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Control chart and data fusion for varietal origin discrimination: Application to olive oil

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Combining data from different analytical sources could be a way to improve the performances of chemometric models by extracting the relevant and complementary information for food authentication. In this study, several data fusion strategies including concatenation (low-level), multiblock and hierarchical models (mid-level), and majority vote (high-level) are applied to near- and mid-infrared (NIR and MIR) spectral data for the varietal discrimination of olive oils from six French cultivars by partial least square discriminant analysis (PLS1-DA). The performances of the data fusion models are compared to each other and to the results obtained with NIR or MIR data alone, with a choice of chemometric pre-treatments and either an arbitrarily fixed limit or a control chart decision rule. Concatenation and hierarchical PLS1-DA fail to improve the prediction results compared to individual models, whereas weighted multiblock PLS1-DA models with the control chart approach provide a more efficient differentiation for most, but not all, of the cultivars. The high-level models using a majority vote with the control chart decision rule benefit from the complementary results of the individual NIR and MIR models leading to more consistently improved results for all cultivars.

1. Introduction

The increasing availability of multivariate data from various analytical sources and large numbers of samples makes it necessary to develop statistical tools that can extract the relevant information from these big datasets. For this purpose, a variety of chemometric models have been developed to predict either quantitative or qualitative properties. In the field of food authentication, the goal is often to determine if the actual characteristics of the samples agree with the information provided by their label. Thus, supervised methods can be used to classify new samples as authentic or non-authentic based on the known characteristics of previous samples. Several approaches have been developed including class-modeling algorithms, which focus on the similarities within a class, and discriminant analysis algorithms, which focus on the differences between classes [1,2]. One of the most relevant methods in this latter approach is the partial least square discriminant analysis (PLS-DA), which has been applied to various authentication cases [3–6] but could be optimized to better deal with complex datasets, as pointed out in a recent review [7]. Amongst the proposed improvements, the choice of a decision rule is of great importance to better interpret the results. In a previous article, a new decision rule based on the control chart approach was developed and allowed better predictions of olive oil varietal origin using data from

chromatographic analyses [8]. Here, this control chart rule is tested on PLS1-DA models using infrared spectroscopic data to confirm its efficiency.

Moreover, since a lot of different analytical techniques can be applied to assess the authenticity of food products [2,9], combining data from complementary analyses is expected to improve the performances of the statistical models. Data fusion strategies are divided into three categories: low-level, mid-level and high-level [10]. Low-level fusion consists in the simple concatenation of the matrices containing the data from the different sources, followed by the analysis of the concatenated data by the chosen chemometric model [11–13]. However, this method suffers from the very large number of variables in the concatenated matrix with the risk of increasing noise which may cancel out the advantages of adding sources of information. To solve this issue, mid-level fusion uses a first step of dimension reduction to extract the relevant information from the original data matrices and only the selected features are then combined to build the chemometric model [14–17]. Finally, high-level fusion builds separate models on each original dataset and these prediction results are then combined for the final decision where the class assignment is made according to probability rules or majority vote [18–20]. Previous studies have reported mixed outcomes from the use of data fusion. In most cases the results were improved by low-level [11–13,15,16,20], mid-level [15,16,20] or high level [18–20]

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strategies, but some studies have also reported that data fusion failed to improve the results compared to models built with individual datasets [14,16,17].

In this article, one low-level strategy with concatenation, two mid-level strategies with hierarchical and multiblock models, and one high-level strategy with majority vote are applied to the discrimination of olive oil varietal origin by PLS1-DA using near- and mid-infrared (NIR and MIR) spectral data. The performances of the data fusion models using either arbitrary fixed threshold or control chart decision rules are compared to each other and to individual models using only NIR or MIR data.

2. Material and methods

2.1. Olive oil samples

A total of 230 samples from six monovarietal extra-virgin olive oils produced over three harvest years (2016, 2017 and 2018) were used for this study. The samples came from olives picked manually in several orchards and whose cultivars were clearly identified: *Aglandau* (AG, $n = 63$), *Cailletier* (CA, $n = 29$), *Olivière* (OL, $n = 30$), *Picholine* (PI, $n = 35$), *Salonenque* (SA, $n = 39$) or *Tanche* (TA, $n = 34$). All the samples were produced using the same extraction method in the pilot plant of the professional association France Olive (Aix-en-Provence, France) and belonged to the extra-virgin olive oil category [21].

