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Effect of the pig rearing system on the final volatile profile of Iberian dry-cured ham as detected by PTR-ToF-MS

J. Sánchez del Pulgar ^{a,b}, C. Soukoulis ^a, A.I. Carrapiso ^c, L. Cappellin ^a, P. Granitto ^d, E. Aprea ^a, A. Romano ^a, F. Gasperi ^a, F. Biasioli ^{a,*}

^a IASMA Research and Innovation Centre, Fondazione Edmund Mach, Food Quality and Nutrition Department, Via E. Mach, 1, 38010 S. Michele a/A, Italy

^b Food Technology, Facultad de Veterinaria, UEx, Campus Universitario s/n, 10003 Cáceres, Spain

^c Food Technology, Escuela de Ingenierías Agrarias, UEx, Carretera de Cáceres s/n, 06071 Badajoz, Spain

^d CIFASIS, French Argentina International Center for Information and Systems Sciences, UPCAM (France)/UNR-CONICET (Argentina), Bv 27 de Febrero 210 Bis, 2000 Rosario, Argentina

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ABSTRACT

The volatile compound profile of dry-cured Iberian ham lean and subcutaneous fat from pigs fattened outdoors on acorn and pasture (*Montanera*) or on high-oleic concentrated feed (*Campo*) was investigated by proton transfer reaction time-of-flight mass spectrometry. In addition to the usual proton transfer ionization the novel switchable reagent ions system was implemented which allows the use of different precursor ions (H_3O^+ , NO^+ and O_2^+). The analysis of the lean and subcutaneous fat volatile compounds allowed a good sample discrimination according to the diet. Differences were evident for several classes of compounds: in particular, *Montanera* hams showed higher concentrations of aldehydes and ketones and lower concentrations of sulfur-containing compounds compared to *Campo* hams. The use of NO^+ as precursor ion confirmed the results obtained with H_3O^+ in terms of classification capability and provides additional analytical insights.

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1. Introduction

Iberian dry-cured ham is a Spanish high-valued product, highly appreciated by consumers due to its unique sensory characteristics that depend on ripening conditions (Ruiz, Ventanas, Cava, Andrés, & García, 1999) and raw meat characteristics – i.e. animals' age, pig genotype and type of feed during the fattening period (Carrapiso, Bonilla, & García, 2003; Pérez-Palacios et al., 2010). Currently different commercial types of Iberian dry-cured hams are available according to the fattening of the pigs (R.D., 1469/2007). The most valuable dry-cured hams are produced from pigs fattened outdoors, with feeding based on acorn and pasture (*Montanera*), whereas the less expensive come from pigs fattened indoors on a concentrated feed. A new Iberian dry-cured ham type (*Campo*) produced from free range fattened Iberian pigs reared on a concentrated feed has been recently added to the Iberian ham commercial grading (R.D., 1469/2007), and several studies about these hams have been performed (Daza, Rey, Ruiz, & López-Bote, 2005; Ventanas, Ventanas, Tovar, García, & y Estévez, 2007).

Different methods have been proposed for the commercial classification of Iberian dry-cured ham, which analyse the subcutaneous fat (SCF) fatty acid profile (Ruiz et al., 1998) to the n-alkanes (Tejeda, Antequera, Martín, Ventanas, & García, 2001), n-alkenes and branched

hydrocarbons of the intramuscular fat (IMF) (Petrón, Tejeda, Muriel, Ventanas, & Antequera, 2005, 2006), the volatile hydrocarbon profile (Narváez-Rivas, Vicario, Alcalde, & León-Camacho, 2010) and the fatty acid profile of the IMF (Pérez-Palacios, Ruiz, Tejeda, & Antequera, 2009). Even if some of these methods, especially the IMF fatty acid profile analysis, allow good discrimination according to the fattening diet, all of them are complex and time-consuming techniques.

Proton transfer reaction mass spectrometry (PTR-MS) is an established method for the rapid, non-destructive detection of volatile organic compounds (VOCs) released from several food matrices or emitted during dynamic processes such as food processing or consumption (Aprea, Biasioli, Carlin, et al., 2007; Aprea, Biasioli, Gasperi, Märk, & van Ruth, 2006; Aprea, Biasioli, Gasperi, et al., 2007; Araghipour et al., 2008; Biasioli et al., 2003, 2006; Gasperi et al., 2009). The technique's fundamentals and its applications to food, environmental or medical purposes have been extensively reviewed (Biasioli, Gasperi, Yeretizian, & Märk, 2011; Biasioli, Yeretizian, Märk, Dewulf, & Van Langenhove, 2011; Blake, Monks, & Ellis, 2009). Very briefly, PTR-MS is based on the reaction of protonated water with VOCs molecules with a proton affinity higher than water's and the successive detection by a quadrupole mass spectrometer. Recently a PTR-MS coupled to a high resolution time-of-flight (ToF) mass analyser was commercialised (Jordan et al., 2009), which partially overcomes the limitations of the slower and low resolution quadrupole version, offering higher mass resolution ($m/\Delta m$ up to 5000), high sensitivity (pptv) and a higher time resolution (0.1 s) (Jordan

* Corresponding author.

E-mail address: franco.biasioli@iasma.it (F. Biasioli).

et al., 2009). PTR-ToF-MS has been successfully applied to the study of the volatile compounds of milk during fermentation (Soukoulis et al., 2010) and cheese (Fabris et al., 2010), and its capacity to discriminate dry-cured hams according to geographical origin and ripening processing has been also demonstrated (Sánchez del Pulgar et al., 2011).

Jordan et al. (2009) proposed a Switchable Reagent Ions (SRI) system which allows the use of different reagent ions in PTR-MS. This system introduces in PTR-MS apparatuses, without loss of sensitivity, a possibility that characterises competing techniques for VOCs detection such as the selected ion flow technique (SIFT-MS) (Spanel & Smith, 1999). Based on SRI, Jordan et al. (2009) demonstrated that the use of NO^+ and O_2^+ precursor ions in the PTR-MS apparatuses improves the analytical performance, particularly for the separation of isobaric compounds (i.e. aldehydes, 2-ketones and alpha-diketones) and for the detection of compounds with proton affinities lower than that of water (e.g. alkanes, alkenes). In general, the usefulness of NO^+ and O_2^+ as precursor ions for the analysis of volatile compounds in meat products has been demonstrated by SIFT-MS (Olivares et al., 2010, 2011).

For these reasons, it is proposed to use NO^+ and O_2^+ as reagent ions for the analysis of the volatile compound profile of dry-cured Iberian ham, to detect separately aldehydes and ketones (NO^+) and linear and branched alkanes (O_2^+), which could be useful to investigate the effect of rearing system and diet (Narváez-Rivas et al., 2010).

