

Establishment of a Spectral Database for Classification of Edible Oils using Matrix-assisted Laser Desorption/Ionization Mass Spectrometry

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Abstract

In this study, we aim to establish a comprehensive spectral database for analysis of edible oils using matrix-assisted laser desorption/ionization mass spectrometry (MALDI-MS). More than 900 edible oil samples, including 30 types of edible oils, were analyzed and compared, and the characteristic peaks and spectral features of each edible oil were obtained. Edible oils were divided into eight groups based on their characteristic spectral patterns and principal component analysis results. An overall correct rate of 97.2% (98.1% for testing set) was obtained for classification of 435 edible oil products using partial least square-discriminant analysis, with nearly 100% correct rate for commonly used edible oils. Differentiation of counterfeit edible oils, repeatedly cooked edible oils and gutter oils from normal edible oils could also be achieved based on the MALDI-MS spectra. The establishment of this spectral database provides reference spectra for spectral comparison and allows rapid classification of edible oils by MALDI-MS.

Keywords: Edible oils; gutter oils; MALDI-MS; classification; characteristic peaks; principal component analysis; partial least square-discriminant analysis.

1. Introduction

Edible oils are one of the major sources of fat for human being. Most of the edible oils contain two essential fatty acids, omega-3 and omega-6 fatty acids, which cannot be synthesized by human body and must be absorbed to maintain human health (Gunstone, 2004). Edible oils are thus essential to our daily life. However, the quality and safety of edible oil products in the market are not secured with the widespread of edible oil frauds in recent years (Hong et al., 2017; Peng et al., 2017). Edible oil frauds typically include edible oil counterfeit, which involves using cheaper edible oils to replace more expensive ones (Hong, et al., 2017; Zhu, Wang, & Chen, 2017), edible oil adulteration, which involves mixing edible oils with cheaper ones (Calvano, De Ceglie, D'Accolti, & Zambonin, 2012; Jimenez-Carvelo, Gonzalez-Casado, & Cuadros-Rodriguez, 2017), and the use of gutter oils (i.e., recycled cooking oils), which involves refining wasted or used cooking oils and labeling them as normal edible oils (Peng et al., 2017). Emergence of gutter oils that have potential health problems has raised many public concerns. However, identification of gutter oils is a challenging analytical task, since gutter oils have very diverse preparation processes and sources and it may not be effective to rely on detection of targeted toxic substances (Centre for Food Safety Hong Kong, 2013) or specific food marker compounds (Jin et al., 2013) for determination of gutter oils.

61 Current practice relies on standard gas chromatography-flame ionization detection (GC-FID)
62 method (International Organization for Standardization, 2014) for analysis of edible oils. In
63 this method, the fatty acid content in an edible oil sample is determined using the GC-FID-
64 based method and matched with the Codex standard (Codex Alimentarius, 2015a, 2015b). The
65 disadvantages of this conventional technique, such as the need of chemical derivatization and
66 time-consuming column separation (Cozzolino & Giulio, 2011), induced the development of
67 simple and rapid analytical techniques to meet the high analytical demand of edible oils.
68 Various techniques, such as Raman spectroscopy (Li et al., 2017) laser-induced fluorescence
69 (Mu et al., 2016), infrared spectroscopy (Rohman, 2016), electronic nose (Xu, Yu, Liu, &
70 Zhang, 2016), low field nuclear magnetic resonance (Zhu, et al., 2017), ion mobility
71 spectrometry (L. Zhang et al., 2016), and mass spectrometric techniques, including direct
72 infusion electrospray ionization (Catharino et al., 2005), direct analysis in real time (DART)
73 (Vaclavik, Cajka, Hrbek, & Hajslova, 2009) and matrix-assisted laser desorption/ionization
74 (MALDI) (Ayorinde, Elhilo, & Hlongwane, 1999; Jakab, Nagy, Heberger, Vekey, & Forgacs,
75 2002; Lay, Liyanage, Durham, & Brooks, 2006; Ng, So, Zheng, & Yao, 2015), have been
76 employed for analysis of edible oils. In particular, MALDI-MS has distinctive features over
77 the other techniques, such as short analysis time, high sensitivity and high tolerance to
78 impurities. In MALDI-MS, several hundreds of samples can be loaded onto different spots of