2.2. Near-infrared spectroscopy

FT-NIR spectra were obtained with an Antaris II spectrometer (Thermo Scientific, Waltham, MA, USA) in transmission mode, in a temperature-controlled room at 21 °C. The oil was poured in a QX Quartz Suprasil 300 cell (Hellma Analytics, Mülheim, Germany) with an optical path of 2 mm and an empty quartz cell was used to take a background reference before each measurement. Between each sample, the quartz cell was cleaned with isoctane, dried with air, rinsed with dichloromethane and dried again with air. Each spectrum was recorded between 10000 and 4500 cm^{-1} by the accumulation of 16 scans with a resolution of 4 cm^{-1} . The analysis was repeated two times for each sample and the resulting spectra were averaged. The NIR range between 10000 and 6100 cm^{-1} , a noisy area containing non-informational or redundant absorbances, was not included in the chemometric models. Thus the remaining NIR spectra consisted of 831 variables between 6100 and 4500 cm^{-1} .

2.3. Mid-infrared spectroscopy

FT-MIR spectra were obtained using a Nicolet Avatar spectrometer (Thermo Scientific, Waltham, MA, USA) with a nitrogen-cooled MCT detector, Ever-Glo source and KBr/Ge beam splitter. The measurements were conducted in a temperature-controlled room at 21 °C and air was taken as a background reference before each spectrum. A drop of EVOO was placed on the diamond crystal of a Golden Gate ATR accessory (Specac, Orpington, UK). Its spectrum was recorded between 4000 and 600 cm^{-1} by the accumulation of 64 scans with a resolution of 4 cm^{-1} . The ATR plate was cleaned with ethanol before each acquisition. This process was repeated three times for each EVOO sample, and the three resulting spectra were averaged prior to data analysis. The MIR range between 4000 and 1800 cm^{-1} , a noisy area containing non-informational or redundant absorbances, was not included in the models. Thus the remaining MIR spectra consisted of 571 variables between 1800 and 700 cm^{-1} .

2.4. Chemometrics

Exploratory analyses were conducted with the Unscrambler X software (CAMO Software, Oslo, Norway), to visualize the repartition of

the samples with a principal component analysis (PCA) [22,23]. The spectra were pre-treated with either standard normal variate (SNV) or Savitzky-Golay (SG) first derivative (polynomial order 2; 3 smoothing points) to improve the signal to noise ratio and reduce the undesirable additive or multiplicative effects. The outliers identified on the influence plot of the PCA were removed (with SNV: 1 *Aglandau*, 2 *Cailletier*, 1 *Olivière*, 1 *Picholine*, 1 *Salonenque*, 1 *Tanche* / with SG first derivative: 1 *Aglandau*, 4 *Cailletier*, 1 *Olivière*, 1 *Picholine*, 1 *Salonenque*). For the rest of the study, the best pre-treatment was selected based on the performance of each prediction model.

Multivariate statistical analyses were performed by several variations of the partial least square – discriminant analysis (PLS1-DA) [3] developed with Matlab R2014b software (The MathWorks, Natick, MA, USA). For each model, the samples were assigned a binary coding indicating if they belonged (value of 1) or not (value of 0) to the modelled cultivar. Two thirds of the samples from each cultivar and each harvest year were randomly selected to compose a calibration set and the remaining third served as a validation set to test the predictive abilities of the models. With the fixed threshold decision rule [24], a sample was considered as belonging to the modelled cultivar if its predicted value was between 0.6 and 1.4, belonging to the other cultivars if predicted between -0.4 and 0.4, and suspect if predicted outside of these boundaries. With the control chart decision rule [8], warning and control limits were calculated at respectively 95% and 99% confidence intervals around the mean calibration scores of the modelled cultivar. A sample was accepted as belonging to the modelled cultivar if its predicted value was inside the warning limits, rejected if predicted outside the control limits, and suspect if predicted between the warning and control limits. The performances of the models were evaluated by three quality parameters, namely sensitivity, specificity and area under the receiver operating curve (AUC) [25]. Moreover, to avoid over-fitting, the calibration included a full leave-one-out cross-validation procedure in order to select the optimal number of latent variables (LV) as the lowest number of LV giving an AUC of cross-validation over 90%.