Therefore, the aims of this study were:

- to identify the possible effect of different rearing systems on the final volatile compounds of dry-cured Iberian hams by a rapid direct injection mass spectrometric technique.
- to set up discrimination methods for the classification of the different products and to obtain qualitative and quantitative analytical information on the volatile compounds useful for that discrimination.
- to evaluate, for the first time in food science, whether the use of different precursor ions (H_3O^+ , NO^+ and O_2^+) in PTR-MS apparatuses allows a better separation and/or provides better analytical information.

2. Material and methods

2.1. Ham samples

Twenty five Iberian dry-cured hams were obtained from twenty Iberian x Duroc 75% pigs (pure Iberian female x Iberian x Duroc male). Ten of these pigs were fattened outdoors on acorn and grass (*Montanera*), while the other fifteen pigs were fattened outdoors on a high-oleic concentrated feed (*Campo*), according to the regulations for Iberian pork products (R.D., 1469/2007). All the hams were simultaneously processed in the same way and ripened for 720 days according to García et al. (1996) and the regulations for Iberian pork products (R.D., 1469/2007) and Dehesa de Extremadura hams (Orden de 2 de Julio de, 1990).

2.2. Sample preparation

The analysis was carried out following the procedure described in Sánchez del Pulgar et al. (2011). From each ham a piece of the *Biceps femoris* muscle containing its corresponding subcutaneous fat (SCF) was taken, vacuum packaged and kept at 2 °C. Just before the analysis of the lean, the superficial layer and the visible fat of each piece of muscle was removed, and 3 cubes of 1 cm³, approximately 1.2 g lean (3 replicates) were prepared. The cubes were introduced into 40 ml Pyrex glass vials for analytical/chromatography use (Supelco, Bellefonte, USA), capped by PTFE/Silicone septa (Supelco, Bellefonte, USA). In the case of the subcutaneous fat, the surface of each piece

was also removed and 3 fat cubes of 1 cm³ (approximately 0.9 g) were prepared from each sample.

The same procedure was followed for the analysis with H_3O^+ , NO^+ and O_2^+ as reagent ions preparing three different sample series of lean.

2.3. Proton transfer reaction-time of flight-mass spectrometry (PTR-ToF-MS)

In order to standardise the measurement, the prepared vials were equilibrated at 37 °C for 30 min in a water bath prior to analysis. Next all the samples were analysed by direct injection of the head space mixture into the PTR-ToF-MS drift tube via a heated (110 °C) peek inlet for 30 s, taking 30 average spectra (Fabris et al., 2010). The PTR-ToF-MS analysis, using H_3O^+ as reagent ion was performed on the ten dry-cured *Montanera* hams and ten *Campo* hams. Due to limitation in sample availability, when using NO^+ and O_2^+ as reagent ion, the same hams measured with H_3O^+ were analysed except for five *Campo* hams, which were replaced with a different five *Campo* hams.

Measurements were carried out using a commercial PTR-ToF-MS 8000 apparatus (Ionicon Analytik GmbH, Innsbruck, Austria), in its standard configuration (V mode). The sampling time per channel of ToF acquisition was 0.1 ns, amounting to 350,000 channels for a mass spectrum ranging up to $m/z = 400$. The selected conditions in the drift tube depended on the primary ion and are reported in Table 1.

2.4. Spectra analysis

The external calibration automatically done by the acquisition program provided a poor mass accuracy, thus internal calibration of ToF spectra was performed off-line (Cappellin et al., 2010). The procedure allowed achievement of a mass accuracy generally better than 0.001Th for the considered mass range, which was in most cases sufficient for sum formula identification. Data pre-processing on ToF spectra was carried out to remove the baseline and noise reduction was achieved by averaging over the 30 consequent ToF spectra corresponding to the same sample, thereby improving the signal-to-noise ratio by about five times (Cappellin et al., 2011). Peak detection and area extraction were performed as described in Cappellin et al. (2011). Throughout this paper VOC concentrations are expressed in ppbv (parts per billion by volume) and were calculated from peak areas according to the formula described by Lindinger, Hansel, and Jordan (1998). A constant reaction rate coefficient of $2 \cdot 10^{-9}$ cm³/s was employed in the calculations. In the case of H_3O^+ as primary ion, this introduces a systematic error of up to 30% that can be accounted for if the actual rate coefficient is known (Cappellin et al., 2012). For NO^+ and O_2^+ the same formula was used but the employed constant rate coefficient cannot be safely assumed to be close to the one of the actual reaction and therefore results must be considered as being expressed in normalized counts.

Table 1

Drift tube conditions and counts per second (cps) of the primary ion isotopes during the analysis with each primary ion.

Primary Ion	Drift voltage (V)	Pressure (mB)	Temperature (°C)	E/N (Td)	Mass 21 ^a (cps)	Mass 31 ^b (cps)	Mass 34 ^c (cps)
H_3O^+	600	2.25	110	154	1744	5	86
NO^+	508	2.38	110	100	662	3116	218
O_2^+	608	2.05	110	137	380	247	4048

^a Mass corresponding to the M + 2 isotopologue of H_3O^+ .

^b Mass corresponding to the M + 1 isotopologue of NO^+ .

^c Mass corresponding to the M + 2 isotopologue of O_2^+ .

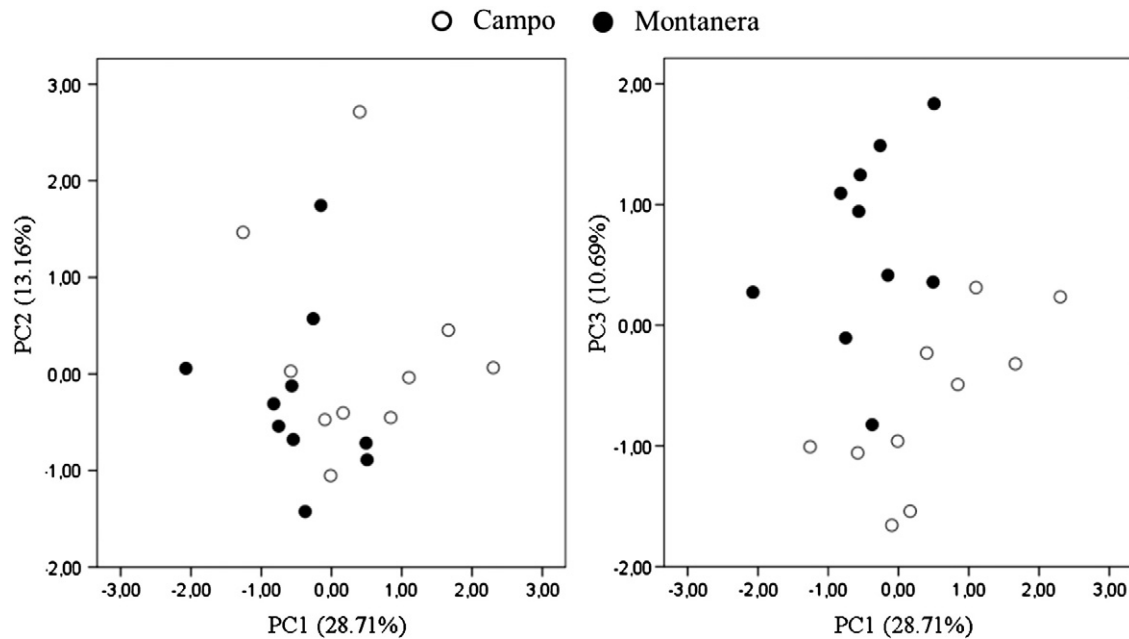


Fig. 1. Score plots obtained by the PCA analysis (PC1 vs PC2 and PC1 vs PC3 respectively) of the PTR-ToF-MS H_3O^+ fingerprint of the headspace of Iberian dry-cured *Montanera* and *Campo* hams lean.