one target plate and introduced into the mass spectrometer for analysis, and spectrum for each sample can be obtained within seconds. Very recently, a simple protocol was developed to allow direct analysis of edible oils, with no memory effect and clear mass spectra for spectral comparison of edible oils (Ng et al., 2015). Studies have proved that MALDI-MS spectra of different edible oil species, especially the triacylglycerides (TAGs) region, were specific and could be used for characterization of edible oils (Lay et al., 2006; Ng et al., 2015; Picariello, Paduano, Sacchi, & Addeo, 2009; Picariello, Sacchi, & Addeo, 2007). Recent studies also demonstrated differentiation of geographical origins of olive oils and detection of olive oil adulteration using MALDI-MS and statistical analysis (Peršurić, Osuga, Galinac Grbac, Peter-Katalinić, & Kraljević Pavelić, 2017; Jergović, Peršurić, Saftić, & Kraljević Pavelić, 2017). Ayorinde and co-workers calculated and compared the fatty acid contents of different edible oils based on their MALDI-MS spectra and GC chromatograms, and concluded that the results obtained from the MALDI-MS and GC methods agreed with each other (Ayorinde, Garvin, & Saeed, 2000; Hlongwane, Delves, Wan, & Ayorinde, 2001).

Although MALDI-MS has been extensively used to analyze edible oils (Ayorinde, Elhilo et al., 1999; Lisa, Holcapek, & Bohac, 2009; Picariello et al., 2007; Picariello et al., 2009; Q. Zhang et al., 2016), so far there has been no systematic investigation and establishment of a

comprehensive MALDI spectral database of various edible oils, which will definitely benefit the technique and the analysis of edible oils. In this study, more than nine hundred edible oil samples were collected and analyzed using the MALDI-MS protocol we developed previously (Ng et al., 2015). The MALDI-MS spectra obtained were stored in an in-house developed system, and a comprehensive MALDI-MS spectral database including more than thirty edible oil species was then established to allow rapid classification of edible oils by comparing the MALDI-MS spectra of the samples with the reference spectra in the database.

2. Experimental section

2.1 Chemicals and materials

Thirty types of edible oil products were collected from collaborators and authentic stores in Hong Kong, mainland China and Taiwan, with origins from various countries or regions (see Table S-1 for the complete list of the oil samples and Table S-2 for their origins). Three bottles were collected for each commercially available oil product. Standards of olive oil, castor oil, corn oil, soybean oil, sesame oil, coconut oil, peanut oil, linseed oil, palm oil, canola oil, cottonseed oil, sunflower seed oil, safflower oil and lard oil were purchased from Supelco and Sigma-Aldrich (St. Louis, USA). The heated oil samples were prepared in the laboratory. The gutter oil samples were provided by Syscan Technology Holdings Limited (Wuhan, China).

All samples were sealed and stored in a dry and dark environment before analysis. MALDI matrices 2, 5-dihydroxybenzoic acid (DHB) and α -cyano-4-hydroxycinnamic acid (CHCA) were purchased from Aldrich (St. Louis, MO, USA). HPLC grade acetone and HPLC grade tert-butyl methyl ether (TBME) were purchased from Acros Organic (Waltham, MA, USA). HPLC grade acetonitrile (ACN) was purchased from Anaqua Chemical Supply (Houston, TX, USA) and HPLC grade dichloromethane (DCM) was purchased from RCI Labscan (Bangkok, Thailand). Polyethylene glycol standards were purchased from Fluka (St. Louis, MO, USA). Sodium iodide (NaI) was purchased from Panreac Química (Barcelona, Spain) and sodium chloride (NaCl) was purchased from Riedel-de Haën (Seelze, Germany). Derivatization reagent, trimethylsulfonium hydroxide (TMSH, 0.25 M solution in methanol) was purchased from Acros Organic, and a mixture of 37 fatty acid methyl standards (FAME) was purchased from Supelco. All chemicals were used directly without further purification.