First, individual PLS1-DA models were developed using either the NIR or the MIR data. Then, several data fusion strategies were applied.

- Low-level:

NIR and MIR data were appended in a single matrix and PLS1-DA models were developed using this concatenated dataset.

- Mid-level:

- o Hierarchical models (PLS-PLS1-DA): individual NIR and MIR datasets were subjected to a first step of dimension reduction by PLS1-DA. Then the scores obtained with the optimal number of LV for each individual model were combined and used to develop the final PLS1-DA model [15].
- o Multiblock models (MB-PLS1-DA): for each iteration the PLS scores were calculated from the individual NIR and MIR data blocks, weighted according to their number of variables, and combined into a « super-matrix » used in the final PLS1-DA step. The super-matrix and the two individual blocks were deflated using the « super-scores » instead of the individual scores [26].

- High-level:

Separate PLS1-DA models were developed with the NIR and MIR data, and the minimum, maximum and average of the predicted values from both models were included in a majority vote to reach the final prediction [18]. For instance, with the fixed threshold, a sample with a predicted value of 0.55 with the NIR model (suspect) and 0.67 with the MIR model (accepted) would have an average value of 0.61 (accepted) and by combining these three values the sample would be accepted as belonging to the modelled cultivar after majority vote. On the contrary, if the predicted values were 0.55 with NIR (suspect) and 0.63 with MIR (accepted), giving an average value of 0.59 (suspect), the majority vote would classify this sample as suspect.

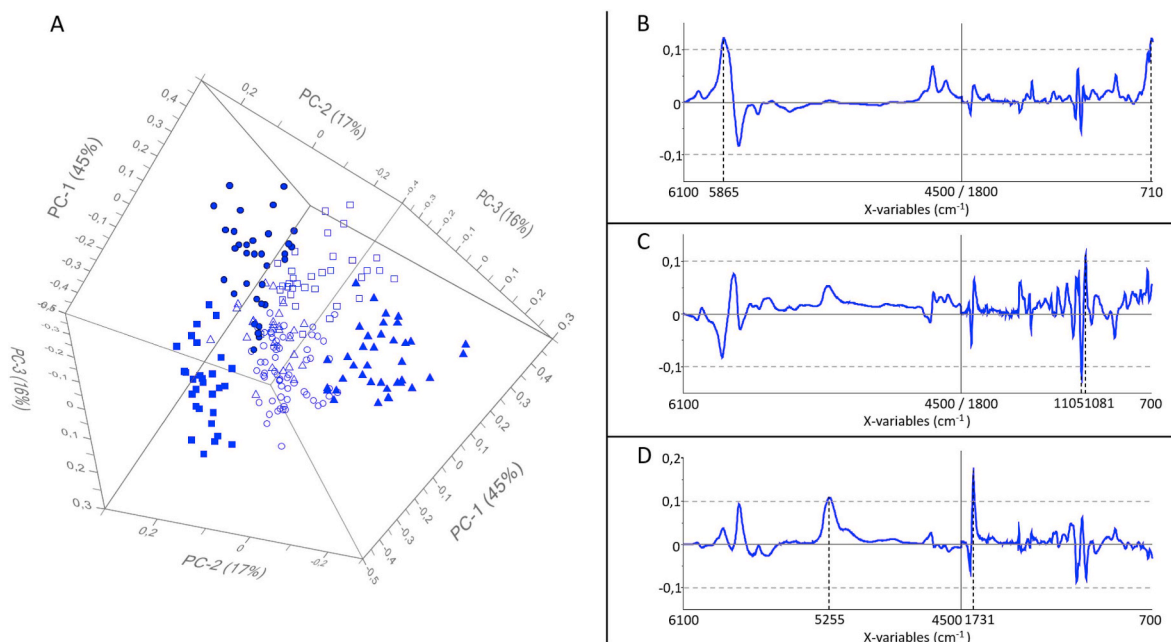


Fig. 1. A: PCA scores, B: loadings for PC1, C: loadings for PC2 and D: loadings for PC3, obtained with the concatenated NIR (6100–4500 cm^{-1}) and MIR (1800–700 cm^{-1}) data after SNV, with samples represented according to their cultivar on the scores plot (○: *Aglandau*, △: *Cailletier*, ■: *Olivière*, □: *Picholine*, ▲: *Salonenque*, ●: *Tanche*) and most influential bands identified on the loadings plots.

