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4	Determination of Polyphenolic Profiles by Liquid Chromatography-Electrospray-
5	Tandem Mass Spectrometry for the Authentication of Fruit Extracts
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#### 27 Abstract

Liquid chromatography-electrospray-tandem mass spectrometry (LC-ESI-28 MS/MS) was applied to the analysis and authentication of fruit-based products and fruit-29 based pharmaceutical preparations. A Kinetex C<sub>18</sub> reversed-phase column 30 under gradient elution with 0.1 % formic acid aqueous solution and methanol mobile phases 31 was used for the simultaneous determination of 26 polyphenols, allowing an acceptable 32 separation in less than 22 min. Instrumental quality parameters such as limits of 33 detection (LOD, values between 12-14 µg/L for 19 of the 26 analyzed polyphenols), 34 linearity ( $r^2 > 0.991$ ), run-to-run and day-to-day precisions (RSD values lower than 9.9 35 and 13.5 %, respectively), and accuracy (relative errors lower than 8 %) were 36 established. A simple extraction method, consisting of a sample sonication with 37 acetone:water:hydrochloric acid (70:29.9:0.1 v/v/v) and centrifugation, was proposed. 38 39 Two calibration procedures, external calibration using standards prepared in water and 40 standard addition, were evaluated for polyphenol quantification in several grape and 41 cranberry fruits and processed fruit products. For a 95 % confidence level, no statistical differences were observed between the two calibration methods (p values between 0.06 42 43 and 0.95), denoting that external calibration was suitable enough for the quantitative analysis of polyphenols in fruit-based products. The proposed LC-ESI-MS/MS method 44 45 was then applied to the analysis of polyphenols in 23 grape-based and cranberry-based natural products and pharmaceutical preparations. Polyphenolic concentration data was 46 47 then analyzed by principal component analysis (PCA) to extract information of the most significant profile data contributing to authentication of natural extracts according to 48 49 their fruit of origin.

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53 KEYWORDS: Polyphenols; Natural products; Food characterization; Food Analysis;
54 Liquid chromatography; Mass spectrometry

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#### 60 **1. Introduction**

The importance of diet on human health and well-being has been widely 61 recognized all over the world. For instance, USA recommends that people consume at 62 least 2.5 cups of vegetables and 2 cups of fruits daily [1], which is based on a general 63 64 diet of 2000 kcal per day. In Europe, instead, the traditional Mediterranean diet has formed the basis for food consumption during the past century, originally settled on 65 Mediterranean agronomical, pastoral, and rural archetypes. The regular consumption of 66 fruits and vegetables, rich in antioxidants and bioactive compounds, has been shown to 67 68 exert an important role in the prevention of many diseases, such as skin pathologies, various types of cancer, cardiovascular disorders, and other age-related degenerative 69 70 pathologies, besides the general health benefits they provide [2-6].

71 Polyphenols usually are related with characteristic metabolic patterns present in all vegetal tissues, as well as in flowers and fruits. Several thousands of plant 72 73 polyphenols are known, including a wide variety of molecules that contain at least one 74 aromatic ring with one or more hydroxyl groups in addition to other constituents. They can be divided in several classes, i.e. phenolic acids (hydroxybenzoic acids and 75 hydroxycinnamic acids), flavonoids (flavonols, flavones, flavanols, flavanones, 76 isoflavones, proanthocyanidins (PACs)), stilbenes, and lignans [7]. Phenolic profile is 77 an important indicator of fruit quality because of their contribution to the taste, color 78 and nutritional properties [8]. In addition, these compounds are considered one of the 79 80 most relevant antioxidants of human diet [9], so over the past ten years food researchers and manufacturers have become increasingly interested in this family of compounds. 81

82 Berries are an excellent source of polyphenols, especially anthocyanins. The consumption of berry fruits associated with their contribution to improved human health 83 is an issue of considerable interest [10]. Cranberry (Vaccinium macrocarpon) and its 84 85 derived products, including juices and nutraceuticals, have shown some beneficial health effects associated to their polyphenolic content [11]. However, the best known 86 bioactivity of cranberry polyphenols deals with their capacity to inhibit the adhesion of 87 pathogenic bacteria to uroepithelial cells of the urinary tract, thus contributing to the 88 prevention of urinary tract infections [12,13]. The most common polyphenols found in 89 cranberries comprise phenolic and benzoic acids, and flavonoids such as anthocyanins, 90 flavonols, and flavan-3-ols [12]. Recently, many commercial products claiming to be 91 manufactured from cranberry-based extracts have appeared in the market. Some of these 92 products are sold as if they had the same health properties of cranberries, but they do 93

not contain the bioactive polyphenols (i.e. A-type proanthocyanidins among other
polyphenols). This fact shows the importance of developing analytical methodologies
for the characterization of natural extracts to achieve correct authentication regarding
the fruit of origin.

98 Liquid chromatography (LC) with photodiode array (PDA) detection or coupled 99 to mass spectrometry (LC-MS) are among the most common techniques used for the 100 identification, characterization, and determination of polyphenolic compounds in a great 101 variety of plants and fruit-based products [14-21]. High resolution mass spectrometry 102 (HRMS) has also been proposed for the analysis and characterization of polyphenols in 103 fruit products [14,17,21-23]. For instance, Iswaldi et al. [22] proposed the use of time-104 of-flight mass spectrometry (TOF-MS) for the study of the phenolic fraction in cranberry syrup, and Vallverdu-Queralt et al. [17] characterized tomato polyphenols by 105 liquid chromatography-electrospray-linear ion trap quadrupole Orbitrap mass 106 107 spectrometry. Although reversed-phase chromatographic methods are very popular for 108 the determination of low molecular mass flavonoids, a large proportion of this family of 109 compounds in fruits and vegetables consists of highly condensed polymeric 110 proanthocyanidins. Under these circumstances, LC-MS and LC-HRMS play an important role to help in the characterization of PACs in natural extracts [24-26]. 111

Characterization and classification of fruit-based products, including some 112 commercial pharmaceutical preparations, can be tackled from the compositional profiles 113 114 as a source of analytical information. Polyphenolic compounds, as well as other low 115 molecular weight organic acids, alcohols, esters, etc., have been also found to be 116 efficient descriptors of some climatic, agricultural and technological features and, thus, 117 the variability of compounds will strongly depend on the fruit of origin [16,27-29]. Therefore, the polyphenolic profile could be a useful platform for reliable 118 119 discrimination between fruit-based products via chemometric methods such as principal 120 component analysis (PCA). Information recovered mathematically might be essential in order to prevent misuses in the production of commercial fruit-based products with 121 122 health-promoting properties.