2.5. Statistical analysis

A one-way ANOVA was carried out in order to find the masses significantly different between dry-cured *Campo* and *Montanera* hams ($p < 0.05$). Principal component (PCA) and discriminant analysis were also performed. A simplified version of the methods used in [Fabris et al. \(2010\)](#) was used for discriminant analysis on PTR-ToF-MS data, using Penalized Discriminant Analysis (PDA) ([Granitto et al., 2007](#)) as classifier. To evaluate the results of the classification method a leave-

group-out (LGO) method was used: the process of leaving a group out as test set and using the rest of the data set to fit the models was iterated. The regularization constant of PDA was selected each time at this step by internal cross validation using only the training data sets. The PDA model was then used to individually classify the samples of the independent test batch. Each individual group in this LGO procedure consisted of the 3 replicates of the same ham. The discrimination results were analysed with confusion matrices, in which rows correspond to the true classes and columns to the predicted ones ([Witten & Frank, 2005](#)).

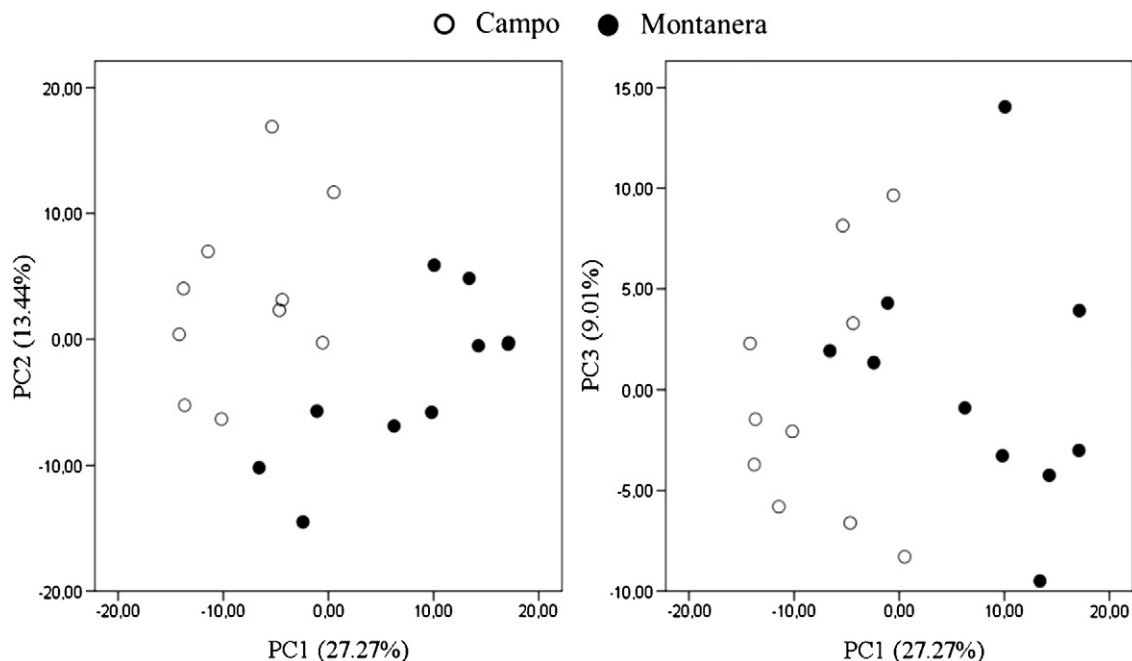


Fig. 2. Score plots obtained by the PCA analysis (PC1 vs PC2 and PC1 vs PC3 respectively) of the PTR-ToF-MS H_3O^+ fingerprint of the headspace of Iberian dry-cured *Montanera* and *Campo* hams subcutaneous fat (SCF).

Table 2

Confusion matrices for the penalized discriminant analysis with the lean and subcutaneous fat analysis data obtained by using H_3O^+ as reagent ion. Numbers in bold indicate correct classified replicates.

Lean			Subcutaneous fat		
	Campo	Montanera	%		
Campo	27	3	95	Campo	28
Montanera	0	30		Montanera	29

%. Percentage of correct classified replicates in both dataset.

3. Results and discussion

3.1. PTR-ToF-MS H_3O^+ headspace analysis of lean and SCF samples

The rapid analysis (30 s) of the headspace of the *Campo* and *Montanera* dry cured hams by PTR-ToF-MS using H_3O^+ as reagent ion resulted in more than 600 mass peaks. Multivariate analysis of PTR-ToF-MS spectra is a useful tool to exploit a rapid VOC fingerprint to classify dry-cured hams according to different criteria. Principal components analysis (PCA) was performed as an exploratory non-supervised data analysis, and the results are displayed in Figs. 1 and 2. In this analysis the average spectra of the three replicates of each sample were used. In Fig. 1, which corresponds to the analysis of the lean data, the three first principal components (PC) explain 53% of the total variance. Although the PC1–PC2 plot does not show a good discrimination of the samples, the PC1–PC3 plot indicates the possibility of a very good discrimination according to the rearing system of the pigs. In Fig. 2, which corresponds to the analysis of the SCF data, the two first PC explain 41% of the total variance and allow a better discrimination of the ham samples according to the fattening diet of the pigs.

A PDA (Penalized Discriminant Analysis) classifier was used to better assess the possibility of distinguishing ham samples according to the rearing system. Table 2 displays the confusion matrices of the PDA for the H_3O^+ lean and SCF analysis: a very good classification was achieved on both datasets, with only 3 replicates being misclassified in each case. In both cases two of the three errors correspond to the same sample. These results confirm the good separability suggested by the PCA analysis.

Tables 3 and 4 show the concentrations of different tentatively identified mass peaks from the spectral data acquired using H_3O^+ as primary ion from the analysis of the lean and the subcutaneous fat (SCF) samples respectively. Among the more than 600 peaks identified, only those corresponding to concentrations higher than 1 ppbv and significantly different ($p < 0.05$) between ham types are listed. For lean (Table 3) and fat headspace (Table 4), aldehydes and ketones with less than 9 carbon atoms were more concentrated in *Montanera* dry-cured Iberian hams than in *Campo* ones. The opposite holds for aldehydes and ketones with 9 and 10 carbon atoms.