In order to explore the influence of both NIR and MIR data and their respective contribution to the discrimination of the studied olive oil varietal origin, variable importance in projection (VIP) values were calculated for each data fusion model using the formula from Mehmood et al. [27]. Since the average of the squared VIP values is equal to 1, variables with a VIP greater than 1 are usually considered as more relevant to the model, but some studies suggest that this threshold could vary [27,28].

3. Results and discussion

The NIR and MIR spectra obtained from all the studied olive oil samples, with identification of the spectral bands, are shown as Supplementary Information (Figure A).

3.1. Exploratory analysis

Fig. 1 presents the scores (Fig. 1-A) and loadings (Fig. 1-B, 1-C and 1-D) of the PCA on concatenated NIR and MIR data after SNV pre-treatment. The best separation according to cultivars was obtained using the first three PCs. The first component (PC1) represents 45% of the information. It is strongly influenced by the NIR band around 5865 cm^{-1} that corresponds to the first overtone of C–H bond vibrations ($-\text{CH}_3$, $-\text{CH}_2$) and could be related to the degree of unsaturation of triacylglycerols ($-\text{CH}=\text{CH}-$), as well as by the MIR band around 710 cm^{-1} that is attributed to fundamental C–H bending vibrations [29]. PC1 separates OL and AG samples which have mostly negative scores, from PI, SA and TA samples which have mostly positive scores on this component, and CA samples which have medium values. The second component (PC2) represents 17% of the information and is more influenced by MIR data, especially by the region between 1105 and 1081 cm^{-1} attributed to the fundamental vibrations of C–O (ester) and C–C bonds, and could also be influenced by the degree and type of unsaturation of fatty acids [29]. PC2 separates TA, OL and CA samples which have mostly positive scores, from SA samples which have negative scores, while AG and PI samples have rather medium values on this component. The third component (PC3) represents 16% of the information. It is mostly influenced by the NIR band around 5255 cm^{-1}

and the MIR region around 1731 cm^{-1} , which can be attributed to vibrations of the C=O bond of carbonyl groups [29]. PC3 separates SA and CA samples which have mostly positive scores from TA samples which have mostly negative scores while AG, OL and PI have medium values. Thus, even though the groups of cultivars are overlapping on these three PCs, chemometric models should be able to discriminate the samples according to their varietal origin. Samples from the AG, CA and PI cultivars may be more difficult to identify since their characteristics place them in the middle of all the other cultivars.

3.2. PLS1-DA on individual data

The results of the prediction models based on the individual NIR (A) or MIR (B) datasets for each cultivar, with the pre-treatment (SNV or SG first derivative) giving the highest AUC, are presented in Fig. 2. The values of sensitivity, specificity and AUC of the best models are also detailed in Table 1. More results with confusion matrices for all the models are available in the Supplementary Information (Tables A and B).

Using the fixed limit decision rule, MIR data gives similar or better results than NIR for most of the cultivars, except *Aglandau*. Indeed, the models built with NIR data have optimal numbers of LV between 7 for *Olivière* and *Salonenque* and 12 for *Cailletier* and *Picholine*, and reach AUC between an acceptable value of 0.86 for *Tanche* and an excellent value of 1.00 for *Salonenque*. The models using MIR data have optimal numbers of LV ranging from 6 for *Salonenque* to 13 for *Cailletier*, and their AUC all reach satisfying values from 0.90 for *Aglandau* and *Tanche* to 1.00 for *Picholine* and *Salonenque*.

Using the control chart decision rule reduces the difference between the results obtained with each source of data, homogenizes the performances of prediction between the 6 cultivars, and can also moderate the optimal number of LV for some models. Regarding the models using NIR data, the control chart approach improves the sensitivity but slightly decreases the specificity. Nevertheless, the overall AUC values are improved for most of the cultivars compared to the fixed limit, with satisfying values between 0.91 for *Olivière* and 0.99 for *Cailletier*, while lower optimal numbers of LV are used (from 4 for *Olivière* to 12 for *Picholine*). The control chart approach brings less improvement to the