This work aims to develop a liquid chromatography-electrospray-tandem mass spectrometry (LC-ESI-MS/MS) method for the identification and determination of polyphenolic profile in fruit-based products and natural extracts. For this purpose, a total of 26 polyphenolic compounds belonging to different families (stilbenes, phenolic acids, and flavonoids) were selected, and a simple sample treatment, consisting of an

extraction by sonication with acetone:water:hydrochloric acid (70:29.9:0.1 v/v/v) and centrifugation, was applied [21]. Different kinds of cranberry-based and grape-based samples were analyzed, including fruits, fruit juices, and raisins, as well as commercial cranberry-based products such as pharmaceutical natural extracts, powder capsules, syrup and sachets. Data corresponding to the polyphenolic composition were considered as a source of potential descriptors to be exploited for the authentication of fruit-based products.

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# 136 **2. Materials and Methods**

## 137 **2.1. Chemicals**

138 Unless otherwise stated, all reagents were of analytical grade. Gallic acid, protocatechualdehyde, (+)-catechin hydrate, gentisic acid, p-salicylic acid, chlorogenic 139 acid, caffeic acid, (-)-epicatechin, (-)-epigallocatechin gallate, syringic acid, 140 syringaldehyde, ethyl gallate, umbelliferon, p-coumaric acid, taxifolin, polydatin, ferulic 141 142 acid, sinapic acid, resveratrol, quercitrin hydrate, fisetin and kaempferol were obtained 143 from Sigma-Aldrich (Steinhein, Germany). Homogentisic acid, protocatechuic acid and 144 vanillic acid were purchased from Fluka (Steinhein, Germany), and quercetin dihydrate 145 from Riedal-de Haën (Seelze, Germany).

Formic acid (98-100 %) was provided by Merck (Darmstadt, Germany). LC-MSgrade methanol and water were purchased from Sigma-Aldrich.

148 Stock standard solutions of all polyphenols (~1000 mg/L) were prepared in 149 methanol in amber-glass vials. Intermediate working solutions were prepared weekly 150 from these stock standard solutions by appropriate dilution with water. All stock 151 solutions were stored at 4 °C for not more than 1 month.

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#### 2.2. Instrumentation and methods

Chromatographic separation was performed on an Accela liquid chromatography 154 system (Thermo Fisher Scientific, San José, CA, USA), equipped with a quaternary 155 pump, an autosampler and a column oven. A Kinetex C<sub>18</sub> reversed-phase column (100 x 156 4.6 mm, 2.6 µm particles) provided by Phenomenex (Torrance, CA, USA) was used for 157 158 the proposed method. Gradient separation was created from solvent A (0.1 % formic acid aqueous solution) and solvent B (methanol) as follows: 0-3 min, linear gradient 159 from 5 to 25 % B; 3-6 min, at 25 % B; 6-9 min, from 25 to 37 % B; 9-13 min, at 37 % 160 B; 13-18 min, from 37 to 54 % B; 18-22 min, at 54 % B; 22-26 min, from 54 to 95 % B; 161

162 26-29 min, at 95 % B; 29-29.15 min, back to initial conditions at 5 % B; and from 29.15
163 to 36 min, at 5 % B. The mobile phase flow rate was 1 mL/min.

The mass spectrometer was a TSQ Quantum Ultra AM (Thermo Fisher 164 165 Scientific) triple quadrupole equipped with heated-electrospray (H-ESI) as ionization source in negative mode. Nitrogen (purity > 99.98 %) was used as a sheath gas, ion 166 167 sweep gas and auxiliary gas at flow-rates of 65, 0 and 40 a.u. (arbitrary units), 168 respectively. Both H-ESI vaporizer temperature and ion transfer tube temperature were 169 set at 350 °C, and the electrospray voltage at -2.5 kV. Full-scan MS acquisition mode (m/z 50-500) in Q1 (mass resolution of 0.7 m/z FWHM, full width half maximum) with 170 171 an scan time of 0.5 s was primarily used for characterization and evaluation. Selected reaction monitoring (SRM) acquisition mode (mass resolution of 0.7 m/z FWHM on 172 173 both Q1 and Q3), with a scan width of 0.5 m/z and a scan time of 0.01 s, was used for 174 quantification purposes by monitoring two SRM transitions. Argon was used as 175 collision gas at 1.0 mTorr and the optimum collision energy (CE) for each transition 176 monitored (quantifier and qualifier) is shown in Table 1. For LC-MS experiments, a 1:1 post-column split of the chromatographic eluent, by means of a Valco zero dead volume 177 178 tee piece, was used.

To optimize both the H-ESI source and tandem mass spectrometry working conditions, 5 mg/L stock standard solution of each compound prepared in methanol:water (1:1 v/v) was infused at a flow-rate of 15  $\mu$ L/min using the syringe pump integrated in the TSQ instrument, and mixed with 500  $\mu$ L/min of a 0.1 % formic acid aqueous solution:methanol (1:1 v/v) mobile phase, by means of a Valco zero dead volume tee piece (Supelco, Gland, Switzerland). Precursor and product ion assignments are also indicated in Table 1.

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## 187 **2.3. Sample treatment**

Different classes of fruit-based products: two fruit samples (cranberry and grapes), five raisin samples (2 based on cranberry and 3 based on grapes), and six juice samples (3 based on cranberry and 3 based on grapes) from different trademarks were purchased from Barcelona markets. In addition, a total of 10 raw extract materials and commercial cranberry-based pharmaceutical preparations presented as powder capsules, syrup, sachets, and natural extracts were provided by Deiters S.L. Company (Barcelona, Spain).