The peak observed at $m/z = 87.081$, corresponding to protonated linear and branched aldehydes and ketones with 5 carbon atoms ($\text{C}_5\text{H}_{11}\text{O}^+$), showed a higher concentration than the other peaks corresponding to aldehydes and ketones, both in lean and SCF, probably due to the presence of 3-methylbutanal, and in less proportion to 2-methylbutanal, pentanal and 2-pentanone. In fact, 3-methylbutanal is the most abundant branched aldehyde found in dry-cured ham, followed by 2-methylbutanal (Andrade, Córdoba, Sánchez, Casado, & Rodríguez, 2009; Andrés, Cava, & Ruiz, 2002; Carrapiso, Jurado, & García, 2003; Dirinck, Van Opstaele, & Vandendriessche, 1992; García et al., 1991; Jurado, García, Timón, & Carrapiso, 2007; Martín, Córdoba, Aranda, Córdoba, & Asensio, 2006; Ruiz, García, Muriel, Andrés, & Ventanas, 2002; Sabio, Vidal-Aragón, Bernalte, & Gata, 1998). 2 and 3-methylbutanal are originates mainly from Strecker reactions of the aminoacids and Maillard reactions (Berdagué, Denoyer, Le Quere, & Semon, 1991; Flores, Grimm, Toldrá, & Spanier, 1997; García et al., 1991; Ruiz et al., 2002; Toldrá, Aristoy, & Flores, 2000; Ventanas et al., 1992). Nevertheless, in the SCF the major contributor to this peak is probably pentanal, followed by 3 and 2-methylbutanal (Timón, Ventanas, Carrapiso, Jurado, & García, 2001). For these isobaric compounds the concentration in dry-cured *Montanera* hams lean was double than in *Campo* hams, while in the SCF the concentration was six times higher in *Montanera* hams. Saturated aldehydes and ketones with 6 carbon atoms (measured at $m/z = 101.098$) were also two-fold higher in *Montanera* than in *Campo* ham lean, and four times in the SCF of *Montanera* than in the SCF of *Campo* hams. In this case the major contributor to the abundance is probably hexanal, the most concentrated aldehyde in dry-cured ham lean (Andrade et al., 2009; Andrés et al., 2002; Carrapiso, Jurado, et al., 2003; Dirinck et al., 1992; García et al.,

Table 3

Peaks from the PTR-ToF-MS H_3O^+ lean analysis significantly different between both ham batches ($p < 0.05$) and with intensity higher than 1 ppbv (mean \pm standard error).

Measured mass (m/z)	Sum Formula	Theoretical mass (m/z)	Tentative identification	Campo (ppbv)	Montanera (ppbv)
33.033	CH_5O^+	33.0335	Methanol	765 ± 25	692 ± 17
55.055	C_4H_7^+	55.0542	C4 aldehydes fragment	221 ± 19	361 ± 43
83.087	$\text{C}_6\text{H}_{11}^+$	83.0855	C6 aldehydes fragment	105 ± 13	211 ± 34
87.081	$\text{C}_5\text{H}_{11}\text{O}^+$	87.0804	C5 aldehydes and ketones	499 ± 62	1057 ± 108
91.058	$\text{C}_4\text{H}_{11}\text{S}^+$	91.0576	Butanethiol/1-(methylthio)-propane	5.3 ± 0.5	3.9 ± 0.3
95.059	$\text{C}_5\text{H}_7\text{N}_2^+$	95.0604	Methylpyrazine	5.3 ± 0.3	4.3 ± 0.2
97.067	$\text{C}_6\text{H}_9\text{O}^+$	97.0648	Ethylfuran	4.0 ± 0.4	5.2 ± 0.3
98.101	$\text{C}_6\text{H}_{12}\text{N}^+$	98.0964	Hexanenitrile	4.4 ± 0.3	5.4 ± 0.3
99.082	$\text{C}_6\text{H}_{11}\text{O}^+$	99.0804	C6 unsaturated aldehydes and ketones	3.1 ± 0.1	3.9 ± 0.3
99.118	$\text{C}_7\text{H}_{15}^+$	99.1168	1-Heptene/Methyl-cyclohexane	2.7 ± 0.2	2.1 ± 0.1
101.098	$\text{C}_6\text{H}_{13}\text{O}^+$	101.0961	C6 aldehydes and ketones	44 ± 5	92 ± 10
105.072	$\text{C}_5\text{H}_{13}\text{S}^+$	105.0732	Pentanethiol	5.7 ± 0.6	3.3 ± 0.3
111.083	$\text{C}_7\text{H}_{11}\text{O}^+$	111.0804	2,4-Heptadienal	2.4 ± 0.1	2.8 ± 0.1
119.089	$\text{C}_6\text{H}_{15}\text{S}^+$	119.0889	1-(Methylthio)-pentane/Hexanthiol	1.1 ± 0.1	0.8 ± 0.1
125.099	$\text{C}_8\text{H}_{13}\text{O}^+$	125.0961	Octadienal/octadienone	2.2 ± 0.1	2.6 ± 0.2
133.106	$\text{C}_7\text{H}_{17}\text{S}^+$	133.1045	Heptanethiol	2.3 ± 0.3	1.4 ± 0.2
139.115	$\text{C}_8\text{H}_{15}\text{O}^+$	139.1117	Nonadienal/pentyl-furan	3.2 ± 0.5	6.3 ± 0.8
141.130	$\text{C}_9\text{H}_{17}\text{O}^+$	141.1274	C9 unsaturated aldehydes and ketones	2.1 ± 0.2	1.3 ± 0.1
143.115	$\text{C}_8\text{H}_{15}\text{O}^+$	107.1066	2,3-octanedione/2,4-octadienal	1.9 ± 0.1	1.4 ± 0.1
143.146	$\text{C}_9\text{H}_{19}\text{O}^+$	143.143	C9 aldehydes and ketones	21 ± 1	11.7 ± 0.9
149.138	$\text{C}_{11}\text{H}_{17}^+$	149.1325	Pentylbenzene	1.0 ± 0.1	0.66 ± 0.05
157.163	$\text{C}_{10}\text{H}_{21}\text{O}^+$	157.1587	C10 aldehydes and ketones	1.7 ± 0.1	1.3 ± 0.1
177.167	$\text{C}_{13}\text{H}_{21}^+$	177.1678	Heptylbenzene	1.2 ± 0.1	0.8 ± 0.1
201.188	$\text{C}_{12}\text{H}_{15}\text{O}_2^+$	201.184	Decanoic acid	1.2 ± 0.1	0.60 ± 0.05

Table 4

Peaks from the PTR-ToF-MS H_3O^+ subcutaneous fat (SCF) analysis significantly different between both ham batches ($p < 0.05$) and with intensity higher than 1 ppbv (mean \pm standard).