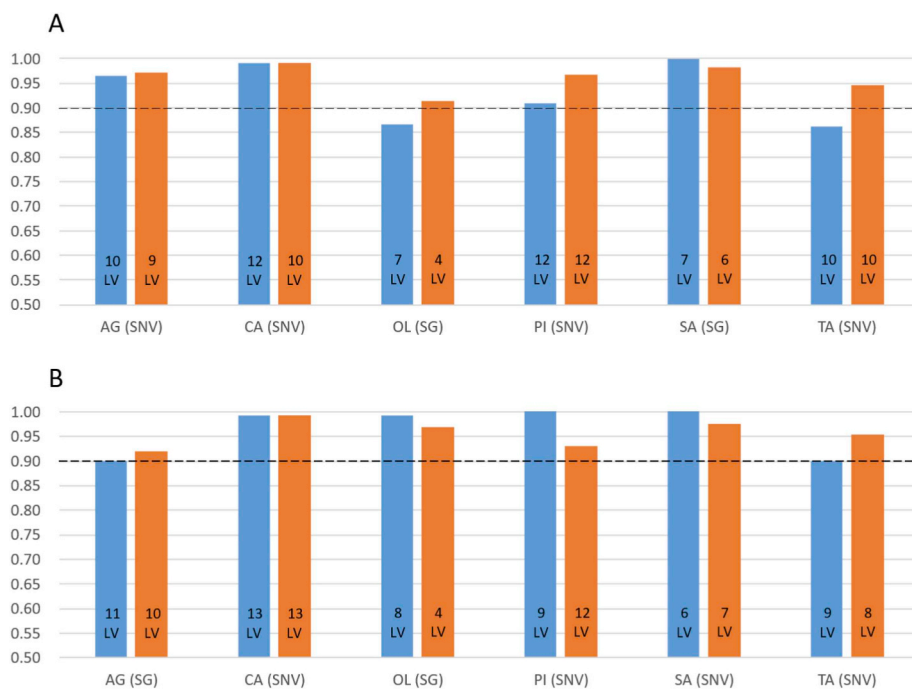


Fig. 2. AUC values for the PLS1-DA models using A: NIR data or B: MIR data, with the best pre-treatment, optimal number of LV, fixed limit (blue, left) or control chart (orange, right) decision rules, for each predicted cultivar (AG: *Aglandau*, CA: *Cailletier*, OL: *Olivière*, PI: *Picholine*, SA: *Salonenque*, TA: *Tanche*). (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

Table 1

Performance parameters of the PLS1-DA models using NIR or MIR datasets with the best pre-treatment and with fixed limit or control chart decision rule (SENS: sensitivity, SPEC: specificity, AUC: area under the curve, AG: *Aglandau*, CA: *Cailletier*, OL: *Olivière*, PI: *Picholine*, SA: *Salonenque*, TA: *Tanche*).

		NIR			MIR		
		SENS	SPEC	AUC	SENS	SPEC	AUC
AG	Fixed Limit	95%	98%	0.97	86%	94%	0.90
	Control Chart	100%	94%	0.97	95%	89%	0.92
CA	Fixed Limit	100%	98%	0.99	89%	97%	0.93
	Control Chart	100%	98%	0.99	100%	98%	0.99
OL	Fixed Limit	75%	98%	0.87	100%	98%	0.99
	Control Chart	88%	95%	0.91	100%	94%	0.97
PI	Fixed Limit	82%	100%	0.91	100%	100%	1.00
	Control Chart	100%	94%	0.97	91%	95%	0.93
SA	Fixed Limit	100%	100%	1.00	100%	100%	1.00
	Control Chart	100%	97%	0.98	100%	95%	0.98
TA	Fixed Limit	73%	100%	0.86	82%	98%	0.90
	Control Chart	91%	98%	0.95	91%	100%	0.95

models based on MIR data. However, the results remain very satisfying with AUC between 0.92 for *Aglandau* and 0.99 for *Cailletier*, and with also lower optimal numbers of LV between 4 for *Olivière* and 13 for *Cailletier*.

With each data source and each decision rule, the high number of LV required for the models predicting the *Cailletier* and *Picholine* cultivars confirms the observation from the exploratory analysis regarding the difficulty of their discrimination. Moreover, even though the performances of the models using either NIR or MIR data are close for most of the cultivars, the main differences can be observed for *Aglandau*, which is better predicted with NIR, whereas *Olivière* is better predicted with MIR.

