06*	0.04*
	pesmor	0.11	0.04	0.09*	0.04	0.06*	0.06*
	psani1	0.06*	0.02	0.06*	0.02	0.03*	0.01
	reso5	-0.08	-0.04	-0.05	-0.03	0.03	0.00
	souche3	0.18	0.00	0.08*	0.03	0.06*	0.03
	symp	-0.03	0.07	0.04	0.04	0.06*	0.03
$X_2$	enlcais2	-0.11*	-0.05	-0.11*	-0.06	-0.07*	-0.08*
	enlcharg	0.00	-0.13*	0.00	-0.13*	-0.03	-0.09*
	enlelev1	-0.18*	-0.02	-0.17*	0.00	-0.07*	0.02
	iplum1	0.07*	0.05	0.07*	0.05	0.06*	0.02
	iras1	-0.05	-0.03	-0.05	-0.03	-0.04*	-0.03
	isolei1	-0.15*	-0.08*	-0.13*	-0.06*	-0.02	-0.06*
	pmenl	-0.04	-0.16*	-0.04	-0.16*	-0.04	-0.13*
	pmortrans	0.14*	-0.12	0.11*	-0.07	0.08*	-0.01
	denstransa	-0.04	0.14	0.00	0.08	0.07*	0.02
	dattentemoy	-0.04	-0.05	-0.04	-0.05	-0.03	-0.06*
	dcais	0.09*	0.01	0.09*	0.00	0.04*	0.00
$X_3$	ievinbr	0.08*	0.14*	0.07*	0.13*	0.04	0.09*
	ilaic1	-0.02	0.06*	-0.02	0.06*	0.01	0.05*
	inspeca3	-0.08	-0.13*	-0.07*	-0.10*	-0.05*	-0.06*
	naxpop1000	-0.03	0.03	-0.04	0.00	-0.05*	-0.03*

matrices. This continuum is easy to grasp and implement because the solutions are derived from an eigenanalysis of a matrix. The practical advantage of the mbCR approach lies in the fact that the tuning parameter makes it possible to explore a wide range of methods in order to find an optimal set of coefficients. This approach gives a unified framework so as to deal with potential multicolinearity problems.

The tuning parameter stands as a regularization parameter as it improves the conditioning of each matrix  $(X_k'X_k)$ . The optimal value of this parameter may be determined through a cross-validation procedure. From the simulation study, we show that multiblock Redundancy Analysis has a better fitting ability than multiblock PLS but has a lower prediction ability for medium and high level of multicolinearity. The continuum approach can be viewed as a ridge-type regularization of multiblock Redundancy Analysis. We show that whatever the size of the sample size and the level of multicolinearity, mbCR has similar or better fitting and prediction ability than mbPLS and mbRA. From the case study, we found that mbCR slightly outperforms mbPLS and mbRA. To summarize, we can advice to use mbCR in lieu of mbRA or mbPLS when the level of multicolinearity is high. When no multicolinearity occurs, the proposed multiblock Redundancy Analysis could be recommended.

Moreover, further research is needed in order to investigate more deeply the benefits gained from introducing the regularization procedure considering that it entails the cost of introducing a new parameter. Another topic for future research is to investigate the connection between the tuning parameter and the number of components to be introduced in the model. Different tuning parameters  $(\gamma_1, \ldots, \gamma_K)$  could also be included in the model, depending of the level of multicolinearity within each datasets  $(X_1, \ldots, X_K)$ .

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# Nuo daugiablokių mažiausių kvadratų iki daugiablokės pertekliškumo analizės: kontinumo metodas

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Daugiablokis dalinis mažiausių kvadratų (DMK) metodas yra dažnai taikomas regresiniuose uždaviniuose, tiriant ir modeliuojant sąryšius tarp duomenų bazės ir kelių duomenų bazių. Šis metodas yra DMK, susiejančio dvi duomenų bazes, apibendrinimas. Darbe yra pasiūlytas DMK plėtinys daugiablokėje formuluotėje. Parodyta, kad daugiablokis DMK ir daugiablokė pertekliškumo analizė maksimizuoja tą patį kriterijų skirtingais ribojimais. Pasirodo, abu sprendiniai priklauso tai pačiai kontinumo aibei, tiriamai darbe.