Measured mass (m/z)	Sum Formula	Theoretical mass (m/z)	Tentative identification	Campo (ppbv)	Montanera (ppbv)
33.034	CH_5O^+	33.0335	Methanol	310 ± 18	218 ± 14
41.039	C_3H_5^+	41.0386	Alkyl fragment	422 ± 27	701 ± 38
43.018	$\text{C}_2\text{H}_3\text{O}^+$	43.0178	Alkyl fragment	130 ± 12	169 ± 8
43.054	C_3H_7^+	43.0542	Alkyl fragment	383 ± 25	590 ± 43
55.055	C_4H_7^+	55.0542	C4 aldehydes fragment	143 ± 20	373 ± 41
57.034	$\text{C}_3\text{H}_5\text{O}^+$	57.0335	C3 aldehydes and ketones	66 ± 5	99 ± 4
69.036	$\text{C}_4\text{H}_5\text{O}^+$	69.0335	Furan	7.8 ± 0.6	10.7 ± 0.7
69.070	C_5H_5^+	69.0698	C5 aldehydes fragment	113 ± 9	260 ± 23
71.086	$\text{C}_5\text{H}_{11}^+$	71.0855	Alkyl fragment	36 ± 4	62 ± 7
73.065	$\text{C}_4\text{H}_9\text{O}^+$	73.0648	C4 aldehydes and ketones	180 ± 13	399 ± 36
79.055	C_6H_7^+	79.0542	Benzene	5.1 ± 0.4	6.5 ± 0.4
80.053	$\text{C}_5\text{H}_6\text{N}^+$	80.0495	Pyridine	1.00 ± 0.05	1.23 ± 0.04
81.071	C_6H_5^+	81.0699	Alkyl fragment	5.8 ± 0.3	7.5 ± 0.4
83.051	$\text{C}_5\text{H}_7\text{O}^+$	83.0491	Methyl-furan	9.4 ± 0.5	12.7 ± 0.6
83.086	$\text{C}_6\text{H}_{11}^+$	83.0855	C6 aldehydes fragment	72 ± 12	186 ± 23
85.066	$\text{C}_5\text{H}_9\text{O}^+$	85.0648	C5 unsaturated aldehydes and ketones	5.6 ± 0.7	8.9 ± 0.7
87.080	$\text{C}_5\text{H}_{11}\text{O}^+$	87.0804	C5 aldehydes and ketones	225 ± 32	1540 ± 285
97.065	$\text{C}_6\text{H}_6\text{O}^+$	96.0648	Ethylfuran	22 ± 2	40 ± 3
98.100	$\text{C}_6\text{H}_{12}\text{N}^+$	98.0964	Hexanenitrile	4.0 ± 0.3	7.1 ± 0.7
99.081	$\text{C}_6\text{H}_{11}\text{O}^+$	99.0804	C6 unsaturated aldehydes and ketones	2.0 ± 0.1	3.7 ± 0.3
101.060	$\text{C}_5\text{H}_9\text{O}_2^+$	101.0597	2,3-pentanedione	9 ± 1	15 ± 2
101.096	$\text{C}_6\text{H}_{13}\text{O}^+$	101.0961	C6 aldehydes and ketones	30 ± 4	130 ± 19
107.050	$\text{C}_7\text{H}_7\text{O}^+$	107.0499	Benzaldehyde	1.5 ± 0.2	2.2 ± 0.1
111.080	$\text{C}_7\text{H}_{11}\text{O}^+$	111.0804	2,4-Heptadienal	3.0 ± 0.3	4.7 ± 0.2
111.117	$\text{C}_8\text{H}_{15}^+$	111.1168	C8 aldehydes fragment	4.1 ± 0.6	6.5 ± 0.9
113.060	$\text{C}_6\text{H}_9\text{O}_2^+$	113.0597	2,5-Dimethyl-3(2 H)-Furanone	0.89 ± 0.03	1.10 ± 0.04
113.096	$\text{C}_7\text{H}_{13}\text{O}^+$	113.0961	C7 unsaturated aldehydes and ketones	2.9 ± 0.2	3.9 ± 0.1
115.074	$\text{C}_6\text{H}_{11}\text{O}_2^+$	115.0754	Hexanedione/5-ethylidihydro-2(3 H)-furanone	2.3 ± 0.1	3.4 ± 0.2
115.112	$\text{C}_7\text{H}_{15}\text{O}^+$	115.1117	C7 aldehydes and ketones	54 ± 6	134 ± 17
123.082	$\text{C}_8\text{H}_{11}\text{O}^+$	123.0804	Dimethyl-phenol	1.3 ± 0.1	1.8 ± 0.2
125.096	$\text{C}_8\text{H}_{13}\text{O}^+$	125.0961	Octadienal/octadienone	4.7 ± 0.7	7.0 ± 0.8
127.112	$\text{C}_8\text{H}_{15}\text{O}^+$	127.1117	1-Octen-3-one	1.4 ± 0.1	2.3 ± 0.2
129.127	$\text{C}_8\text{H}_{17}\text{O}^+$	129.1274	C8 aldehydes and ketones	18 ± 2	27 ± 3
136.022	$\text{C}_7\text{H}_6\text{NS}^+$	136.0215	Benzothiazole	1.75 ± 0.04	1.59 ± 0.04
143.142	$\text{C}_9\text{H}_{19}\text{O}^+$	143.143	C9 aldehydes and ketones	7 ± 1	4.0 ± 0.4

1991; Jurado et al., 2007; Martín et al., 2006; Ruiz et al., 2002; Sabio et al., 1998) and fat (Timón et al., 2001). An hexanal fragment caused by the loss of a water molecule was also tentatively identified at $m/z = 3.087$, and its concentration in the lean was double in the dry-cured *Montanera* hams than in *Campo* hams, while in the SCF its concentration was almost triple in *Montanera* hams. The origin of pentanal and hexanal is fatty acid oxidation, mainly the oxidation of linoleic acid (Larick, Turner, Schoenherr, Coffey, & Pilkington, 1992; Ruiz et al., 2002), which seems to be more concentrated in dry-cured hams from pigs fattened on acorn and grass than in pigs fattened on a high-oleic concentrated feed (Pérez-Palacios et al., 2010). Also saturated aldehydes and ketones with 7 and 8 carbon atoms (measured at $m/z = 115.112$ and 129.127, Table 4) were more concentrated in the SCF of *Montanera* hams than in *Campo* ones. In both cases the linear aldehydes (heptanal and octanal respectively) are probably the major contributors to the peak concentration, but in the case of

the peak at $m/z = 115.112$ (Table 4), 2-heptanone, which was also found at a high concentration in the SCF of dry-cured Iberian ham (Timón et al., 2001), can play a role. These compounds are produced by the oxidation of unsaturated fatty acids, like oleic acid (Ruiz et al., 2002b), the most concentrated fatty acid in pork, and also more concentrated in hams from pigs fattened on acorn and grass than in hams from pigs fattened on concentrated feed (Carrapiso, Jurado, Martín, & García, 2007; Carrapiso, Jurado, Timón, & García, 2002; Ventanas et al., 2007), even when the concentrated feed is enriched with oleic acid (Pérez-Palacios et al., 2010). Peaks corresponding to unsaturated aldehydes and ketones with less than 9 carbon atoms were also found in higher concentration in *Montanera* than in *Campo* hams, both in lean (Table 3) and SCF (Table 4), probably due to the higher concentration in these hams of the unsaturated fatty acids that are their precursors (Pérez-Palacios et al., 2010), but the concentration in both kinds of ham was much lower than that of

Table 5

Odour-impact compounds tentatively identified from the PTR-ToF-MS H_3O^+ analysis of the dry-cured Iberian ham lean headspace and their mean concentration (ppbv) in *Campo* and *Montanera* hams.