Prior to sample treatment, fruits, raisins and liquid samples (juices and cranberry
pharmaceutical syrup) were freeze-dried to achieve a fully lyophilized products with a
texture similar to that of natural extracts and commercial pharmaceutical samples
(powdered samples). To this end, samples remained 24 h inside a lyophilizer from -80
°C to room temperature, and then were kept for 6.5 h at 40 °C.

Sample treatment was carried-out following a previously described method with some modifications [21,30]. Briefly, 0.1 g of sample were dispersed in 10 mL of acetone:water:hydrochloric acid (70:29.9:0.1 *v:v:v*) and sonicated for 30 min. After that, the mixture was centrifugated for 15 min at 3500 rpm, and the extracts were stored at -4 °C until analyzed. Before injection extracts were filtered through 0.45  $\mu$ m nylon filters (Whatman, Clifton, NJ, USA).

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# 207 2.4. Data analysis

MATLAB (Version 6.5) was used for calculations. Principal component analysis
(PCA) was from the PLS-Toolbox (Eigenvector Research Inc., Mason, WA, USA) [31].
A detailed description of this method is given elsewhere [32].

211 The data matrix to be treated consisted of concentration values of quantified polyphenols in the different samples under study (see section 2.3). The dimension of the 212 213 matrix was 23 samples x 26 analytes). Since concentrations of some pharmaceutical samples were 3 orders of magnitude higher than those occurring in the fruit samples 214 215 (fruit, raisins and juices), normalization pretreatment with respect to the overall polyphenolic concentration was required to provide similar weights to all the samples. 216 217 The plot of scores showing the distribution of the samples on the principal components 218 (PCs) revealed patterns that may be correlated to sample characteristics, such as source 219 fruit in this case. The study of the distribution of variables from the loading plot 220 provided information dealing with their correlations as well as dependences of 221 polyphenols on fruit product properties.

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#### 223 **3. Results and discussion**

# 224 **3.1. Chromatographic separation**

The chromatographic separation was carried out with a Kinetex  $C_{18}$  reversedphase (100 x 4.6 mm, 2.6  $\mu$ m particles) column and gradient elution with 0.1 % formic acid aqueous solution and methanol mobile phases, as previously established by HPLC with UV absorbance detection [21]. In comparison with the previous HPLC-UV 229 method, a 1:1 post-column split of the chromatographic eluent was applied using a 230 Valco T piece to make compatible chromatographic and MS conditions. As a result, the flow rate of mobile phase entering into the MS instrument was 500  $\mu$ L min<sup>-1</sup>. Under 231 232 these conditions, an acceptable chromatographic separation of the 26 polyphenolic 233 compounds was obtained (Figure 1) in less than 22 min. However, several full or partial 234 co-elutions occurred such as those of caffeic acid, epicatechin and epigallocatechin 235 gallate (peaks 10, 11 and 12), taxifolin, polydatin, ferulic acid and sinapic acid (peaks 236 18, 19, 20 and 21), and quercitrin hydrate and fisetin (peaks 23 and 24).

237 However, baseline chromatographic separation is not mandatory because co-238 elutions can be selectively resolved by MS using the appropriate SRM transitions (see Table 1) if no ion suppression effects were present. To study the ion suppression effect, 239 these co-eluting compounds were analyzed by triplicate with the proposed method both 240 individually and in the corresponding co-eluting mixtures. As an example, Figure 1S 241 242 (supplementary material) shows the signals obtained for caffeic acid, epicatechin and 243 epigallocatechin gallate compounds (peaks 10, 11 and 12) at a concentration of 500 244 µg/L. For all evaluated compounds, analysis of variance (ANOVA) was applied 245 showing that for a 95% confidence level polyphenolic peak signals when analyzed individually and in co-eluting mixtures were not significantly different (p values always 246 247 higher than 0.05), so no ion-suppression effects were observed.

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#### 249 **3.2. LC-MS/MS conditions**

250 The liquid chromatographic system was coupled to a triple quadrupole mass 251 spectrometer using an H-ESI source in negative mode. Full scan MS spectra from m/z252 50-500 were recorded. For all polyphenols under study the base peak was the 253 deprotonated molecule [M-H]. Also, neither adducts nor in-source collision-induced 254 dissociation ions were observed in the MS spectra of the compounds at significant intensities except for taxifolin and polydatin which showed in-source fragmentation at 255 relative intensities above 30 % and 50 %, respectively. Thus, the deprotonated molecule 256 257 was selected as precursor ion for all the studied compounds in tandem MS 258 fragmentation experiments.

The fragmentation of these compounds in the triple quadrupole was studied under tandem MS conditions. For the correct product ion assignment, collision energy curves (5-80 eV) were studied. Some similarities were found in the fragmentation of the studied families of polyphenols. For instance, the compounds belonging to the flavonoid

family ((+)-catechin, (-)-epicatechin, (-)-epigallocatechin gallate, taxifolin, fisetin, 263 264 quercetin and kaempferol) presented highly fragmented product ion spectra, with most product ions arising from cross-fragmentation of the aromatic rings in their structures. 265 266 For this reason, sensitivity for these compounds was expected to be lower than for the 267 rest of the polyphenols studied. Typically, most phenolic acids showed the loss of CO<sub>2</sub> 268 in their product ion scan spectra, along with the losses of radical •CH<sub>3</sub> and/or CH<sub>2</sub>O 269 when methoxy substituents were present in the aromatic ring of the compounds (as 270 happens with ferulic acid, sinapic acid, syringic acid, vanillic acid and syringaldehyde). Moreover, the phenolate and hydroxyphenolate ions (m/z 93 and 108, respectively), 271 272 which are characteristic of polyphenolic compounds, were encountered in most product 273 ion scan spectra, although they were not always the most intense product ions. Lastly, 274 polydatin lost the glycoside ring to yield the resveratrol deprotonated molecule (m/z)275 227) which would then produce product ions resulting from the losses of  $C_2H_2O$ . After 276 studying the product ion scan spectra of the compounds, the most intense and 277 characteristic transitions were selected for both quantitative and confirmation purposes. The assignments for the precursor ion and the two most intense product ions for each 278 279 compound, which were selected as quantifier and qualifier SRM transitions, are given in Table 1, and optimal collision energies for both quantifier and qualifier SRM transitions 280 281 are also indicated.