3.3. PLS1-DA with data fusion

NIR and MIR spectra contain some redundant but also some complementary information and combining both data is expected to improve the prediction of olive oil cultivars. The results of the prediction models with the pre-treatment giving the highest AUC are presented in

Fig. 3, while sensitivity, specificity and AUC values are detailed in **Table 2**. Confusion matrices for all the models are available in the Supplementary Information (**Tables C to F**).

3.3.1. Low-level concatenation

Applying simple PLS1-DA to a low-level concatenated dataset containing both NIR and MIR data does not bring much improvement to the results compared to the individual datasets (**Fig. 3-A**).

Indeed, with the fixed limit decision rule the optimal numbers of LVs are still ranging from 7 for *Olivière* and *Salonenque* to 13 for *Cailletier*, and the AUC values are between 0.90 for *Aglandau* and 0.98 for *Olivière*. The only improvement is obtained for the model predicting the *Tanche* cultivar, with an AUC of 0.94 instead of 0.90 with MIR and 0.86 with NIR data only.

The control chart decision rule improves the sensitivity for most cultivars, but decreases the specificity for some, resulting in overall higher AUC values for *Aglandau*, *Cailletier* and *Tanche* but lower values for *Olivière* and *Salonenque*. The AUC remain satisfying for all the cultivars, between 0.93 for *Salonenque* and 0.99 for *Cailletier*, even though only the prediction of the *Tanche* cultivar is improved compared to the models using individual NIR or MIR data.

The variable importance in projection (VIP) scores for the PLS1-DA models using concatenated NIR and MIR data with the control chart decision rule (Supplementary Information - Figure B) indicate that both sources provide some useful information, but the most important variables differ depending on the modelled cultivar. For instance, NIR data seems to have more influence on the model predicting the *Olivière* cultivar, whereas MIR data brings more information to the model predicting *Cailletier*. The most influential bands for each model can also be identified [29]. For the NIR spectra part, the area bringing the most information to all the models is between 5910 and 5750 cm^{-1} corresponding to the first overtone of C-H vibrations. Its influence is even stronger for the *Salonenque*, *Olivière* and *Tanche* cultivars. The region between 5290 and 5220 cm^{-1} , related to the combination of O-H vibrations from water, also plays a part in most models except for the *Olivière* and *Salonenque* cultivars. The region between 4685 and 4645 cm^{-1} , attributed to the combination of C-H vibrations from unsaturation, brings additional information, mostly for the *Olivière* and *Tanche* cultivars. As for the MIR spectra part, the two most influential

