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Study of phenolic extractability in grape seeds by means of ATR-FTIR and Raman spectroscopy



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ABSTRACT

Near infrared hyperspectral imaging has been applied to grape seeds in order to select a representative subset of samples according to their spectral features in the 900–1700 nm range. Afterwards, selected grape seeds have been classified according to their total phenol and flavanol extractabilities. In this way, samples were sorted in three different groups identified as low, medium and high extractability levels.

In order to establish the chemical structures which can be responsible for the different extractabilities, vibrational spectroscopy has been applied to the non-extracted material after seed extractions. Attenuated total reflectance Fourier transform infrared (ATR-FTIR) and Raman spectra of non-extracted seed material have been recorded and their main spectral features have been linked to extractabilities of flavanolic and total phenolic compounds.

The vibrational spectroscopic analysis confirms that grape seed phenolic extractability is influenced by the cell wall composition (polysaccharides, lignins, pectins) and by the degree of esterification of pectins.

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1. Introduction

Although grape seeds represent only 0–6% of berry weight, they effect the sensory properties of wine. Depending on varieties, they contain between 20 and 55% of the phenolic compounds of the berry (Ribéreau-Gayon et al., 2006). Phenolic compounds are characterized by having at least one aromatic ring with one or more hydroxyl groups attached. Flavonoids are the most numerous amongst phenolics. The generic structure of the major flavonoids comprises fifteen carbons (C6–C3–C6), with two aromatic rings (C6) connected by a heterocyclic pyran ring of three-carbon bridge (C3). Flavonoids are secondary metabolites that have important and varied roles (Crozier, Clifford, & Ashihara, 2006), including well-known health benefits. They possess ideal structural

Abbreviations: ATR-FTIR, attenuated total reflectance Fourier transform infrared; DAD, diode array detector; DMACA, 4-dimethylaminocinnamaldehyde; IR, infrared; H, Mahalanobis distance; MSC, multiplicative scatter correction; NEM, non-extracted material; NH, neighborhood Mahalanobis distance; NIR, near infrared; NIRS, near infrared spectroscopy; PC, principal component; PCA, principal component analysis.

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chemistry for free radical-scavenging activities, and they have been shown to be more effective antioxidants *in vitro* than vitamins E and C, on a molar basis (Rice-Evans, Miller, & Paganga, 1997).

The main subclasses of flavonoids present in grapes (*Vitis vinifera* L.) are flavonols, flavanols and anthocyanins, and they are transferred to the wine during the fermentation process. Flavonols and anthocyanins are principally localized in the skin, whereas flavanols are synthesized primarily in seeds and stems (Jackson, 2000). Flavanols constitute the most abundant phenolic family in grape seeds and play a relevant role in the sensory characteristic of red wines. They can increase the wine body, but also the sensations of astringency and bitterness (Waterhouse, 2002). Moreover, flavanols, like the main phenolic families, can increase and stabilize the wine color by means of the copigmentation phenomenon (Boulton, 2001; Gordillo et al., 2015; Nogales-Bueno, Baca-Bocanegra, Jara-Palacios, Hernández-Hierro, & Heredia, 2017).

The importance of controlling the amount of phenolic compounds that may be extracted from grapes to wine lies in the fact that the phenolic content wine depends mainly on the amount of phenolic compounds released from grape to wine. It is well known that the extractabilities of phenolic compounds from the solid parts of grapes (seeds and skins) depends on the grape maturity.

However, in skins, phenolic extractability increases with maturity, whereas, in seeds it decreases. (Ribéreau-Gayon et al., 2006).

The effect of cell wall degradation on the degree of extraction of phenolic compounds from grape skins has already been widely studied. Several studies have been developed using this matrix in order to control and understand the relationship between cell wall composition and the extraction degree of phenolic compounds. An increase in the sugar content (degree of ripening) has been correlated with decreasing amounts of cell wall material, galactose, cellulose and mannose, accompanied by a decrease in the degree of methylation of pectic polysaccharides or pectins (Ortega-Regules, Ros-García, Bautista-Ortín, López-Roca, & Gómez-Plaza, 2008). These factors could be responsible for the different extractabilities of phenolic compounds, bearing in mind that differences on thickness or density of the skin cell-wall could also play a role.

Regarding grape seeds, extractability decreases during ripening, from about 25% at the first stages to about 5% in the later (Rodríguez-Pulido et al., 2014). It might be inferred that this phenomenon could be attributed to the presence of the middle integument, which is waterproof and very hard when seeds ripen and lignify, and could prevent the extraction (Cadot, Minana-Castello, & Chevalier, 2006). On the other hand, changes in the cell wall polysaccharide structure could affect the solubility of the flavanols (Bautista-Ortín et al., 2013). These authors found that maceration enzymes, normally used in the wine elaboration process to degrade skin cell wall in order to increase color, can also favour the extraction of flavanols from the seeds. These oenological products easily degrade the cell walls of seeds (especially if they are unripe and show low lignification), promoting the release of seed compounds into the must-wine, increasing the wine body, but also the sensations of astringency and bitterness. It is therefore of interest to evaluate how cell wall structure changes among seed samples of different grape varieties or with different levels of phenolic compounds extractability.