2.2 MALDI-MS analysis

The edible oil samples were analyzed using the protocol we reported previously (Ng et al., 2015) with slight modification. Briefly, each edible oil sample was directly transferred by a medical cotton tip onto the spot on the MALDI plate, which was pre-deposited with DHB matrix layer. The MALDI plate was then inserted into an ultrafleXtreme MALDI-TOF/TOF

133 mass spectrometer (Bruker, Billerica, MA, USA) equipped with a 355 nm smartbeam-II laser
134 for the MALDI-MS analysis. The mass spectrometer was operated at a frequency of 2000 Hz
135 and in positive and reflectron mode. The matrix suppression cut-off was set to m/z 500 and the
136 spectra were acquired with an m/z range of 500-2000 Da. The ion pulse excitation was set to
137 140 ns. The ion source voltage 1, ion source voltage 2, lens voltage, reflector voltage 1 and
138 reflector voltage 2 were set to 20 kV, 17.75 kV, 7 kV, 21.1 kV and 10.85 kV, respectively. The
139 reflector base detector voltage was set to 2.32 kV and the sample rate was set to 5.0 GS/s. The
140 mass spectrometer was calibrated with the PEG solution mixture
141 (PEG600/PEG1000/PEG2000/NaI = 1/2/2/5 (v/v)). Each mass spectrum was acquired
142 manually by irradiating at least 5 random positions (1000 laser pulse per one position) on the
143 sample spot. The sub-spectra were accepted if the signals in the TAGs region (typically at the
144 region of m/z 570-750 or m/z 850-920) were observed and the absolute intensity of the top peak
145 of the final spectra obtained by combining at least 5 positive sub-spectra reached 1×10^4 . The
146 mass spectra were processed in flexAnalysis 1.4 program (Bruker, Billerica, MA, USA) under
147 “centroid” peak detection algorithm, “SavitzkyGolay” peak smoothing algorithm (the width
148 equal to 0.2 m/z and the cycles equal to 1) and baseline subtraction algorithm “TopHat”. Each
149 edible oil sample was analyzed in triplicate. The sodium adducted TAGs were observed in the
150 MALDI-MS mass spectra and their identification were based on the results of the previous

studies (Ayorinde, Elhilo et al., 1999; Lay et al., 2006; Picariello et al., 2009) and further confirmed with MS/MS analysis in this study (see the Supporting Information). For the TAGs in rapeseed oil, the peak assignments were based on the theoretical calculation of fatty acid contents from the GC-FID results.

2.3 Statistical analysis

The data obtained from three samples of the same edible oil products were averaged. The normalized intensities of monoisotopic peaks of TAGs and cyclolinopeptides in the spectra (absolute intensity of the peak observed / total absolute intensity of all the peaks observed in the mass spectrum) were input into the statistics software (Umetrics Simca 14.0, Andover, MA, USA) for principal component analysis (PCA) and partial least square-discriminant analysis (PLS-DA). The PCA and PLS-DA models were established using “simple mode” and “autofit” (automatic cross-validation) with the default settings. In the PLS-DA, 198 edible oil products, including the products with different brands, collected from different locations and possessed different TAGs patterns, were selected as the training set for establishment of the PLS-DA models. One “grouping” model using the groups as the dummy Y variables and seven sub-models with the species as the dummy Y variables were established. The remaining products were used as testing set to test the performance of the established PLS-DA models.