Measured mass (m/z)	Sum formula	Theoretical mass (m/z)	Tentative identification	Mean concentration (ppbv)
49.011	CH_5S^+	49.0106	Methanethiol	44.1
75.045	$\text{C}_3\text{H}_7\text{O}_2^+$	75.0440	Propanoic acid	21.3
87.047	$\text{C}_4\text{H}_7\text{O}_2^+$	87.0441	2,3-butanedione	26.1
89.061	$\text{C}_4\text{H}_9\text{O}_2^+$	89.0597	Butanoic acid	45.5
103.078	$\text{C}_5\text{H}_{11}\text{O}_2^+$	103.0754	3-methylbutanoic acid	5.6
105.039	$\text{C}_4\text{H}_9\text{OS}^+$	105.0369	3-(methylthio)-propanal	1.0
109.077	$\text{C}_6\text{H}_9\text{N}_2^+$	109.0760	2,6-dimethylpyrazine	8.9
121.067	$\text{C}_6\text{H}_9\text{O}^+$	121.0648	Benzeneacetaldehyde	20.6
129.057	$\text{C}_6\text{H}_9\text{O}_3^+$	129.0546	2,5-dimethyl-4-hydroxy-—3(2 H)-furanone	0.6
131.110	$\text{C}_7\text{H}_{15}\text{O}_2^+$	131.1067	2-methylbutanoic acid, ethyl ester + 3-methylbutanoic acid, ethyl ester	1.7

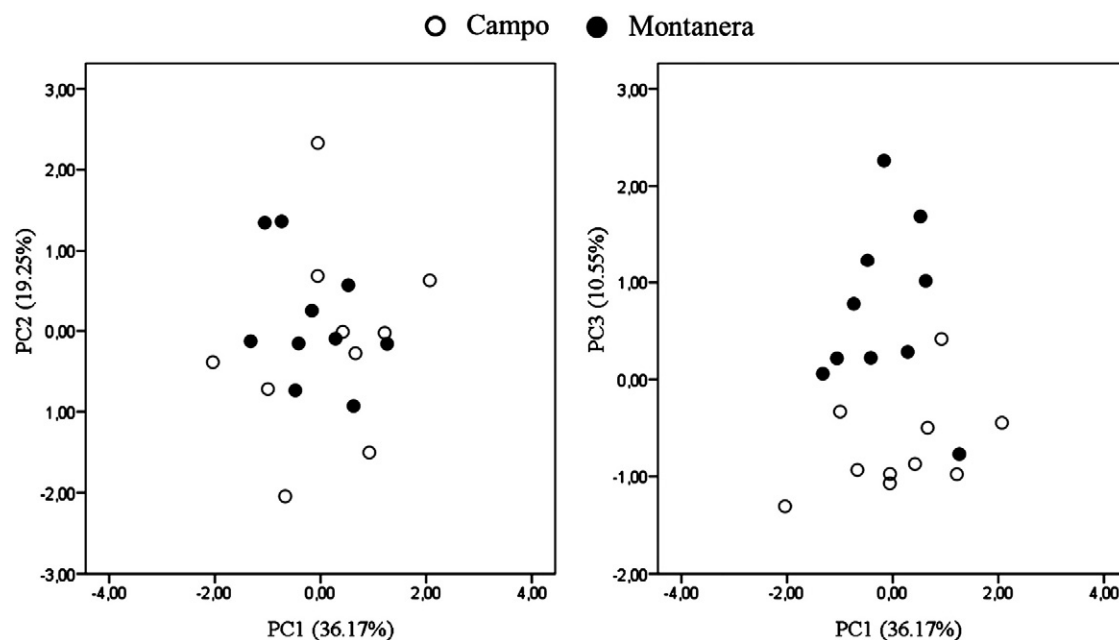


Fig. 3. Score plots obtained by the PCA analysis (PC1 vs PC2 and PC1 vs PC3 respectively) of the PTR-ToF-MS NO^+ fingerprint of the headspace of Iberian dry-cured *Montanera* and *Campo* hams lean.

the saturated aldehydes and ketones, which agree with previous results in dry-cured Iberian ham (Andrés et al., 2002; Martín et al., 2006; Ruiz et al., 2002).

Aldehydes and ketones with 3 and 4 carbon atoms were also tentatively identified at $m/z = 59.051$ and 73.065 respectively. The peak at $m/z = 59.051$, with propanone as probable major contributor (Ruiz et al., 2002), was saturated and the concentration was estimated on the basis of its isotopologue at $m/z = 60.053$. It showed no significant differences between ham batches. In the case of the aldehydes and ketones with 4 carbon atoms ($m/z = 73.065$) the major contributors to the peak signal were probably butanal and methylpropanal, found at higher concentrations than their corresponding ketones in previous studies (Andrés et al., 2002; Carrapiso, Jurado, et al., 2003; Ruiz et al., 2002). This peak was significantly different between both ham batches in the fat headspace (Table 4) but

not in the lean headspace (Table 3), while the concentration of the butanal fragment, tentatively identified at $m/z = 55.055$, was double in both lean and SCF of dry-cured *Montanera* hams than in *Campo* hams (Tables 3 and 4).

On the contrary, peaks corresponding to aldehydes and ketones with 9 carbon atoms in lean and SCF (measured $m/z = 143.146$ and 143.142 respectively) and to aldehydes and ketones with 10 carbon atoms in lean (measured $m/z = 157.163$) were more concentrated in dry-cured *Campo* hams.

According to Sánchez del Pulgar et al. (2011), many peaks corresponding to alkyl fragments were found in the head-space of dry-cured Iberian ham, some of them being significantly different between *Montanera* and *Campo* hams. Peaks at $m/z = 41.038$, 43.054 and 71.085 were more concentrated in *Montanera* hams, and are related to alkyl fragments probably from the split off of water from

Table 6

Peaks from the PTR-ToF-MS NO^+ lean analysis significantly different between both ham batches ($p < 0.05$) and with intensity higher than 1 normalized count (mean \pm standard error).