Near infrared spectroscopy (NIRS) has been used in order to screen total or extractable phenolic compounds in grapes obtaining quite good results (Ferrer-Gallego, Hernández-Hierro, Rivas-Gonzalo, & Escribano-Bailón, 2011; Kempf, Leon, Best, De Baerdemaeker, & De Ketelaere, 2010; Nogales-Bueno, Baca-Bocanegra, Rodríguez-Pulido, Heredia, & Hernández-Hierro, 2015; Nogales-Bueno, Hernández-Hierro, Rodríguez-Pulido, & Heredia, 2014; Rodríguez-Pulido et al., 2014; Torchio, Río Segade, Giacosa, Gerbi, & Rolle, 2013). However, it is not possible to easily interpret the relationship between cell wall composition and phenolic compounds extractability based on near infrared (NIR) region features. The near infrared region contains absorption bands corresponding to overtones and combinations of fundamental C—H, O—H and N—H vibrations. Additionally, the presence of Fermi resonances can also increase the complexity of the NIR spectra. Therefore, many band assignments can only be tentative or unresolved which limits the usefulness of the near infrared region in structural determination. In comparison, mid infrared spectra exhibit sharp and narrow peaks essentially related to fundamental modes of vibration which can be easily assigned to chemical structures (Bokobza, 1998). Mid infrared spectroscopy has previously been used in order to relate skin cell wall composition to anthocyanin extractability (Femenia, Sánchez, Simal, & Rosselló, 1998; Hernández-Hierro et al., 2014). In these studies, mid infrared spectral features could be directly linked to the esterification of pectins, which might be responsible for the different extractabilities of the aforementioned phenolic compounds. Moreover, this spectroscopic technique has been applied to the study and quantitation of phenolic compounds extracted during winemaking (Di Egidio, Sinelli, Giovanelli, Moles, & Casiraghi, 2010; Fragoso, Acena, Guasch, Mestres, & Busto, 2011). However, in these cases, cell wall composition has not been taken into account.

Cell wall structure of grape and other fruits has been widely studied by means of vibrational spectroscopy. Vibrational spectroscopy is a subset of spectroscopy which comprises IR and Raman spectroscopy and analyses vibrations within a molecule (or material) (Byrne et al., 2014). These two vibrational spectroscopic techniques are complementary. Whereas electric dipole transitions of IR (and UV–visible) absorption require a change of the dipole moment of the material as a result of the transition, Raman scattering requires a change in the polarizability of the bond as a result of the transition. Thus, access to molecular level information by means of two different physical processes can be achieved (Byrne, Sockalingum, & Stone, 2011).

Mid infrared spectroscopy has been employed to study cell wall structure of grapes and other fruits (Acebes et al., 2014; Fasoli et al., 2016; Kyomugasho, Christiaens, Shpigelman, Van Loey, & Hendrickx, 2015; Wilson et al., 2000; Zietsman et al., 2015). However, studies which employ mid infrared spectroscopy to study the cell wall structure of grapes seeds have not been performed. Raman spectroscopy has also been employed to study cell wall structure in fruits, although, no studies have been found for grapes (Chylińska, Szymańska-Chargot, & Zdunek, 2014; da Silva, Vandenabeele, Edwards, & Cappa de Oliveira, 2008; Gierlinger, Keplinger, & Harrington, 2012; Szymańska-Chargot et al., 2016). Therefore, although both mid infrared and Raman spectroscopy have proved to be useful and reliable techniques for the study of the cell wall structure in plant and fruit tissues, they have not yet been applied to the study of grape seeds.

In the present study, near infrared hyperspectral imaging has been applied to grape seed samples in order to select a representative subset of samples according to their near infrared spectral features. Afterwards, total phenol and flavanol extractabilities have been obtained for these selected samples and they were sorted into three different groups, identified as low, medium and high extractability levels. Next, Raman and attenuated total reflectance Fourier transform infrared (ATR-FTIR) spectra have been collected for non-extracted material after grape seed extractions. Finally, ATR-FTIR and Raman spectra have been analyzed in order to relate their more important features to phenolic extractability levels in grapes seeds. Moreover, spectral features are also related to the principal sample attributes. To our knowledge, this is the first time that ATR-FTIR and Raman spectroscopy are jointly applied to grape seed non-extracted material samples in order to relate their more important spectral features to their phenolic extractability levels.