169

170 **3. Results and discussion**

171 **3.1 MALDI-MS spectra and spectral database of edible oils**

172 The rapid and simple protocol we previously developed for MALDI-MS analysis of edible oils
173 (Ng et al., 2015) was employed in this study. This protocol allows direct loading of edible oil
174 samples on the predeposited DHB matrix layers that can be stable over one week, and offers
175 mass spectra of edible oils with high quality and high reproducibility. The typical spectra of all
176 the 30 edible oils can be found in Figure S-1. To our knowledge, MALDI-MS spectra of some
177 edible oils, e.g., perilla oil, pine nut oil and rice bran oil, have not been reported yet. For those
178 edible oils previously investigated by MALDI-MS, the spectra obtained in the study were
179 consistent with those in the literatures (Ayorinde, Elhilo et al., 1999; Ayorinde, Eribo, Balan,
180 Johnson, & Wan, 1999; Jakab et al., 2002; Lay et al., 2006; Picariello et al., 2007).

181

182 For each oil product from the market, i.e., a specific edible oil of a specific brand, typically
183 three bottles of edible oils were collected and analyzed in this study. The mass spectra obtained
184 from different bottles of the same oil products were highly similar to each other, with an
185 example of the spectra for peanut oil as shown in Figure S-2. However, for the same kind of
186 edible oils of different brands, variations were observed for the spectra obtained. For example,

as shown in Figure S-3, similar ion peaks were detected for the mass spectra obtained for the peanut oil samples from three different brands, but their intensity ratios varied, e.g., different intensity ratios between m/z 903.7 (LLO), m/z 905.8 (LOO) and m/z 907.8 (OOO) obtained from the three different brands of samples. This was consistent with the varied range of C18:1 (35.0-69.0%) and C18:2 (12.0-43.0%) contents of peanut oils as described in the Codex standards (Codex Alimentarius, 2015a). Variations of TAG patterns were also observed for other edible oils but at different extents. Some of the edible oils such as peanut oil and olive oil could have larger variations, while some of the edible oils such as canola oil could have smaller variations. However, as illustrated below, such variations within the same oil species were normally smaller than the differences between different oil species, allowing differentiation of different oil species.

Special consideration was given to canola oil (i.e., low erucic acid rapeseed oil) and rapeseed oil that are two closely related edible oil species. Their fatty acid contents were very similar, except that canola oil contains less than 2% C22:1 (see Figures S-1.5 and S-1.23 for their typical mass spectra). The TAGs of C22:1 were observed at m/z 935.8-1019.9 and their suggested compositions are listed in Table S-7. The collected rapeseed oil samples were confirmed by conventional GC-FID analysis, and it was found that out of the 21 collected

“rapeseed oil” products, only 6 contained more than 2% C22:1 and confirmed as rapeseed oil.

The remaining samples were counted as canola oil in this study.

As shown in Table S-1, more than 900 edible oil samples, including 30 species of edible oils with different brands for each species, were analyzed to construct the spectral database. For the collected edible oil samples that showed MALDI-MS spectra obviously different from those of the same species, confirmatory analysis using the conventional GC-FID method for FAME analysis was performed, and the edible oil samples with fatty acid contents failed to match the Codex standards were rejected and were not included in the database. Some edible oil samples with high levels of oxidized/thermal products were also rejected. The sample information (oil type, origin of manufacture, origin of raw materials, and source of collection), full spectra and the spectra of TAGs region of the authentic samples were then exported and stored into a database on an Apache server. As shown in Figure S-4, an in-house developed user interface (HTML/JavaScript at front-end and PHP at back-end) allows users to retrieve individual spectra grouped by the type of oils.

3.2 Characteristic peaks and PCA of the MALDI-MS spectra

Spectra of different edible oils were compared and characteristic peaks have been found for various edible oils. The mass spectra of castor oil (Figure S-1.6) and fish oil (Figure S-1.10) were easily distinguished from those of other edible oils. Castor oil showed a very simple mass spectrum with a major peak at m/z 955.7 that corresponds to TAG of 12-OH-C18:1 (Ayorinde, Elhilo et al., 1999). Fish oil possessed a complex mass spectrum which contained abundant signals for both TAGs commonly observed in vegetable oils (i.e. TAGs from m/z 850 to m/z 920) and TAGs of C20:5 and C22:6 (Codex Alimentarius, 2014) such as m/z 981.8 and m/z 1009.8, which were only observed in fish oil. These specific MALDI-MS patterns allowed castor oil and fish oil to be easily distinguished based on their spectra.