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# **3.3. Instrumental quality parameters**

284 Instrumental quality parameters of the proposed LC-ESI-MS/MS method under 285 optimal conditions were calculated for the 26 polyphenolic compounds and the figures 286 of merit are given in Table 2. Limits of detection (LODs), based on a signal-to-noise 287 ratio of 3:1, were calculated using standard solutions at low concentration levels, and 288 values down to 12-14  $\mu$ g/L were achieved for 19 of the 26 studied polyphenols, in the range 26-68 µg/L for 5 polyphenols, and only fisetin and kaempferol compounds 289 showed LODs at around 110 µg/L. Limits of quantification (LOQs), based on a signal-290 291 to-noise ratio of 10:1, between 40 and 387  $\mu$ g/L were obtained.

292 Calibration curves based on peak area at concentrations above LOQ to 100 mg/L 293 were established. Good linearity was observed for all compounds with correlation 294 coefficients ( $r^2$ ) higher than 0.991.

Run-to-run and day-to-day precisions for compound quantifications were calculated at three concentration levels, low level (LOQ), middle level (500  $\mu$ g/L), and

high level (10 mg/L). For both fisetin and kaempferol, compounds which showed the 297 highest LOQs, only two concentration levels (low and high ones) were evaluated. In 298 order to obtain the run-to-run precision, five replicate determinations for each 299 300 concentration level were carried out. Day-to-day precision was estimated from 15 301 replicate determinations at each concentration level on 3 non-consecutive days (5 302 replicates each day). For run-to-run precision, relative standard deviations (RSD) values 303 in the ranges 1.2-9.9 %, 1.6-6.8 % and 0.6-8.2 % for low, middle and high concentration 304 levels, respectively, were obtained. In general, good and similar precisions regardless the concentration level evaluated were obtained. Day-to-day precision worsened 305 306 slightly, but RSD values were lower, in any case, than 13.5 %. As a conclusion, good precision was attained for the proposed method even at LOQ levels. 307

As no reference material is available, accuracy was evaluated at the three concentration levels by comparing spiked with calculated concentrations using external calibration. Results were excellent, with relative prediction errors (%) lower than 8.0 %.

The results obtained showed that the proposed LC-MS/MS method was satisfactory in terms of sensitivity, precision and accuracy for the determination of polyphenols.

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# 315 3.4. Determination of polyphenols in fruit-based products and pharmaceutical 316 preparations

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The applicability of the proposed LC-MS/MS method was evaluated in the 318 319 determination of 26 polyphenols in 23 fruit-based products, including fruits, raisins, 320 juices, cranberry-based raw extract materials and commercial products (syrup, sachets, 321 powder capsules and natural extracts). A simple sample treatment, consisting of an 322 extraction by sonication with acetone:water:hydrochloric acid (70:29.9:0.1 v/v/v) and 323 centrifugation, was performed. As an example, Figure 2 shows the LC-ESI-MS/MS 324 chromatograms obtained for the 14 most abundant polyphenolic compounds found in 325 the analysis of a cranberry pill pharmaceutical sample.

Prior to the analysis of fruit-based product samples, two different quantification methods were evaluated: (i) external calibration using standards prepared in water, and (ii) standard addition. For comparison, the analysis of three cranberry and three grape samples with different matrices (fruit, juice and raisin) was carried out by triplicate with both external calibration and standard addition. Results are given in electronic

supplementary material Table 1S. Compound identification was based on retention 331 332 times and ion-ratios between quantifier and qualifier SRM transitions. In general, external calibration provided results similar to those obtained by standard addition. Only 333 334 slightly differences for some compounds were encountered. Results from the two 335 approaches were compared statistically using a paired t-test. For a 95 % confidence level, the results were not significantly different, with p values higher than 0.05 (see 336 electronic supplementary material Table 1S). Hence, external calibration was suited to 337 tackle the quantitative determination of polyphenols in fruit-based products. 338

Table 3 shows the concentration levels of polyphenols found in the analyzed 339 samples. For data simplification, only results for 9 of the 23 analyzed samples are 340 depicted in the table, together with the concentration range observed for each 341 polyphenol. Umbelliferon, resveratrol and fisetin polyphenolic compounds were not 342 343 detected in any of the cranberry-based or grape-based analyzed samples. Among the 344 other studied polyphenols, homogentisic acid, gentisic acid, syringaldehyde, ethyl 345 gallate, sinapic acid and kaempferol were neither detected in grape-based samples. Some differences in the polyphenolic compounds detected when comparing natural 346 347 and/or processed cranberry- and grape-based products were also observed. As an example, Figure 3a compares the concentration level of the 10 most relevant 348 349 polyphenols detected in cranberry and grape fruits, as well as their juices and raisins. As can be seen, chlorogenic acid, epicathechin, coumaric acid, quercitrin and quercetin are 350 351 more characteristic polyphenolic compounds in cranberry fruit and fruit-processed 352 products, while other such as gallic acid and catechin tend to be more abundant in grape 353 fruit products. Regarding cranberry pharmaceutical preparations, higher concentrations 354 of some polyphenolic compounds were found in comparison to fruit and related food 355 samples. As an example, Figure 3b compares the concentration level of the 10 most 356 relevant polyphenols detected in these samples. Catechin, chlorogenic acid, epicatechin, 357 epigallocatechin, quercetin and quercitrin were found at very high concentrations levels with catechin, epicatechin and quercitrin being the most abundant ones (with 358 359 concentrations higher than 3000 mg/kg in some of the samples).

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The interesting differences observed among concentration levels of polyphenolic compounds suggest that polyphenolic profile derived from LC-ESI-MS/MS analysis 361 could be proposed as a feature well-suited for the authentication of fruit-based products. 362

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#### 364 **3.5.** Principal component analysis

The polyphenolic concentrations of the samples under study, determined by the 365 366 proposed LC-ESI-MS/MS method, were analyzed chemometrically to draw relevant patterns dealing with the characteristics of natural and processed products. As the first 367 368 issue to be considered, polyphenolic contents in the extracts and pharmaceutical 369 preparations were 100- to 1000-fold higher than those occurring in the fruits and related 370 food samples. Hence, normalization pretreatment with respect to the overall 371 polyphenolic concentration was required in order to provide similar influences on the 372 chemometric model to all the samples.