2. Material and methods

2.1. Samples

V. vinifera L. cv. Syrah and Tempranillo red grape samples were collected from two vineyards located in the Condado de Huelva Designation of Origin D.O. (Andalusia, Spain). Grapes were collected on two different dates, when Tempranillo and Syrah vineyards were respectively harvested (August 7 and 11, 2014). A total of two hundred Syrah and Tempranillo grapes were collected, one hundred per variety. In order to achieve representative samples sets, single grapes were collected from the top, middle and bottom of the cluster and from the sunlight and shade side. The samples were refrigerated and immediately transported to the laboratory.

2.2. Sample selection

Grape seeds were removed from the whole grapes and left to dry at room temperature. Then individual near infrared hyperspectral images were recorded for grape seeds belonging to the same

grape and the near infrared spectrum was saved (between 950 and 1650 nm) as described elsewhere (Rodríguez-Pulido et al., 2014). Afterwards, sample selection was carried out as described in Nogales-Bueno et al. (2015). Briefly, an unsupervised pattern recognition technique, principal component analysis (PCA), was used to select representative samples from the spectral data set. Eight principal components were taken into account in order to explain ninety per cent of the spectral variability of original spectral matrix. Mahalanobis distances (H) for each sample were calculated and samples were grouped according to a neighborhood H criterion ($NH \leq 0.6$). Thus, 66 groups with different spectral characteristics were created and one sample from every group was selected. These 66 selected samples were used in all the subsequent analyses. Grape seeds were weighed, immediately frozen and stored at -20°C until analyses were performed.

2.3. Model wine and exhaustive extractions

Two different extractions were carried out: a model wine extraction and an exhaustive extraction.

For model wine extraction, grape seeds were immersed in a model wine hydroalcoholic solution (4 g L^{-1} tartaric acid, 12.5% (v/v) ethanol, adjusted at pH 3.6 with NaOH 0.5 M) for a maceration period of 72 h. Then, supernatants were used into the subsequent analyses.

Afterwards, grape seed samples were freeze-dried, grounded and macerated in methanol:water 75:25 (v/v), sonicated during 15 min (JP Selecta, Barcelona, Spain) and centrifuged ($830\times g$, 15 min). These extractions were repeated twice in order to achieve an exhaustive extraction of phenolic compounds. The methanolic extracts were combined and finally made up to a final volume of 50 mL with methanol.

As a by-product of model wine and exhaustive extractions, non-extracted material (NEM) was obtained from each grape seed sample. These NEM samples were freeze-dried and then stored in a desiccator until further use.

2.4. Total phenol and flavanol extractability determinations

Total phenol and flavanol extractabilities were determined for grape seed samples. They were determined for each supernatant coming from the model wine and exhaustive extractions. Then, extractabilities of each sample were evaluated, respectively, as the fractions of total phenols and flavanols extracted by the model wine solution with respect to the exhaustive extraction. Finally, grape seed samples were sorted according to their phenolic extractability levels expressed as percentages.

Total phenol contents were determined using the Folin–Ciocalteu method (Singleton & Rossi, 1965). Two hundred microliters of exhaustive or model wine extractions were mixed with 1.5 mL of sodium carbonate (20% w/v), 500 μL of Folin reagent and made up to 10 mL with ultrapure water.

Flavanol contents were determined following a modification of the method of Vivas, Glories, Lagune, Saucier, and Augustin (1994). Ten or twenty microliters of exhaustive or model wine extractions were mixed with 190 or 180 μL of methanol respectively and 1 mL of DMACA (4-dimethylaminocinnamaldehyde) reagent. The DMACA reagent was prepared immediately before use, containing 0.1% (w/v) DMACA in a mixture of HCl:methanol (1:10, v/v).

Both Folin and DMACA analyses were performed on an Agilent 8453 UV–visible spectrophotometer (Palo Alto, USA), equipped with diode array detection (DAD), measuring absorbance at 765 and 640 nm respectively. The extract volumes were appropriately modified for samples which needed it.

2.5. ATR-FTIR data collection

ATR-FTIR spectra were recorded with the Perkin Elmer (MA, USA) Spotlight 400 N Universal Attenuated Total Reflectance (UATR) accessory of the spectrometer, which employs a 9-bounce diamond top-plate for this analysis. Spectral data were the result of 32 scans, with a spectral resolution of 1 cm^{-1} and covering the spectral range between 600 and 4000 cm^{-1} .

All samples were measured with a force gauge of 140 units. NEM from grape seed samples were powders and 6 absorption spectra were collected for each grape. A background spectrum was also recorded and automatically subtracted by the software.