The TAGs regions of butter (Figure S-1.3) and coconut oil (Figure S-1.7) were mainly the peaks lower than m/z 850 with abundant peaks such as m/z 661.5 and m/z 689.6 which corresponded to TAGs with C12:0 and C14:0, respectively (Saraiva, Cabral, Eberlin, & Catharino, 2009). These TAGs were only found in butter and coconut oil and could be used for differentiation of butter and coconut oil from other edible oils. Comparison between the spectra of butter and coconut oil showed that the spectral pattern of these two species were different. Coconut oil had higher intensities of peaks at m/z 605.5 and m/z 633.5 while butter had higher intensities of peaks at m/z 771.6 and m/z 827.6, i.e., coconut oil contained more short chain

240 fatty acids such as C12:0. Margarine is a mixture of vegetable oils but is physically similar to
241 butter. It is not easy to differentiate them by naked eyes. As shown in Figure S-5, the mass
242 spectra of margarine showed strong signals of higher mass TAGs such as m/z 879.8 and m/z
243 903.7 (i.e. higher level of C18:1 and C18:2), while the mass spectrum of butter contained
244 significant lower mass TAGs peaks as discussed previously. The remarkably different mass
245 spectra of margarine and butter allowed them to be easily distinguished from each other.

246

247 As shown in the MALDI-MS spectra of flaxseed (Figure S-1.11) and perilla oil (Figure S-1.20),
248 two specific peaks at m/z 895.7 (LnLnLn) and m/z 897.7 (LnLnL) were observed and could
249 serve as the markers of these two oils. Peak at m/z 899.7 for flaxseed oil showed obviously
250 higher abundance than for perilla oil, allowing differentiation between these two oils. In
251 addition, some characteristic peaks such as m/z 999.5 and m/z 1062.6 were observed only in
252 flaxseed oil (see Figure S-6a and S-6b). The peaks found in such high mass region for edible
253 oils usually corresponded to thermal/oxidation products which were formed after prolonged
254 heating (Picariello et al., 2009; Q. Zhang et al., 2016). However, accurate mass measurements,
255 database searching and MS/MS analysis (see Figure S-6c and S-6d for the results) indicated
256 that the most abundant peak (m/z 1062.6) in the spectrum corresponded to sodium adducted
257 ion of cyclolinopeptide A, a natural product found in flaxseed oil extract (Gaymes, Cebrat,

Siemion, & Kay, 1997), and the other peaks at m/z 1080.7, m/z 1096.7, m/z 999.6 and m/z 983.5 corresponded to the sodium adducted ions of cyclolinopeptides B, C, E and J (Stefanowicz, 2004). For the 28 flaxseed oil products collected for the analysis, the cyclolinopeptide peaks were detected in 25 products, but not in the other 3 products that were confirmed also as flaxseed oil based on their TAGs distribution. Although the cyclolinopeptides were not observed in all the flaxseed oil products, the presence of such peaks could provide additional information for characterization of flaxseed oil.

Palm oil is one of the most popular edible oil species used in the food industry (Gunstone, 2011). There are seven types of palm oil derivatives that possess different fatty acid contents (Codex Alimentarius, 2015a). However, only two types of palm oil derivatives, i.e., palm oil (Figure S-1.17) and palm superolein (Figure S-1.18), were found in the local markets. Palm oil showed higher abundances of peaks at m/z 855.7 (PPO) and m/z 881.8 (POO) than other common vegetable oils such as canola oil and olive oil. Compared with palm oil, palm superolein had weaker peak at m/z 855.7 but stronger peak at m/z 881.8, as it contains less C16:0 but more C18:1. Lard shared similar MALDI-MS spectral pattern with that of palm superolein.