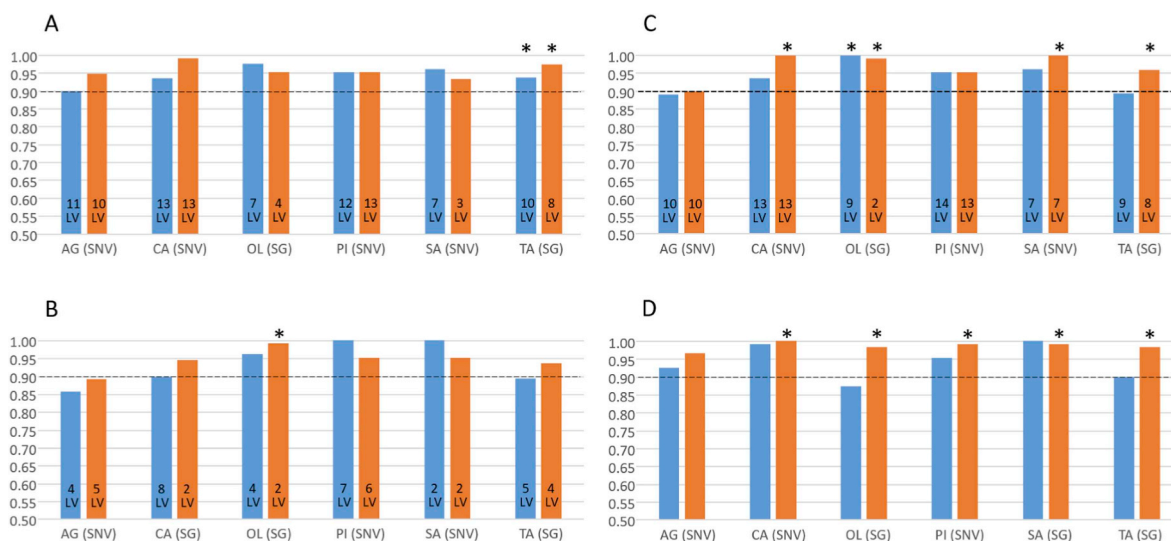


Fig. 3. AUC values for the data fusion models using A: concatenation, B: hierarchical, C: multiblock or D: majority vote, with the best pre-treatment, optimal number of LV, fixed limit (blue, left) or control chart (orange, right) decision rules, for each predicted cultivar (AG: *Aglandau*, CA: *Cailletier*, OL: *Olivière*, PI: *Picholine*, SA: *Salonenque*, TA: *Tanche*). A star (*) indicates that the results are better than with only NIR or MIR data. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

Table 2

Performance parameters of the PLS1-DA models using data fusion strategies with the best pre-treatment and with fixed limit or control chart decision rule (SENS: sensitivity, SPEC: specificity, AUC: area under the curve, AG: *Aglandau*, CA: *Cailletier*, OL: *Olivière*, PI: *Picholine*, SA: *Salonenque*, TA: *Tanche*).

		Concatenation			Hierarchical			Multiblock			Majority Vote		
		SENS	SPEC	AUC	SENS	SPEC	AUC	SENS	SPEC	AUC	SENS	SPEC	AUC
AG	Fixed Limit	80%	100%	0.90	75%	96%	0.86	80%	98%	0.89	85%	100%	0.93
	Control Chart	90%	100%	0.95	80%	98%	0.89	80%	100%	0.90	95%	98%	0.97
CA	Fixed Limit	89%	98%	0.94	89%	91%	0.90	89%	98%	0.94	100%	98%	0.99
	Control Chart	100%	98%	0.99	100%	89%	0.95	100%	100%	1.00	100%	100%	1.00
OL	Fixed Limit	100%	95%	0.98	100%	92%	0.96	100%	100%	1.00	75%	100%	0.88
	Control Chart	100%	91%	0.95	100%	98%	0.99	100%	98%	0.99	100%	97%	0.98
PI	Fixed Limit	91%	100%	0.95	100%	100%	1.00	91%	100%	0.95	91%	100%	0.95
	Control Chart	91%	100%	0.95	100%	90%	0.95	91%	100%	0.95	100%	98%	0.99
SA	Fixed Limit	92%	100%	0.96	100%	100%	1.00	92%	100%	0.96	100%	100%	1.00
	Control Chart	100%	87%	0.93	100%	90%	0.95	100%	100%	1.00	100%	98%	0.99
TA	Fixed Limit	91%	97%	0.94	82%	97%	0.89	82%	97%	0.89	82%	98%	0.90
	Control Chart	100%	95%	0.98	100%	87%	0.94	100%	92%	0.96	100%	97%	0.98

areas are between 1760 and 1710 cm^{-1} , related to C=O stretching vibrations, for all the cultivars but *Salonenque*, and between 1200 and 1020 cm^{-1} , attributed to C-O deformation vibrations, especially for *Salonenque*. Other important information comes from the 1520-1430 cm^{-1} region, attributed to C-H deformation vibrations, which has less influence for *Picholine*, as well as 925-905 cm^{-1} , that could be related to *cis* unsaturation, which is less useful for the *Olivière* and *Tanche* cultivars, 845-800 cm^{-1} attributed to C-C deformation vibrations, which is more useful for the *Aglandau* and *Salonenque* cultivars, and finally 750-700 cm^{-1} , attributed to C-H deformation vibrations, which is less useful for the *Olivière* and *Tanche* cultivars. Very similar VIP scores are obtained with the fixed limit decision rule. However, the lack of improvement observed with the low-level concatenation model may be due to the fact that the relevant information is concealed in the very large dataset, since only a few of the 1402 variables are really useful in the discrimination of the cultivars. Thus, mid-level models with an additional step condensing the appropriate information could have better prediction abilities.

3.3.2. Mid-level hierarchical PLS-PLS1-DA

PLS-PLS1-DA models, using the scores obtained with the optimal numbers of LV from NIR and from MIR individual models as a reduction dimension step, do not lead to better prediction results for most of the

cultivars (Fig. 3-B).