Measured mass (m/z)	Sum Formula	Theoretical mass (m/z)	Tentative identification	Campo (nc) ^a	Montanera (nc)
33.033	CH_4OH^+	33.0335	Methanol	1331 ± 50	1142 ± 52
57.033	$\text{C}_3\text{H}_5\text{O}^+$	57.0335	C3 aldehydes	98 ± 10	128 ± 9
58.042	$\text{C}_3\text{H}_6\text{O}^+$	58.0413	C3 ketones	241 ± 19	385 ± 30
72.057	$\text{C}_4\text{H}_8\text{O}^+$	72.057	C4 aldehydes	67 ± 5	81 ± 4
76.004	CH_2NO_3^+	76.0029	Cluster formic acid	4.1 ± 0.2	4.6 ± 0.2
80.062	C_6H_8^+	80.0621	Cyclohexadiene	3.5 ± 0.2	4.6 ± 0.3
82.076	$\text{C}_6\text{H}_{10}^+$	82.0777	Cyclohexene/methyl-cyclopentene	8.2 ± 0.8	12.6 ± 1.2
86.072	$\text{C}_5\text{H}_{10}\text{O}^+$	86.0726	C5 ketones	89 ± 11	197 ± 23
87.080	$\text{C}_5\text{H}_{10}\text{OH}^+$	87.0804	C5 aldehydes and ketones	1380 ± 176	2440 ± 248
99.080	$\text{C}_6\text{H}_{11}\text{O}^+$	99.0804	C6 aldehydes	68 ± 8	116 ± 13
99.114	$\text{C}_6\text{H}_{13}\text{N}^+$	99.1043	Methyl-piperidine	4.9 ± 0.3	6.2 ± 0.3
100.082	$\text{C}_6\text{H}_{12}\text{O}^+$	100.0883	C6 ketones	15 ± 2	25 ± 2
101.096	$\text{C}_6\text{H}_{12}\text{OH}^+$	101.0961	C6 aldehydes and ketones	82 ± 12	145 ± 15
102.056	$\text{C}_4\text{H}_8\text{NO}_2^+$	102.055	cluster C4 ketones	22 ± 1	32 ± 2
112.084	$\text{C}_7\text{H}_{12}\text{O}^+$	112.0883	C7 unsaturated aldehydes	8.2 ± 0.8	11.4 ± 0.9
114.060	$\text{C}_5\text{H}_8\text{NO}_2^+$	114.055	Cluster C5 unsaturated ketones	4.1 ± 0.3	5.6 ± 0.4
116.070	$\text{C}_5\text{H}_{10}\text{NO}_2^+$	116.0706	cluster C5 ketones	56 ± 9	171 ± 23
130.085	$\text{C}_6\text{H}_{12}\text{NO}_2^+$	130.0863	cluster C6 ketone	8.3 ± 1.1	16.2 ± 1.8
144.063	$\text{C}_6\text{H}_{10}\text{NO}_3^+$	144.0655	Cluster hexanedione/ethylidihydro-2(3H)-furanone	3.7 ± 0.3	5.9 ± 0.6

^a Normalized counts.

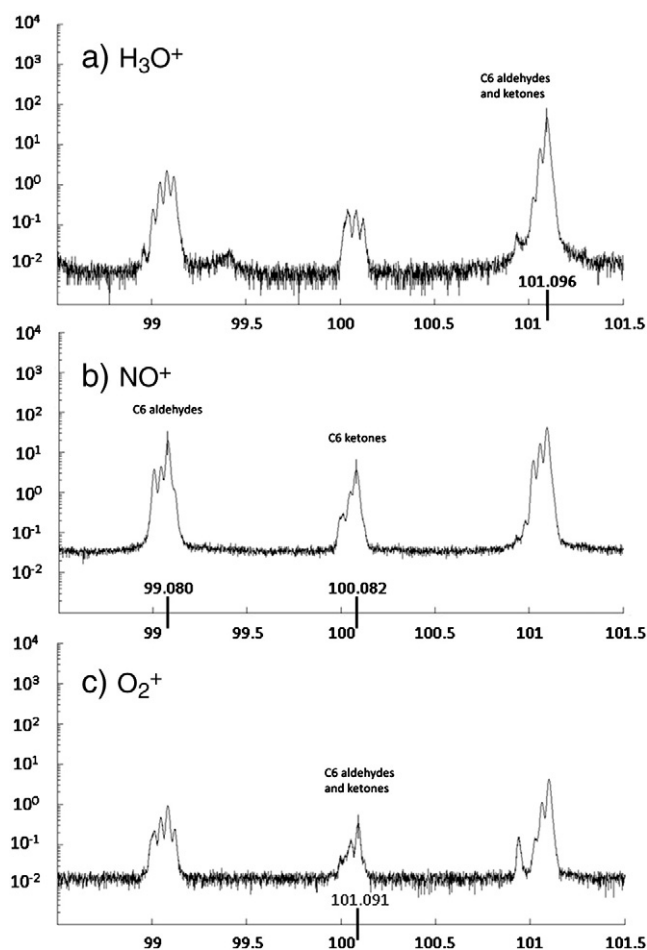


Fig. 4. Example of peaks related to aldehydes and ketones measured by using different reagent ions. a: H_3O^+ . b: NO^+ . c: O_2^+ .

linear and branched saturated alcohols (Aprea, Biasioli, Mark, & Gasperi, 2007).

The peak at $m/z = 43.018$ in the SCF samples was also more concentrated in *Montanera* hams SCF (Table 4), and corresponds to an alkyl fragment from the split off of a water molecule from acetic acid (Aprea, Biasioli, Gasperi, et al., 2007).

Dry-cured Iberian *Campo* ham lean showed higher concentration for the peaks at m/z 91.058, 105.072, 119.089 and 133.106 (Table 3), corresponding to the sum formula of the sulphur-containing compounds $\text{C}_4\text{H}_{11}\text{S}^+$, $\text{C}_5\text{H}_{13}\text{S}^+$, $\text{C}_6\text{H}_{15}\text{S}^+$, and $\text{C}_7\text{H}_{17}\text{S}^+$ (tentatively identified as butanethiol/1-(methylthio)-propane, pentanethiol/1-(methylthio)-butane, hexanethiol/1-(methylthio)-pentane and heptanethiol/1-(methylthio)-hexane respectively), while the SCF of *Campo* hams showed higher amounts of benzothiazol ($m/z = 136.022$, $\text{C}_7\text{H}_6\text{NS}^+$) (Table 4). Sulphur-containing compounds arise from Strecker reactions of amino-acids and Maillard reactions (Ruiz et al., 2002) and are potent odorants, major contributors to meat odour (Carrapiso et al., 2002; Ruiz et al., 2002; Sánchez del Pulgar, García, Reina, & Carrapiso, 2012). Moreover, the peak detected at $m/z = 95.059$ (tentatively identified as methylpyrazine – $\text{C}_5\text{H}_7\text{N}_2^+$ –) in lean (Table 3), with the same origin as sulphur compounds and the rest of the nitrogen compounds (Ruiz et al., 2002), was more concentrated in dry-cured *Campo* hams than in *Montanera* ones. Pyrazines are also potent odorants with nutty and roast notes (Ruiz et al., 2002; Sánchez del Pulgar et al., 2012). However, *Montanera* hams SCF showed higher signals of the peak at $m/z = 80.053$, corresponding to the formula $\text{C}_5\text{H}_6\text{N}^+$ and tentatively identified as pyridine.