2.6. Raman data collection

A Horiba Jobin-Yvon LabRAM HR800 spectrometer with an external 300 mW diode laser operating at 785 nm as source was used throughout this work. For the measurements, a $\times 100$ objective (MPlanN, Olympus) was employed, providing a spatial resolution of $\sim 1\text{ }\mu\text{m}$ at the sample. The confocal hole was set at 100 μm , the specified setting for confocal operation. The system was spectrally calibrated to the 520.7 cm^{-1} spectral line of silicon and the intensity response function was corrected using the Standard Reference Material (SRM) No. 2243 of the National Institute of Standards, Boulder, Colorado, USA (NIST SRM 2243, 2242, 2241). The LabRAM system is a confocal spectrometer that contains two interchangeable gratings (300 and 900 lines per mm respectively). In the following experiments, the 300 lines per mm grating was used, giving a spectral dispersion of $\sim 1.5\text{ cm}^{-1}$ per pixel. The detector used was a 16-bit dynamic range Peltier cooled CCD detector.

All spectra were recorded over the spectral range between 400 and 3500 cm^{-1} and with a spectral resolution of $\sim 0.9\text{--}1.6\text{ cm}^{-1}$. Six spectra were collected for each NEM from grape seed samples.

2.7. Data analysis

K-means cluster analysis of near infrared hyperspectral data was carried out in order to sort grape seed samples according to their phenolic extractability levels (total phenol, flavanol extractabilities). Initial cluster centers were chosen in order to maximize the initial between-cluster distances. Samples were sorted into three groups, low, medium and high phenolic extractability levels. K-means analysis was carried out by means of Statistica v.8.0 software (StatSoft Inc., OK, USA, 2007).

Both ATR-FTIR and Raman raw data needed spectral pre-treatments. A multiplicative scatter correction (MSC) was applied to the ATR-FTIR raw data in order to remove the scattering caused mainly by particle size and compaction. Win ISI (v1.50) (Infrasoft International, LLC, Port. Matilda, PA, USA, 2000) software was used for this aim. Baseline correction was applied to Raman raw data. This correction was carried out using MATLAB R2012b (The Mathworks, Natick, MA, USA, 2012) and following the algorithm described elsewhere by Mazet, Carteret, Brie, Idier, and Humbert (2005). This algorithm estimates the background of a spectrum by means of a non-quadratic cost function. Afterwards, this cost function was subtracted from the raw spectra. Asymmetric truncated quadratic was the cost function which gives the best results to estimate background in Raman spectra of NEM from grape seed samples. Moreover, the backgrounds were estimated by a 5-order polynomial and with thresholds of 0.01.

Peak heights were calculated for ATR-FTIR and Raman pre-treated spectra and PCA was applied to both whole spectral and peak height matrices. Win ISI (v1.50) software was used for PCA of whole spectral matrices, whereas PCA for peak heights matrices was performed by means of Statistica v.8.0 software (StatSoft Inc., OK, USA, 2007). The aforementioned peaks heights for ATR-FTIR

and Raman were previously obtained from spectra and pre-treated spectra respectively using MATLAB R2012b (The Mathworks, Natick, MA, USA, 2012). Furthermore, univariate analysis of variance (Tukey *post hoc* test) was applied to find differences in the peak heights (dependent variables) among different factors (extractability levels or grape varieties). The statistically significant level was considered at $\alpha = 0.05$. Statistica v.8.0 software was used to develop these analyses.

3. Results and discussion

3.1. Total phenol and flavanol extractability levels

K-means cluster analysis of near infrared hyperspectral data sorted grape seed samples in three different groups. The analysis allocated samples into groups according to their total phenols and flavanols extractability levels and, afterwards, these groups were named as low, medium and high extractability levels. Table 1 shows the extractability levels of total phenols and flavanols for grape seed samples and the number of samples classified in each cluster by the k-means method.

3.2. ATR-FTIR data

Raw average spectra of NEM from grape seed samples with low, medium and high extractabilities are shown in Fig. 1A. MSC pre-

treatment was applied to ATR-FTIR spectra and average spectra were calculated for each grape seed NEM sample. Then, the spectra were vector normalized and mean centered. Afterwards, a PCA was applied to this spectral matrix. Overall, the spectral variability explained was 99% using 13 principal components and Mahalanobis distances for each sample were calculated. Samples were ranked in order of their H (Mahalanobis) distance from the mean spectrum of the entire sample set and the $H > 3$ criterion was applied in order to look for spectral outliers. Only one H-outlier was found, and it was not taken into account hereafter. Fig. 1B shows the scores of the NEM from grape seed samples in the space defined by the first and second principal components, which described 44.31% (PC1) and 35.72% (PC2) of the variability in the data. In the scores plot, the samples are represented by a color code indicating the different extractability levels. It can be seen that medium extractability samples overlap low and high extractability samples. In fact, if medium extractability samples are removed from this plot, it is possible to find some degree of separation between low and high extractability samples (Fig. 1C).