Rapeseed oil (Figure S-1.23), as discussed in the previous section, shared similar MALDI-MS spectrum with canola oil, except that peaks at higher mass region such as m/z 991.9 and m/z 1019.9, which corresponded to TAGs of C20:1 and C22:1, were observed only in rapeseed oil. However, the intensities of these additional peaks varied at different rapeseed oil products, consistent with the reported C22:1 contents from 2% to 60% in rapeseed oil (Codex Alimentarius, 2015a). The permitted C22:1 content in edible oil products has been regulated in some regions. For example, the threshold level is 5% by weight of fatty acid content in Hong Kong (Harmful Substances in Food Regulations, 1983). This MALDI-MS approach is potentially useful for rapid screening of the illegal edible oil products by detecting the TAGs peaks of C22:1 (Table S-7).

Except for the above edible oil species that showed very obvious characteristic peaks, MALDI-MS spectra of the other edible oil species mainly consisted of similar peaks such as m/z 877.7 (PLL), m/z 879.7 (PLO), m/z 901.7 (LLL), m/z 903.7 (LLO), m/z 905.8 (LOO) and m/z 907.8 (OOO), but with different intensity ratios. Among these edible oil species, highest peak abundances were observed for m/z 877.7 in cottonseed oil (Figure S-1.9), m/z 879.7 in rice bran oil (Figure S-1.24), m/z 901.7 in safflower oil (Figure S-1.25), soybean oil (Figure S-1.27), walnut oil (Figure S-1.29) and wheat germ oil (Figure S-1.30), m/z 903.7 in corn oil (Figure S-

1.8), pine nut oil (Figure S-1.21), pumpkin seed oil (Figure S-1.22) and sunflower oil (Figure S-1.28), m/z 905.8 in sesame oil (Figure S-1.26), m/z 907.8 in almond oil (Figure S-1.1), avocado oil (Figure S-1.2), camellia oil (Figure S-1.4), canola oil (Figure S-1.5), hazelnut oil (Figure S-1.13), high oleic acid sunflower oil (Figure S-1.14) and olive oil (Figure S-1.16). The most abundant peaks in peanut oil varied as discussed before (Figure S-3), which could be either m/z 903.7, m/z 905.8 or m/z 907.8. These edible oil species could be differentiated by comparing the peak intensity ratios in their mass spectra. For example, the mass spectra of both olive oil (Figure S-1.16) and canola oil (Figure S-1.5) had abundant peaks at m/z 907.8, but olive oil showed higher abundance of peak at m/z 881.8, while canola oil showed higher abundance of peak at m/z 905.8 and additional peaks at m/z 899.7, m/z 901.7 and m/z 903.7). Different intensity ratios of m/z 881.8 and m/z 907.8 were observed for olive oil, avocado oil, hazelnut oil and high oleic acid sunflower oil, allowing differentiation among them.

PCA, a commonly used tool for statistical analysis, is used to facilitate the differentiation and classification of edible oils. Excluding castor oil and fish oil samples, which showed spectra that were observably different from those of other oil species, spectra of 435 edible oil products were analyzed using PCA. As shown in Figure 1a, butter and coconut oil were clustered very separately from each other and other edible oils, in consistent with their observably different

TAG patterns. PCA after excluding coconut oil and butter revealed two groups of edible oils, with one group containing flaxseed oil and perilla oil and the other group containing palm oil, lard and palm superolein, which were clustered separately from the remaining edible oils (see Figure 1b). A new PCA score plot (Figure 2a) was obtained for PCA of the remaining edible oils, which could be divided into 4 regions. Based on these PCA results, the edible oils were divided into seven different groups and one additional group of castor oil and fish oil that showed very specific peaks, which are listed with their characteristic peaks in Table 1. The edible oils in groups 2, 3, 4, 5, 6 and 7 could be further differentiated with further PCA as shown in Figures 1c-d and 2b-e. It is noticed that although the MALDI-MS spectra of some edible oils such as peanut oil showed obvious variations within the same species as discussed previously, such variations were normally still smaller than the differences between different oil species, allowing them to be clearly differentiated from other oil species. However, 3 pairs of edible oils, i.e., olive oil and avocado oil, canola oil and almond oil, and corn oil and pumpkin seed oil, were found difficult to be differentiated based on their TAGs patterns (see Figures S-1.2 and S-1.16; S-1.1 and S-1.5; and S-1.8 and S-1.22, and PCA results in Figures 2b, 2c and 2e).