Hexanenitrile, already identified in dry-cured ham by PTR-ToF-MS (Sánchez del Pulgar et al., 2011), was more concentrated in *Montanera*

hams than in *Campo* ones, both in lean and SCF (Tables 3 and 4). Nitrile compounds have been detected in dry-cured ham (Ruiz et al., 1998), and their most probable formation occurs at the expense of the corresponding aldehydes during lipid oxidation involving nitrite (Mottram, Croft, & Patterson, 1984), so the higher concentration of hexanal found in *Montanera* hams could explain also the higher concentration of hexanenitrile.

The PTR-ToF-MS analysis of the dry-cured Iberian ham headspace allowed the detection of more compounds than those included in Tables 3 and 4, some of them identified as major odorants of dry-cured Iberian ham in previous studies (Carrapiso & García, 2004; Carrapiso et al., 2002; Sánchez del Pulgar et al., 2012), but at concentrations not significantly different between batches. These compounds and their mean concentration in dry-cured *Campo* and *Montanera* hams are listed in Table 5.

3.2. Switching reagent ion system

PCA and discriminant analysis (PDA) were also performed on data obtained with NO^+ and O_2^+ for the lean headspace. For NO^+ (Fig. 3), the first three PCs explain 66% of the total variance. Although the PC1–PC2 plot shows a poor discrimination of the samples, the PC1–PC3 plot suggests a good discrimination of the dry-cured Iberian ham samples according to the rearing system of the pigs. On the contrary, spectra obtained using O_2^+ as precursor ion do not suggest a simple discrimination of the ham batches.

In the case of the NO^+ dataset, PDA classifies correctly 83.3% of the replicates. It is worth mentioning that 5 out of the ten misclassified replicates correspond to five different samples, and the other five correspond to only two samples. In the case of the O_2^+ dataset, even if the PCA does not suggest an easy discrimination of the batches, PDA allows the discrimination of 78.3% of the replicates.

As shown in Table 6, the analysis of the samples with NO^+ allowed the detection of several peaks significantly different between both hams and corresponding to aldehydes and ketones. In agreement with the results from the ham lean analysis using H_3O^+ (Table 3), aldehydes and ketones were more concentrated in dry-cured *Montanera* hams than in dry-cured *Campo* hams. As mentioned in the Introduction, NO^+ allows the detection in different peaks of the isobaric aldehydes and ketones due to the ionization reaction that occurs in both cases (Fig. 4). Except for aldehydes and ketones with 3 carbon atoms, each aldehyde was more concentrated than its corresponding ketone. Matching the absence of differences in the concentration of the 5 carbon atom aldehyde fragment in the lean analysis with H_3O^+ , aldehydes with 5 carbon atoms showed no differences between batches in the NO^+ analysis. No difference was found in the concentration of 4 carbon atom aldehydes, although the butanal fragment was more concentrated in *Montanera* hams than in *Campo* hams in the lean analysis with H_3O^+ . In addition, ketones with 4 and 5 carbon atoms were more concentrated in dry-cured *Montanera* hams than in *Campo* ones. As a first approach to the VOC analysis by PTR-ToF-MS using NO^+ as reagent ion in a complex matrix, the data indicate that also H_3O^+ has intervened in the ionization of the compounds present in the sample headspace. In fact, Table 6 shows significant differences in the concentration of the peaks corresponding to protonated aldehydes and ketones with 5 and 6 carbon atoms, this was also found in the peaks corresponding to other protonated aldehydes and ketones (data not included). Therefore, the PTR-ToF-MS analysis with H_3O^+ allowed the detection of the isobaric aldehydes and ketones in the same peak, and, due to the high energy into the drift tube, also aldehyde fragments caused by the loss of a water molecule (Jordan et al., 2009). Nevertheless, the use of NO^+ as reagent ion allowed the detection of aldehydes and ketones in separated peaks, permitting confirmation of the results of the H_3O^+ analysis, and even indicating which isobaric compound was responsible for the differences between batches. It was

checked that the sum of the intensity of the peaks corresponding to aldehydes and ketones with 6 carbon atoms in the H_3O^+ analysis (ppbv of the hexanal fragment peak and the peak for C6 saturated aldehydes and ketones) and in the NO^+ analysis (normalized counts corresponding to the aldehyde peak, the ketone, the cluster ketone- NO^+ and the peak of protonated aldehydes and ketones). In both analyses the intensity ratio of the C6 aldehydes and ketones between *Campo* and *Montanera* hams is almost the same.

The use of O_2^+ as reagent ion in the PTR-ToF-MS analysis of the dry-cured Iberian ham lean showed no peaks significantly different between ham batches, with the exception of the peak at $m/z = 82.079$, tentatively identified as alkenes with 6 carbon atoms, and the peak at $m/z = 86.075$, tentatively identified as saturated aldehydes and ketones with 5 carbon atoms, which were more concentrated in dry-cured *Montanera* than in *Campo* hams.

The data demonstrate the feasibility and usefulness of the SRI system in food samples. Nevertheless, it was not possible to obtain for NO^+ and O_2^+ the same purity of the primary ion beam achieved with H_3O^+ (Table 1). The presence of the signals produced by H_3O^+ proton transfer during NO^+ and O_2^+ analysis also makes data analysis more difficult and less conclusive.

4. Conclusions

Pig rearing system strongly affects the volatile compound profile of dry-cured Iberian ham.

The rapid PTR-ToF-MS analysis of the headspace of dry-cured Iberian ham, both lean and subcutaneous fat, allows the discrimination of the hams according to the rearing system: *Campo* and *Montanera*.

Many peaks are significantly different between *Montanera* and *Campo* Iberian hams. In particular *Montanera* samples show higher amounts of aldehydes and ketones than *Campo* probably due to the higher amounts of their precursor fatty acids, while dry-cured *Campo* hams have higher concentrations of peaks corresponding to sulphur compounds.

In this study the switching reagent ion system in PTR-MS instruments was applied for the first time to food products providing the possibility of separating the contribution of isobaric compounds (aldehydes and ketones), however the same purity of the precursor ion beam achieved with H_3O^+ was not obtained and this limits classification efficiency and data interpretation.

The rapid and high sensitive PTR-ToF-MS technique coupled to appropriate chemometric analysis provides both classification models and analytical information. It could be useful for the dry-cured Iberian ham industry. In particular the rapid characterisation of subcutaneous fat seems suited for the implementation of a non-destructive on-line quality control monitoring.

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