Fig. 1D shows the MSC pre-treated average spectra of NEM from grape seed samples with low and high extractabilities, which are quite similar. Moreover, Fig. 1D shows the loadings of PC1 and PC2. PC1 and PC2 loadings show large variations in several spectral regions which might influence the grape seeds phenolic extractability. For example, in the scores plot, almost all high extractability samples are located in the negative PC2 region

Table 1
Extractability levels of total phenols and flavanols for grape seed samples. Means and standard deviations.

Samples	n ^a	Extractabilities (expressed as percentage)			
		Total phenols		Flavanols	
		Mean	SD	Mean	SD
All	66	28.09	20.15	42.60	25.76
Low	25	8.57	5.21	15.25	9.63
Medium	28	31.16	7.72	50.45	11.03
High	13	59.04	12.54	78.28	7.24

^a Number of samples.

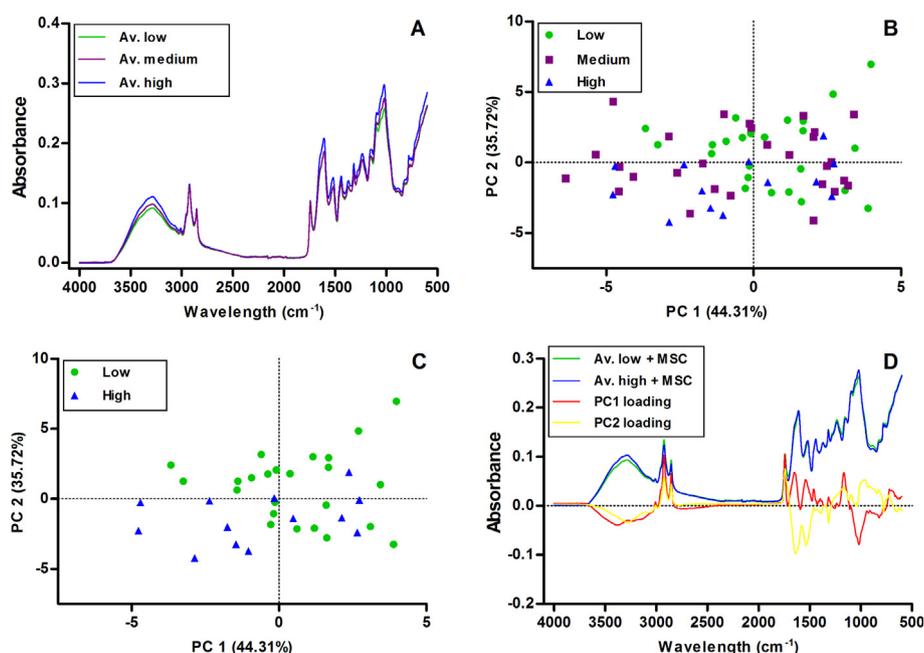


Fig. 1. (A) Raw average spectra of NEM from grape seed samples with low, medium and high extractabilities. (B) Score plot of the first two principal components after PCA performed on ATR-FTIR spectra recorded from grape seed NEM samples. The individual data points have been color coded according to the results of k-means cluster analysis; low, medium and high phenolic extractability. (C) Score plot without medium extractability samples. (D) PC1 and PC2 loading plot and average MSC pre-treated ATR-FTIR spectra for low and high phenolic extractability grape seed NEM samples.

Table 2
Main functional groups assigned to the different vibrations present in the ATR-FTIR spectra of non-extracted material (NEM) from grape seed samples.

Assignment ^a	Absorption bands (cm ⁻¹) (Intensity ^b)	Component	References
$\nu(\text{O-H})$	3285 (m, b)	Polysaccharides, lignins	Heredia-Guerrero et al. (2014), Lupoi, Singh, et al. (2015)
$\nu_a(\text{CH}_2)$	2924 (m)	Lignins, lipids	Fasoli et al. (2016), Lupoi, Singh, et al. (2015)
$\nu_s(\text{CH}_2)$	2854 (w)	Lignins, lipids	Fasoli et al. (2016), Lupoi, Singh, et al. (2015)
$\nu(\text{C=O})_{\text{ester}}$	1743 (m)	Polyesters, pectins, lignins	Fasoli et al. (2016), Szymanska-Chargot and Zdunek (2013), Wilson et al. (2000)
$\nu_a(\text{COO}^-)$ and $\nu(\text{C=C})_{\text{aromatic}}$	1610 (s)	Pectins, phenolics	Gao et al. (2015), Ricci et al. (2015), Szymanska-Chargot and Zdunek (2013), Wilson et al. (2000)
$\nu(\text{C-C})_{\text{aromatic}}$	1519 (w)	Phenolics	Heredia-Guerrero et al. (2014), Lupoi, Singh, et al. (2015)
$\nu(\text{C-C})_{\text{aromatic}}$	1441 (w)	Phenolics	Foo (1981), Heredia-Guerrero et al. (2014)
$\delta(\text{CH}_2)_{\text{out of plane}}$	1370 (vw)	Polysaccharides, pectins	Lupoi, Singh, et al. (2015), Szymanska-Chargot and Zdunek (2013), Wilson et al. (2000)
$\delta(\text{CH}_2)_{\text{scissoring}}$	1318 (w)	Polysaccharides, pectins	Szymanska-Chargot and Zdunek (2013), Wilson et al. (2000)
$\nu(\text{C-O})$ and $\delta(\text{O-H})$	1235 (w)	Polysaccharides, pectins, lignins	Lupoi, Singh, et al. (2015), Szymanska-Chargot and Zdunek (2013), Wilson et al. (2000)
$\nu(\text{C-H})_{\text{aromatic}}$	1154 (vw)	Phenolics	Foo (1981), Lupoi, Singh, et al. (2015)
$\nu(\text{C-O})$ and $\nu(\text{C-C})$	1020 (vs)	Polysaccharides, pectins	Lupoi, Singh, et al. (2015), Szymanska-Chargot and Zdunek (2013), Wilson et al. (2000)
$\delta(\text{CH}_2)_{\text{rocking}}$	781 (vw)	Phenolics	Foo (1981)