With the fixed limit, the final numbers of LV are between 2 for *Salonenque* and 8 for *Cailletier* and the AUC values for prediction are between 0.86 for *Aglandau* and 1.00 for *Picholine* and *Salonenque*. With the control chart, the final numbers of LV are reduced, between 2 for *Cailletier*, *Olivière* and *Salonenque* and 6 for *Picholine*. The sensitivity is improved for most models, but at the expense of the specificity, resulting in AUC values between 0.89 for *Aglandau* and 0.99 for *Olivière*, which is the only improvement compared to individual NIR and MIR models (Table 2).

The VIP scores of the PLS-PLS1-DA models with the control chart decision rule indicate both sources of information are used, although NIR data is more important than MIR data in all these models (Supplementary Information – Figure C). A similar pattern is observed with the fixed limit decision rule. This imbalance between the contributions of the two datasets could be corrected by another mid-level approach using weighted multiblock models.

3.3.3. Mid-level multiblock MB-PLS1-DA

Applying a more complex mid-level MB-PLS1-DA algorithm gives somewhat similar results to concatenation with the fixed limit decision rule but brings more improvement with the control chart decision rule (Fig. 3-C).

Indeed, using the fixed limit the optimal numbers of LV range from 7 for *Salonenque* to 14 for *Picholine* and the AUC values for prediction are between 0.89 for *Aglандаu* and *Tanche* and 1.00 for *Olivière*. This latter model is the only one for which the performances are improved compared to the individual NIR or MIR models. On the other hand, with the control chart approach the numbers of LV are reduced, between 2 for *Olivière* and 13 for *Cailletier* and *Picholine*, while the AUC values are improved for most of the cultivars, ranging from 0.90 for *Aglандаu* to 1.00 for *Cailletier* and *Salonenque*. The predictions for the *Cailletier*, *Olivière*, *Salonenque* and *Tanche* cultivars are also better than with the models using either NIR or MIR data alone.

The global VIP scores of the MB-PLS1-DA models with the control chart decision rule show that, contrary to the PLS-PLS1-DA models, the most important information comes from the MIR data block for all the multiblock models (Supplementary Information – Figure D). This difference is especially important for the models predicting the *Aglандаu* and *Picholine* cultivars, for which the results are not improved, confirming that a good balance between the amount of information from the two data blocks is important to achieve better predictions.

3.3.4. High-level majority vote

The high-level models, based on a majority vote between the predicted values from the individual NIR and MIR models and their average, give a good compromise for all the cultivars, especially with the control chart decision rule (Fig. 3-D). Since the fusion takes place at the decision level in these models, the optimal numbers of LV are those used for the separate NIR and MIR models.

When using the fixed limit, the AUC values of prediction are intermediate between those of the individual NIR and MIR models, ranging from 0.88 for *Olivière* to 1.00 for *Salonenque*. However, with the control chart, better performances than those of the individual models can be reached for all the cultivars except *Aglандаu*. In this case, excellent AUC values are obtained, ranging from 0.97 for *Aglандаu* to 1.00 for *Cailletier*. This fusion method gives the best overall results and is also the easiest to implement.

4. Conclusion

NIR and MIR spectral data can be used separately to discriminate monovarietal olive oils from six French cultivars, and the performance of the models are influenced by the choice of chemometric pre-treatment and decision rule. Indeed, the control chart decision rule allows for overall more satisfying results with less heterogeneity between the modelled cultivars. Combining NIR and MIR data does not always improve the performances of the models. When using low-level concatenation, the complementary information appears to be lost in the very large number of variables. Even mid-level hierarchical PLS1-DA, with a dimension reduction first step, fails to improve the prediction results. Using mid-level weighted multiblock PLS1-DA could be more appropriate, especially with the control chart decision rule, provided that a good balance of information from the two data blocks can be reached. However, this strategy gives heterogeneous results, improving the discrimination for most cultivars but worsening it for others. Finally, the high-level models using a majority vote benefit from the complementary results of the individual NIR and MIR models. This data fusion strategy, associated with the control chart decision rule, achieves better and more homogeneous predictions for all the cultivars and is also more user-friendly than the mid-level strategies.

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