3.3 Classification of edible oils using PLS-DA

PLS-DA is a supervised technique that correlates the *class* (i.e., the edible oil species in this study) and *X-variables* (i.e., the relative intensities of peaks in the MALDI-MS spectra) through the scores calculated by the statistical software (Berrueta, Alonso-Salces, & Heberger, 2007). The *class* of future samples could be predicted after the establishment of PLS-DA models using training samples. This cannot be achieved using the PCA method (Ballabio & Consonni, 2013; Berrueta et al., 2007).

Samples from 198 representative edible oil products were used as the training set for establishment of PLS-DA models in this study. The training set included edible oil products of different species, different collection sources and different MALDI-MS spectral patterns. Based on the PCA results, a two-step approach was employed for the PLS-DA analysis. The samples were firstly classified into the groups (groups 1-7 as shown in Table 1) using the “grouping” PLS-DA model and then further classified into single edible oil species using 7 sub-models (model groups 1-7). The data of samples from the training set were input into the SIMCA software for PLS-DA processing to establish a “grouping” model with the groups as the dummy Y variables. Seven sub-models with the species as the dummy Y variables were established using the training data of the edible oil samples in each group.

The data of 198 edible oil products in the training set were used to establish and cross-validate the model automatically by the software, and 237 edible oil products in the testing set were input into the established models for the external validation. The predicted scores (PS) for the matching of each oil species was calculated for each edible oil sample, and the edible oil sample was then classified as the edible oil species which gave the highest score. For example, the PLS-DA analysis of a sample gave predicted scores of 0.01, -0.02, -0.01, 0.06, 0.18, -0.15 and 0.92 for groups 1, 2, 3, 4, 5, 6 and 7, respectively, indicating that this sample belonged to group 7. Further analysis using the model of group 7 gave predicted scores of 1.16, 0.03, 0.04, -0.04, and -0.19 for the five edible oils in group 7, i.e., olive oil, camellia oil, high oleic acid sunflower oil, avocado oil, and hazelnut oil, respectively. The oil sample was then determined as olive oil.

As shown in Table S-3, for the training set, a correct classification rate of 96.0% was obtained, indicating the effectiveness of the established PLS-DA models and approach for classification of edible oils. It was noticed that for the 8 edible oil samples that were misclassified, they were mainly the oil species with sample sizes equal to or less than 3 (e.g. almond oil and high oleic acid sunflower oil), indicating that further sample collection for these oil species might be needed to improve the model establishment and the classification. The misclassification of

366 grapeseed oil was due to the high similarity of the MALDI-MS spectra among grapeseed oil,
367 safflower oil and sunflower oil. As shown in Table 2, for the testing set, the correct
368 classification rate was 98.3%, with only 4 tested samples misclassified, indicating the
369 established models could predict external samples as good as the training samples. All the
370 misclassifications occurred in the grouping step, with 1 sample from group 4 misclassified as
371 group 5 and 2 samples from group 5 and 1 sample from group 8 misclassified as group 4. The
372 spectra of the misclassified samples were re-examined. It was found that although the mass
373 spectra of the two misclassified samples from group 5 (both were sunflower oils) were similar
374 to those of the others in the group, it showed slightly higher peaks at m/z 879.7 and m/z 905.8
375 than the others (see Figures S-7a and S-7b for the spectral comparison). As shown in the PLS-
376 DA loading plot (Figure S-7c), these two peaks were the important factors for classification of
377 samples in group 4, thus causing the misclassification. Similar explanation could be applied to
378 the other misclassified samples. Such re-inspecting the spectra of the problematic PLS-DA
379 results might help to reduce the chance of misclassification. Overall, including both training
380 set and testing set, 97.2% of the tested samples were classified correctly using PLS-DA and
381 only 12 out of the 435 tested samples were misclassified. For commonly used edible oils such
382 as olive oil, peanut oil and canola oil, correct classification rates of 100% were achieved. Such
383 classification could be considered as highly accurate with the high correct classification rate.