Normalized data was treated by PCA and the corresponding results are given in 373 374 Figure 4. As shown in the plot of scores (Figure 4a), grape and cranberry products 375 appeared in different zones so that PCA was basically able to distinguish among the two 376 fruits of origin. In particular, grape and related samples were located to the top-left part 377 of this graph. In contrast, cranberry samples were mainly spread out on the bottom area. 378 A group of cranberry samples was to the left, close to the area of distribution of grape 379 samples. This finding might indicate that compositions in percentages could be rather similar for the two groups. Conversely, it suggested that there were significant 380 381 qualitative differences in the compositional profiles of some cranberry products.

Regarding the map of loading (Figure 4b), it was found that gallic acid and 382 polydatin were characteristic of grape-related samples so they were present in higher 383 proportions in this class of products. Analytes located to the right part of PC1 (e.g., 384 385 sinapic, ferulic, coumaric and chlorogenic acids and quercitrin) were comparatively more abundant in cranberry samples. These results agree with those previously reported 386 387 in the literature, where these last mentioned compounds are relatively more abundant 388 and available in berry products, although no levels in cranberry were reported [7]. 389 Catechin was found to the left on PC1. In fact, it has been reported in grape samples at 390 levels between 30-175 mg/kg [7]. Indeed, catechin could be released from the degradation of polymeric condensed tannins, typically occurring in high amounts in 391 392 cranberries, so this component might be a potential index of decay processes.

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#### **4.** Conclusions

The results obtained in this work show that the developed LC-ESI-MS/MS method, using a simple external calibration, can be proposed as a suitable method for the determination of polyphenols in fruit-based products and pharmaceutical preparations. LC-ESI-MS/MS showed a good performance, with low limits of detection for most of the studied compounds (down to 12-14  $\mu$ g/L), and with very good precisions (RSD lower than 13.5 %) and accuracies (relative errors lower than 8.0 %). The method was applied to the analysis of 23 grape-based and cranberry-based products and pharmaceutical preparations after a simple sample extraction procedure consisting of an acetone:water:hydrochloric acid extraction by sonication.

Among the 26 polyphenolic compounds analyzed, only three (umbelliferon, resveratrol and fisetin) were not detected in any of the analyzed samples, and other polyphenols such as homogentisic, gentisic and sinapic acids, syringaldehyde, ethyl gallate, and kaempferol were neither detected in grape-based products.

408 Regarding cranberry-based pharmaceutical preparations, extremely higher concentration of some polyphenolic compounds such as catechin, epicatechin and 409 quercitrin were found in comparison to fruit and related food products. The interesting 410 411 differences observed among concentration levels of some polyphenolic compounds between grape-based and cranberry-based products, as well as between pharmaceutical 412 413 preparations and related food products, suggest that polyphenolic concentrations 414 determined by LC-ESI-MS/MS could be proposed as a suitable source of potential 415 descriptors to be exploited for the authentication of fruit-based products. Results from 416 PCA proved that such polyphenolic concentration data allowed the analyzed samples to be clustered according to their source fruit. 417

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Figure captions
Fig. 1. LC-ESI-MS chromatogram for a mixture of the 26 analyzed polyphenols at 500
$\mu$ g/L. Peak identification as in Table 1.
Fig. 2. LC-ESI-MS/MS chromatograms of 14 selected polyphenols found in a cranberry
pill pharmaceutical sample.
Fig. 3. Concentration levels of 10 selected polyphenols in (a) cranberry and grape fruit,
juices and raisins, and (b) cranberry-based pharmaceutical preparations.
Fig. 4. PCA results using normalized concentrations as the analytical dada. (a) Scatter
plot of scores of PC1 and PC2; Grape samples in green circles, cranberry samples in red
circles. F: fruit; J: juice; R: raisin (dried sample); E: extract; S: sachet; P: pill; and Sy:
syrup. (b) Scatter plot of loadings of PC1 and PC2. Dashed line indicates the separation
among cranberry- and grape-based samples

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Peak	Compound	Precursor ions	Product ion assignment (quantifier/qualifier)	Collision energy (CE, eV)
1	Gallic acid	169.0 [M-H] <sup>-</sup>	125.1 [M-H-CO <sub>2</sub> ] <sup>-</sup>	15
			79.0 $[M-H-C_2H_2O_4]^{-1}$	23
2	Homogentisic acid	167.1 [M-H] <sup>-</sup>	123.0 [M-H-CO <sub>2</sub> ] <sup>-</sup>	13
			122.1 [M-H-CHO <sub>2</sub> ]	23
3	Protocatechuic acid	153.0 [M-H] <sup>-</sup>	109.0 [M-H-CO <sub>2</sub> ] <sup>-</sup>	16
			108.0 [M-H-CHO <sub>2</sub> ] •-	24
4	Protocatechualdehyde	137.0 [M-H] <sup>-</sup>	108.0 [M-H-CHO] *-	22
			92.0 [M-H-CHO <sub>2</sub> ]	25
5	(+)-Catechin hydrate	289.1 [M-H] <sup>-</sup>	245.1 $[M-H-C_2H_4O]^-$	15
			$203.1 [M-H-C_4H_6O_2]^{-1}$	20
5	Gentisic acid	153.1 [M-H] <sup>-</sup>	108.0 [M-H-CHO <sub>2</sub> ] •-	22
			109.0 [M-H-CO <sub>2</sub> ] <sup>-</sup>	14
7	p-Salicylic acid	137.0 [M-H] <sup>-</sup>	93.0 [M-H-CO <sub>2</sub> ] <sup>-</sup>	16
			$65.0 [M-H-C_2O_3]^{-1}$	30
8	Chlorogenic acid	353.0 [M-H] <sup>-</sup>	191.1 $[M-H-C_9H_6O_3]^-$	21
			85.0 [M-H-C <sub>12</sub> H <sub>12</sub> O <sub>7</sub> ] <sup>-</sup>	44
)	Vanillic acid	167.1 [M-H] <sup>-</sup>	152.1 [M-H-CH <sub>3</sub> ] •-	15
			$108.0 [M-H-C_2H_3O_2]^{-1}$	18
10	Caffeic acid	179.1 [M-H] <sup>-</sup>	135.1 [M-H-CO <sub>2</sub> ] <sup>-</sup>	16
			134.1 [M-H-CHO <sub>2</sub> ] •-	25
11	(-)-Epicatechin	289.1 [M-H] <sup>-</sup>	245.1 $[M-H-C_2H_4O]^-$	16
			$203.1 [M-H-C_4H_6O_2]^{-1}$	20
12	(-)-Epigallocatechin gallate	457.0 [M-H] <sup>-</sup>	$169.0 [M-H-C_{15}H_{12}O_6]^{-1}$	19
			125.1 [M-H-C <sub>16</sub> H <sub>12</sub> O <sub>8</sub> ] <sup>-</sup>	39
13	Syringic acid	197.0 [M-H] <sup>-</sup>	182.1 [M-H-CH <sub>3</sub> ] •-	14
			123.0 [M-H-C <sub>2</sub> H <sub>2</sub> O <sub>3</sub> ] <sup>-</sup>	24
14	Syringaldehyde	181.0 [M-H] <sup>-</sup>	166.0 [M-H-CH <sub>3</sub> ] <sup>•-</sup>	13