^a Assignment: ν , stretching; δ , bending; s, symmetric; a, asymmetric.

^b Intensity: vw, very weak; w, weak; m, medium; s, strong; vs very strong; b, broad.

(Fig. 1C), accordingly, the negative features in the PC2 loading (1250–1700 and 3000–3500 cm⁻¹ in Fig. 1D) might have significant influence on phenolic compound extractability. The main features presented for grape seed NEM samples are described in Table 2. These features are mainly ascribed to polysaccharides, lignins, lipids, pectins and phenolic compounds (Fasoli et al., 2016; Foo, 1981; Heredia-Guerrero et al., 2014; Lupoi, Singh, Parthasarathi, Simmons, & Henry, 2015; Szymanska-Chargot & Zdunek, 2013; Wilson et al., 2000).

With the aim of identifying what spectral regions have a major influence on phenolic extractability, a univariate analysis of variance was carried out. Peak heights were measured in MSC pre-treated grape seed NEM spectra. Thirteen peaks were taken into account (Table 2). A univariate analysis of variance, a statistical method used to analyze the differences among group means and their associated procedures, was performed. Peak heights were used as dependent variables, whereas extractability level, shown in Table 1, was used as factor or independent variable. This analysis is described in detail in Table 3, which shows the heights (mean and standard error) for all peaks described in Table 2. Moreover, in Table 3, values are marked with letters which indicate the existence or not of statistical differences between extractability levels. No significant differences were found for the peak at 1743 cm⁻¹, ascribed to (C=O)_{ester}, and for the peaks at 2854 and 2924 cm⁻¹,

Table 3
Peak heights (log(1/R)) for different levels of phenolic compounds extractability. Means \pm standard errors of means. For each peak, different letters in the same row indicate statistical differences (Tukey test, $\alpha = 0.05$).

Peak wavenumber (cm ⁻¹)	Extractability level		
	Low	Medium	High
3285	0.121 \pm 0.003 ^a	0.130 \pm 0.002 ^b	0.143 \pm 0.003 ^c
2924	0.168 \pm 0.003 ^a	0.163 \pm 0.003 ^a	0.162 \pm 0.003 ^a
2854	0.102 \pm 0.002 ^a	0.098 \pm 0.002 ^a	0.097 \pm 0.002 ^a
1743	0.109 \pm 0.002 ^a	0.109 \pm 0.002 ^a	0.112 \pm 0.002 ^a
1610	0.239 \pm 0.005 ^a	0.240 \pm 0.004 ^a	0.267 \pm 0.006 ^b
1519	0.150 \pm 0.005 ^a	0.153 \pm 0.004 ^a	0.172 \pm 0.004 ^b
1441	0.149 \pm 0.003 ^a	0.150 \pm 0.002 ^a	0.163 \pm 0.003 ^b
1370	0.127 \pm 0.002 ^a	0.129 \pm 0.002 ^a	0.140 \pm 0.003 ^b
1318	0.150 \pm 0.003 ^a	0.148 \pm 0.003 ^a	0.166 \pm 0.004 ^b
1235	0.167 \pm 0.002 ^a	0.167 \pm 0.002 ^a	0.179 \pm 0.003 ^b
1154	0.154 \pm 0.002 ^{ab}	0.152 \pm 0.002 ^a	0.162 \pm 0.002 ^b
1020	0.267 \pm 0.005 ^a	0.291 \pm 0.004 ^b	0.312 \pm 0.006 ^c
781	0.035 \pm 0.001 ^{ab}	0.034 \pm 0.001 ^a	0.038 \pm 0.001 ^b

ascribed to CH₂ stretches. Thus, they appear not to have any influence on phenolic compounds extractability. Significant differences ($p < 0.05$) were found for the peaks at 781 and 1154 cm⁻¹, but only