384 Most misclassifications were related to the species with small sample sizes, such as pine nut
385 oil and avocado oil in the training set. These could be improved by collecting and analyze more
386 samples for those oil species.

387

388 The predicted scores of the tested samples indicated how well the samples were matched with
389 the training samples. Predicted scores closer to 1 indicated the better fitting between the models
390 and the tested samples. The ranges, means and the standard deviations of the correctly
391 classified edible oil products are listed in Table S-4. The predicted scores of butter, coconut oil,
392 flaxseed oil and perilla oil which possessed characteristic peaks in their MALDI-MS spectra
393 were very close to 1, indicating the high agreement between the established models and the
394 tested samples. The predicted scores of most edible oil species were higher than 0.7 except for
395 grapeseed oil, camellia oil and hazelnut oil. The low predicted scores of some edible oil species
396 were normally due to the lack of characteristic spectral patterns of the species, increasing the
397 difficulty in classification of those species. The range of the predicted scores reflected the
398 variations of the mass spectral patterns of the edible oil species. As shown in Table S-4, peanut
399 oil, walnut oil and soybean oil showed larger variations among all the oil species.

400

Five collected edible oil samples (1 flaxseed oil, 2 sesame oils, 1 sunflower oil and 1 safflower oil), which failed to pass the confirmation using the GC-FID analysis as discussed previously (Figure S-9), were analyzed by the established PLS-DA models. As shown in Table S-5, all these 5 samples were not classified as the claimed edible oil species. These results were consistent with those obtained using the GC-FID analysis, demonstrating the reliability of the PLS-DA models.

3.4 Analysis of repeatedly cooked edible oils and gutter oils

Spectra of the repeatedly cooked edible oils and gutter oil samples are shown in Figure S-8. In these samples, peaks corresponding to oxidized products, such as peaks at m/z 915-955 (Picariello et al., 2009; Q. Zhang et al., 2016), were observed, and the TAGs contents in some of the gutter oil samples were relatively low. Gutter oils have diverse compositions and correspondingly there is no universal spectral pattern for their identification, but the differences in the TAGs contents and the presence of extra peaks of the oxidized products could provide clues for distinguishing gutter oils from normal edible oils. As gutter oils have to be labeled as normal edible oils in order to be sold in the market, identification of mislabeling is a reasonable strategy to screen out gutter oils. In this study, the spectra of the repeatedly cooked edible oils and gutter oil samples were examined and analyzed using the established PLS-DA models,

and the results are shown in Table S-6. Compared with the predicted scores obtained from the authentic edible oil samples (see Table S-4), some of the tested gutter oil samples such as s470 and s471 gave either too high or too low predicted scores for the classification, and some of the tested gutter oil samples such as s472 and s473, although classified normally, showed low TAGs content or high level of oxidized products in their spectra (see Figures S-8d and S-8e), which were obviously different from those of the authentic edible oil samples. Similar results were obtained for the repeatedly cooked edible oils, which showed additional oxidized products and TAGs patterns that no longer matched with soybean oil.

4. Conclusions

In this study, 30 types of edible oils with more than 900 oil samples were analyzed using MALDI-MS. Spectral patterns for various edible oils were obtained, and a comprehensive MALDI-MS spectral database was built for analysis of edible oils. Based on the spectral patterns and PCA results, edible oils were divided into 8 groups with characteristic peaks for each group. PLS-DA models were established for classification of edible oils, with overall 97.2% (96.0% for training set and 98.1% for testing set) of edible oil products correctly classified from 435 edible oil products and nearly 100% correct classification rates for commonly used edible oils. Counterfeit edible oils and gutter oils could also be differentiated

from normal edible oils using the models. Improved analysis could be achieved with incorporation of more edible oil products, particularly for those species with small sample sizes, into the database. The results of this study showed that MALDI-MS analysis combined with the established spectral database and statistical analysis could be an effective approach for rapid classification of edible oils.

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