197.0 [M-H]<sup>-</sup>

161.0 [M-H]<sup>-</sup>

163.1 [M-H]<sup>-</sup>

303.0 [M-H]<sup>-</sup>

389.1 [M-H]<sup>-</sup>

193.1 [M-H]<sup>-</sup>

223.0 [M-H]-

227.0 [M-H]<sup>-</sup>

447.0 [M-H]<sup>-</sup>

285.0 [M-H]<sup>-</sup>

301.1 [M-H]<sup>-</sup>

285.0 [M-H]<sup>-</sup>

151.0 [M-H-CH<sub>2</sub>O]<sup>-</sup>

169.0 [M-H-C<sub>2</sub>H<sub>4</sub>]<sup>-</sup>

133.0 [M-H-CO] 105.0 [M-H-C<sub>3</sub>H<sub>4</sub>O]<sup>-</sup>

119.1 [M-H-CO<sub>2</sub>]

285.0 [M-H-H<sub>2</sub>O]

93.1 [M-H-C<sub>3</sub>H<sub>2</sub>O<sub>2</sub>]<sup>-</sup>

175.0 [M-H-C<sub>6</sub>H<sub>8</sub>O<sub>3</sub>]<sup>-</sup>

227.1 [M-H-C<sub>6</sub>H<sub>10</sub>O<sub>5</sub>]<sup>-</sup> 185.1 [M-H-C<sub>8</sub>H<sub>12</sub>O<sub>6</sub>]<sup>-</sup>

134.1 [M-H-C<sub>2</sub>H<sub>3</sub>O<sub>2</sub>]

164.1 [M-H-C<sub>2</sub>H<sub>3</sub>O<sub>2</sub>] -

143.1 [M-H-C<sub>4</sub>H<sub>4</sub>O]<sup>-</sup>

185.0 [M-H-C<sub>2</sub>H<sub>2</sub>O]<sup>-</sup>

300.1 [M-H-C<sub>6</sub>H<sub>11</sub>O<sub>4</sub>]

271.0 [M-H-C7H12O5]

135.0 [M-H-C<sub>8</sub>H<sub>6</sub>O<sub>3</sub>]<sup>-</sup>

121.1 [M-H-C<sub>9</sub>H<sub>4</sub>O<sub>5</sub>]<sup>-</sup>

151.1 [M-H-C<sub>8</sub>H<sub>6</sub>O<sub>3</sub>]<sup>-</sup>

179.0 [M-H-C<sub>7</sub>H<sub>6</sub>O<sub>2</sub>]<sup>-</sup>

185.0 [M-H-C<sub>4</sub>H<sub>4</sub>O<sub>3</sub>]<sup>-</sup>

117.0 [M-H-C<sub>7</sub>H<sub>6</sub>O<sub>3</sub>]<sup>-</sup>

178.1 [M-H-CH<sub>3</sub>]

208.0 [M-H-CH<sub>3</sub>]