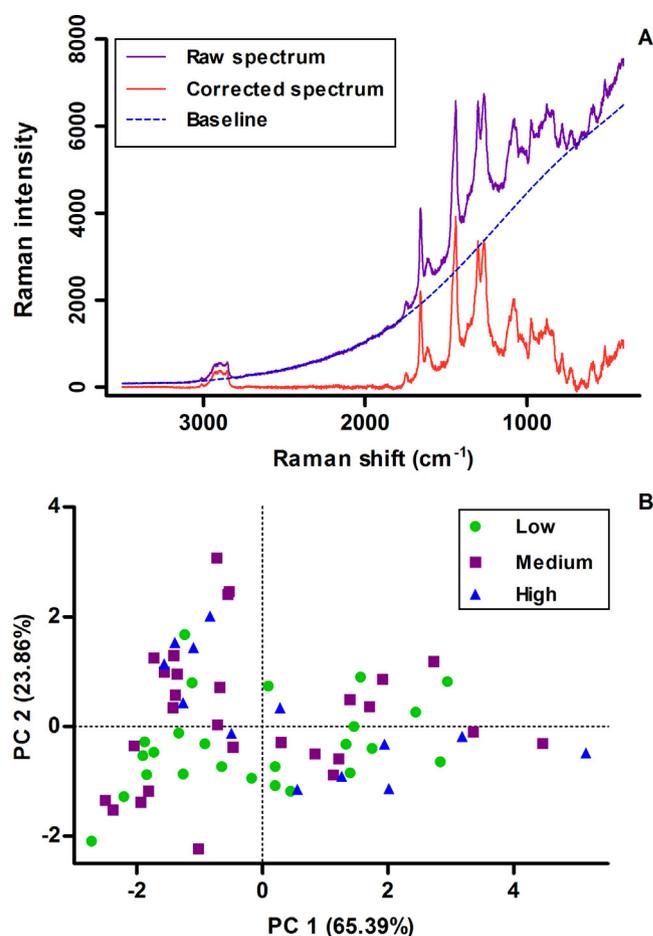


Fig. 2. (A) Raw Raman spectrum, baseline and corrected spectrum for a grape seed NEM sample. (B) Score plot of the first two principal components after PCA performed on peak heights of Raman spectra recorded from grape seed NEM samples. The individual data points have been color coded according to the results of k-means cluster analysis; low, medium and high phenolic extractability.

Table 4

Main functional groups assigned to the different vibrations present in the Raman spectra of non-extracted material (NEM) from grape seed samples.

Assignment ^a	Raman shifts (cm ⁻¹) (Intensity ^b)	Component	References
$\nu(\text{CH}_2)$	2916 (w, b)	Polysaccharides	Chylińska et al. (2014), Lupoi, Gjersing, et al. (2015), Szymańska-Chargot et al. (2016)
$\nu(\text{C}=\text{C})$	1655 (s)	Lignins, fatty acids	da Silva et al. (2008); Seidler-Lozykowska et al. (2010)
$\nu(\text{C}=\text{C})_{\text{aromatic}}$	1609 (w)	Lignins, phenolics	da Silva et al. (2008), Gierlinger and Schwanninger (2006), Seidler-Lozykowska et al. (2010), Sene, McCann, Wilson, and Grinter (1994)
$\delta(\text{CH}_2)_{\text{scissoring}}$	1442 (vs)	Lignins, lipids	Gierlinger and Schwanninger (2006), Lupoi, Gjersing, et al. (2015), Szymańska-Chargot et al. (2016)
$\delta(\text{=CH})_{\text{rocking}}$	1269 (vs)	Lipids	da Silva et al. (2008), Gierlinger and Schwanninger (2006), Lupoi, Gjersing, et al. (2015)

^a Assignment: ν , stretching; δ , bending.^b Intensity: w, weak; s, strong; vs very strong; b, broad.