124.0 [M-H-C<sub>3</sub>H<sub>5</sub>O<sub>2</sub>] -

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Ethyl gallate

Umbelliferon

Taxifolin

Polydatin

Ferulic acid

Sinapic acid

Resveratrol

Kaempferol

Fisetin

Quercitrin hydrate

Quercetin dihydrate

p-coumaric acid

Peak	Compound	LOD	LOQ	Working range	run-to-run	precision, %RSD	(n=5)	day-to-day p	Accuracy <sup>a</sup>		
		(µg/L)	(µg/L)	(mg/L)	Low level	Middle level High level		Low level	Middle level High level		(% relative error)
					(LOQ)	(500 μg/L)	U	(LOQ)	(500 μg/L)	(10 µg/L)	
1	Gallic acid	13	43	0.043-100	4.8	5.5	3.0	8.3	4.7	4.5	0.6-6.1
2	Homogentisic acid	12	40	0.040-100	4.4	2.5	3.4	5.6	7.7	4.0	1.9-7.8
3	Protocatechuic acid	14	47	0.047-100	4.9	1.6	1.6	9.0	6.2	4.7	1.1-3.5
4	Protocatechualdehyde	14	47	0.047-100	3.7	4.5	2.9	8.7	7.8	3.8	0.5-1.3
5	(+)-Catechin hydrate	13	43	0.043-100	5.9	3.6	1.3	6.2	8.2	4.8	0.2-7.7
6	Gentisic acid	14	47	0.047-100	8.2	5.6	2.8	13.4	8.2	4.9	0.3-2.1
7	p-Salicylic acid	13	43	0.043-100	7.3	2.1	5.2	8.5	6.7	7.2	0.1-4.6
8	Chlorogenic acid	13	43	0.043-100	7.8	3.5	2.5	9.6	6.5	5.8	4.1-5.8
9	Vanillic acid	14	47	0.047-100	9.9	5.0	4.1	12.9	12.5	7.5	0.6-5.1
10	Caffeic acid	31	103	0.103-100	6.8	1.6	1.3	8.8	3.8	3.5	0.4-2.1
11	(-)-Epicatechin	26	87	0.087-100	4.8	1.6	2.1	8.1	5.6	4.1	0.3-4.2
12	(-)-Epigallocatechin gallate	32	107	0.107-100	1.2	2.3	1.7	9.6	4.5	6.2	3.5-5.0
13	Syringic acid	14	47	0.047-100	6.3	5.8	2.1	13.5	7.9	7.4	1.8-8.0
14	Syringaldehyde	13	43	0.043-100	3.2	1.9	3.2	6.8	5.2	5.2	0.4-2.5
15	Ethyl gallate	13	43	0.043-100	6.3	2.6	0.6	7.7	3.5	4.9	0.3-5.6
16	Umbelliferon	13	43	0.043-100	4.0	4.6	3.4	5.4	5.4	3.6	0.1-0.4
17	p-coumaric acid	34	113	0.113-100	9.2	2.3	2.6	8.7	5.6	4.9	1.2-3.3
18	Taxifolin	13	43	0.043-100	6.7	4.5	3.8	8.6	6.3	4.4	1.5-2.4
19	Polydatin	13	43	0.043-100	6.7	2.8	3.2	10.0	7.7	5.4	1.4-2.9
20	Ferulic acid	13	43	0.043-100	4.1	6.8	8.2	7.3	6.1	8.1	1.3-4.8
21	Sinapic acid	14	47	0.047-100	3.4	6.1	1.3	6.3	5.6	4.2	0.1-0.9
22	Resveratrol	68	227	0.227-100	2.8	3.2	1.8	7.2	6.4	3.1	1.2-4.9
23	Quercitrin hydrate	14	47	0.047-100	3.1	3.9	5.1	11.9	7.5	6.7	3.1-5.6
24	Fisetin	116	387	0.387-100	6.6	-	4.0	6.8	-	5.6	3.1-4.2
25	Quercetin dihydrate	39	130	0.130-100	4.9	3.0	3.6	5.3	4.4	3.8	1.1-2.6
26	Kaempferol	111	370	0.370-100	4.4	-	4.7	5.3	-	4.8	0.2-2.8

 Table 2. Instrumental quality parameters

<sup>a</sup> Accuracy range for all evaluated concentration levels

Peak	Compound	Cranberry-based samples						Grape-base				
		juice 2	syrup	raw extract 1	sachet 1	capsules 1	concentration	juice 2	juice 3	raisin 2	raisin 3	concentration
							range					range
1	Gallic acid	LOD	10.9±0.4	150.4±0.8	10.0±0.6	38.0±0.8	2.2-235.6	15.7±0.8	70.9±1.6	4.4±0.3	6.0±0.2	3.3-99.7
2	Homogentisic acid	nd	nd	6.4±0.2	nd	$9.2\pm0,1$	1.2-11.0	nd	nd	nd	nd	nd
3	Protocatechuic acid	22.4±0.9	$172.8 \pm 1.1$	$904.8 \pm 4.1$	$230.8 \pm 1.8$	370.6±2.0	6.5-904.8	8.1±0.2	$22.4\pm0.5$	$10.6 \pm 0.2$	LOQ	3.0-22.4
4	Protocatechualdehyde	LOD	LOD	13.7±0.4	LOD	45.5±0.5	1.2-125.2	LOD	LOD	LOD	LOD	1.3-2.6
5	(+)-Catechin hydrate	4.3±0.2	nd	$142.8 \pm 1.6$	13.5±0.2	3363.0±4.6	2.6-7383.3	$10.0\pm0.1$	$70.5 \pm 0.5$	nd	7.1±0.1	7.1-109.4
6	Gentisic acid	nd	LOD	16.0±0.3	LOD	LOD	1.1-36.6	nd	nd	nd	nd	nd
7	p-Salicylic acid	nd	LOD	5.6±0.1	nd	LOD	1.5-5.6	nd	nd	nd	nd	5.0
8	Chlorogenic acid	$14.0\pm0.3$	$88.2 \pm 0.8$	368.5±1.1	44.7±0.7	22.1±0.6	6.1-368.5	nd	nd	nd	nd	6.8
9	Vanillic acid	nd	$16.0\pm0.4$	135.0±0.8	LOD	nd	1.7-135.0	nd	nd	nd	nd	4.2-7.1
10	Caffeic acid	nd	19.6±0.1	248.5±0.8	LOD	$16.9 \pm 0.1$	3.0-248.5	nd	LOD	nd	nd	4.3-5.6
11	(-)-Epicatechin	13.7±0.3	nd	$1038.8 \pm 3.4$	$344.8 \pm 0.8$	3239.6±2.6	1.3-7297.6	2.3±0.3	23.7±0.5	nd	nd	1.3-43.2
12	(-)-Epigallocatechin gallate	nd	nd	nd	nd	128.9±0.1	12.5-1425.7	nd	nd	nd	nd	6.0
13	Syringic acid	nd	LOD	33.6±0.3	LOD	nd	1.6-304.2	8.7±0.2	8.1±0.1	nd	nd	8.1-13.4
14	Syringaldehyde	nd	nd	10.1±0.4	nd	nd	6.4-10.1	nd	nd	nd	nd	nd
15	Ethyl gallate	nd	nd	nd	nd	nd	2.3-303.2	nd	nd	nd	nd	nd
16	Umbelliferon	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
17	p-coumaric acid	$14.1 \pm 0.7$	206.0±0.8	760.4±1.9	LOD	43.3±0.7	1.5-760.4	nd	4.3±0.1	nd	nd	4.3-4.8
18	Taxifolin	nd	nd	273.5±0.7	$8.6\pm0.1$	75.3±0.5	3.6-273.5	nd	nd	nd	nd	nd
19	Polydatin	nd	nd	5.4±0.4	nd	8.7±0.7	5.4-16.5	LOD	12.9±0.3	nd	LOD	1.3-12.9
20	Ferulic acid	LOD	33.8±0.7	93.0±1.0	LOD	6.4±0.1	1.2-93.0	nd	LOD	LOD	nd	1.0-1.7
21	Sinapic acid	nd	19.9±0.5	24.1±0.5	LOD	LOD	1.9-50.9	nd	nd	nd	nd	nd
22	Resveratrol	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
23	Quercitrin hydrate	16.7±0.5	89.1±0.8	$1228.6 \pm 1.2$	nd	24.3±0.6	3.6-1857.5	6.7±0.1	LOD	nd	nd	1.6-6.7
24	Fisetin	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
25	Quercetin dihydrate	21.8±0.4	635.1±0.9	3356.1±3.2	91.6±0.1	572.3±1.1	3.9-3526.7	nd	$14.8\pm0.2$	LOD	5.6±0.1	3.7-14.8
26	Kaempferol	nd	nd	$16.8\pm0.4$	nd	53.2±0.3	16.8-130.0	nd	nd	nd	nd	nd

Table 3. Polyphenol concentration levels in cranberry-based and grape-based products<sup>a</sup>

<sup>a</sup> All concentrations are in mg/kg. Quantifications performed by triplicate (n=3); results are expressed as mean of samples analyzed  $\pm$  standard deviation; nd, not detected.















Figure 1S. Ion suppression study for three coeluting polyphenols (caffeic acid, epicatechin and epigallocatechin gallate).



Peak	Compound	Cranberry fruit		Cranberry juice		Cranberry raisins		Grape fruit		Grape jui	ce 1	Grape raisins	
		EC	SA	EC	SA	EC	SA	EC	SA	EC	SA	EC	SA
1	Gallic acid	nd	nd	LOD	LOD	LOD	LOD	$14.0 \pm 1.2$	12.4±0.2	99.7±4.3	99.5±1.4	LOD	LOD
2	Homogentisic acid	LOD	LOD	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
3	Protocatechuic acid	6.5±0.1	3.9±0.4	26.7±2.0	29.6±0.8	9.4±1.5	$11.9\pm0.8$	LOD	LOD	$18.7 \pm 2.6$	16.2±0.2	LOD	LOD
4	Protocatechualdehyde	nd	nd	nd	nd	LOD	LOD	LOD	LOD	LOD	LOD	nd	nd
5	(+)-Catechin hydrate	$46.5 \pm 2.5$	50.4±0.5	nd	nd	nd	nd	$109.4{\pm}10.8$	112.9±10.2	$57.0\pm5.1$	$54.6 \pm 4.4$	9.2±0.1	$13.4 \pm 0.2$
6	Gentisic acid	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
7	p-Salicylic acid	nd	nd	nd	nd	nd	nd	nd	nd	$5.0\pm0.8$	3.9±0.9	nd	nd
8	Chlorogenic acid	$160.4 \pm 5.6$	$165.2 \pm 1.1$	$14.2 \pm 1.3$	17.7±1.1	6.3±0.8	8.2±1.1	nd	nd	6.8±0.7	3.7±0.3	nd	nd
9	Vanillic acid	5.1±0.5	$4.4\pm0.4$	6.9±0.6	7.2±0.5	LOD	LOD	LOD	LOD	7.1±0.5	4.3±0.2	nd	nd
10	Caffeic acid	4.4±0.2	7.4±0.5	nd	nd	nd	nd	nd	nd	$5.6\pm0.6$	7.4±0.2	nd	nd
11	(-)-Epicatechin	296.8±11.7	257.3±0.2	9.3±1.1	8.6±0.2	LOD	LOD	43.2±2.9	37.7±2.3	26.1±0.6	30.6±0.3	nd	nd
12	(-)-Epigallocatechin gallate	nd	nd	nd	nd	nd	nd	6.0±0.4	3.0±0.6	nd	nd	nd	nd
13	Syringic acid	nd	nd	nd	nd	nd	nd	nd	nd	$13.4\pm0.8$	17.6±0.3	nd	nd
14	Syringaldehyde	$6.4\pm0.4$	4.3±0.2	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
15	Ethyl gallate	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
16	Umbelliferon	nd	nd	nd	nd	nd	dn	nd	nd	nd	nd	nd	nd
17	p-coumaric acid	$12.4{\pm}1.2$	15.9±0.8	14.7±1.7	$18.8\pm0.8$	LOD	LOD	nd	nd	4.6±0.1	9.9±0.7	nd	nd
18	Taxifolin	LOD	LOD	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
19	Polydatin	nd	nd	nd	nd	nd	nd	8.9±0.9	9.6±0.7	7.4±0.3	3.4+0.1	nd	nd
20	Ferulic acid	4.3±0.4	5.5±0.6	$4.4\pm0.6$	$5.5 \pm 1.1$	nd	nd	nd	nd	LOD	LOD	nd	nd
21	Sinapic acid	LOD	LOD	LOD	LOD	nd	nd	nd	nd	nd	nd	nd	nd
22	Resveratrol	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
23	Quercitrin hydrate	97.9±4.7	93.1±1.2	23.6±1.8	25.0±1.2	3.6±0.2	7.5±1.2	2.2±0.4	5.0±0.7	nd	nd	nd	nd
24	Fisetin	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
25	Quercetin dihydrate	30.0±1.6	26.2±0.4	7.3±0.5	6.6±0.4	14.1±3.3	$10.4 \pm 0.4$	4.3±0.1	4.3±0.1	$7.0\pm0.1$	3.8±0.3	9.1±0.3	4.3±0.4
26	Kaempferol	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd	nd
<i>p</i> value	e <sup>b</sup>	0.38		0.06		0.54		0.73		0.77		0.95	

Table 1S. Comparison of calibration procedures for polyphenol quantification in fruit-based products by LC-ESI-MS/MS<sup>a</sup>

<sup>a</sup> All concentrations are in mg/kg. Quantifications performed by triplicate (n=3); results expressed as concentration mean of samples analyzed ± standard deviation. EC, external calibration; SA, standard addition; nd, not detected <sup>b</sup> For a 95 % confidence level.