between medium and high extractability levels. However, no significant differences were found between low and high extractability levels. These two peaks are related to phenolic compounds, and therefore, this fact can show some kind of differentiation between phenolic compounds in medium and high extractability samples, but not between these and low extractability samples. Finally, there are significant differences ($p < 0.05$) in the remaining peaks, 1020 and 3285 cm⁻¹, between the three extractability levels and between low/medium levels and high level for peaks at 1235, 1318, 1370, 1441, 1519 and 1610 cm⁻¹. It is to be noted that almost all these peaks fall within the spectral regions that principal component analysis showed to have significant influence on phenolic compound extractability. Despite the obvious fact that peaks related to the remaining phenolic compounds (1441 and 1519 cm⁻¹) (Foo, 1981; Heredia-Guerrero et al., 2014) influence phenolic extractability, these results confirm that phenolic extractability is also influenced by the cell wall composition (peaks at 1020, 1235, 1318, 1370 and 3285 cm⁻¹ ascribed to polysaccharides, lignins and pectins) and by the degree of esterification of pectins (1610 cm⁻¹, non-esterified pectins) (Gao, Fangel, Willats, Vivier, & Moore, 2015; Szymanska-Chargot and Zdunek, 2013; Wilson et al., 2000). However, this last peak is also ascribed to C=C aromatic stretching in phenolic compounds by other authors (Ricci, Olejar, Parpinello, Kilmartin, & Versari, 2015).

In addition, using grape variety as factor, a new analysis of variance was carried out and significant differences were found for peaks at 1020, 1370, 1441 and 3285 cm⁻¹ among Tempranillo and Syrah samples. All these peaks also have significant influence on the extractability of phenolic compounds. Therefore, these results show that ATR-FTIR spectral features are linked to grape variety and it can be confirmed that this spectroscopic tool can be used to study the relation between grape variety and phenolic compound extractability of grape seeds.

3.3. Raman data

In order to confirm or add more information to the above findings, Raman spectra were measured for NEM from grape seeds. A baseline correction procedure was carried out for each grape seed NEM spectrum (Fig. 2A) and 5 peaks were selected, those at 1269, 1442, 1609, 1655 and 2916 cm⁻¹ (Table 4) (Chylińska, Szymanska-Chargot, & Zdunek, 2014; da Silva et al., 2008; Gierlinger & Schwanninger, 2006; Lupoi, Gjersing, & Davis, 2015; Seidler-Lozykowska, Baranska, Baranski, & Krol, 2010; Szymanska-Chargot et al., 2016). These peaks were selected because they had a good signal to noise ratio and were present in most samples. Then, PCA was performed, both of whole spectra (data not shown) and peak heights (Lupoi, Singh, et al., 2015). The average peak heights were calculated for each sample and PCA was carried out using these data. Fig. 2B shows the scores of the NEM from grape

seed samples in the space defined by the first and second principal components, which described 65.39% (PC1) and 23.86% (PC2) of the variability in the data. In the scores plot, the samples are represented by a color code indicating the different extractability levels. No trends were found, as samples are overlapped in this space, and thus, PCA cannot be used to interpret the differences between phenolic extractability levels in this case.

Afterwards, one-way univariate analyses of variance were performed. Peak heights were used as dependent variables whereas grape seed variety and extractability levels (Table 1) were used as independent variables or factors in the different analyses. Significant difference ($p < 0.05$) among grape varieties was found for the peak at 1655 cm⁻¹. This peak is related to the C=C stretch in lignins and fatty acids. Moreover, significant differences were found for the peaks at 1609, 1655 and 2916 cm⁻¹ when extractability levels were taken into account. These bands are respectively due to phenolic compounds, lignins and polysaccharides (Gierlinger & Schwanninger, 2006; Lupoi, Gjersing, et al., 2015; Seidler-Lozykowska et al., 2010). The band at 1609 cm⁻¹, assigned to aromatic C=C skeletal stretching, has a strong Raman activity, whereas it only weakly active in infrared spectroscopy. However, the anti-symmetric stretching of COO⁻ only has infrared activity (Sene, McCann, Wilson, & Grinter, 1994). This allows an assignment of this Raman feature (1609 cm⁻¹) to phenolic compounds and not to pectins. Therefore, Raman spectroscopy enables confirmation of the linkage between cell wall components and phenolic extractability in grape seeds.

4. Conclusion

ATR-FTIR and Raman spectroscopy have been proven to be effective and reliable tools to relate the more important spectral features to phenolic extractability levels in grape seed and other sample attributes. ATR-FTIR data analysis confirmed that grape seed phenolic extractability is influenced by the cell wall composition (polysaccharides, lignins and pectins) and by the degree of esterification of pectins, despite the obvious fact that peaks related to the remaining phenolics compounds also influence on phenolic extractability. Moreover, the results obtaining from the infrared spectra have been confirmed by Raman spectroscopy. Peaks linked to cell wall composition have been shown to have significant influence on grape seed extractability.

Although FTIR spectroscopy has been applied to grape samples in several studies, to our knowledge, Raman spectroscopy had not been tested in grape seeds in order to study their cell wall structure. Raman spectroscopy has been shown to be a reliable tool for studying cell wall structure in grape seed tissue. Nonetheless, a comprehensive study should be made in order to improve spectra features, i.e., signal to noise ratio or the background produced by fluorescence.

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Conflict of interest

The authors declare no potential conflict of interest.

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