

Statistical Evaluation of Triacylglycerol Composition in Plant Oils Based on High-Performance Liquid Chromatography—Atmospheric Pressure Chemical Ionization Mass Spectrometry Data

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The statistical evaluation of triacylglycerol profiles in plant oils based on high-performance liquid chromatography mass spectrometry (HPLC/MS) analysis enables the differentiation of various plant oils on the basis of the multidimensional data matrix. A data set of 93 oil samples from 60 varieties of plants composed from 355 triacylglycerols is evaluated using the principal component analysis. Analyzed samples are resolved in the principal component analysis plot, and similarities among some types of plant oils are visualized by the formation of clusters. The authentication of plant oils is tested with model samples of olive oil adulterated with sunflower oil at different concentration levels. Our HPLC/MS method using the statistical multivariate data analysis of a large data matrix enables a clear identification of adulterated olive oils already from 1% of added sunflower oil as an adulterant.

KEYWORDS: High-performance liquid chromatography; mass spectrometry; atmospheric pressure chemical ionization; triacylglycerol; plant oil; adulteration; authentication; statistics; principal component analysis

INTRODUCTION

Plant oils are an important commodity in world markets because of their widespread utilization in many branches of industry, cosmetics, and nutrition. They are produced from oil plants representing almost 10% of the world production of all crops according to the Food and Agriculture Organization of the United Nations (1). The annual production of edible plant oils has increased in the past decade by more than 50% to 127 million tonnes a year (1) and is still increasing annually. Edible plant oils are mixtures of lipids composed mainly from triacylglycerols (TGs) with the content up to 95%. They serve as an important source of fatty acids in the human diet, mainly essential fatty acids necessary for the biosynthesis of long-chain polyunsaturated fatty acids important for the synthesis of cell membranes in the human body. A diet with 70 g of fat per day for female adults and 90 g for male adults corresponding to 30–35% of daily energy coming from fats is now considered as consistent with good health (2). In reality, the consumption of oils and fats in USA and EU is about 130 g per day per person (3).

Prices of plant oils are given by many parameters, mainly by the production cost and the quality of plant oils. Higher prices of high-quality plant oils can lead to the effort of falsification by cheaper oils with a lower quality and less beneficial nutritional properties (e.g., expensive virgin olive oil adulterated by cheaper

sunflower oil); therefore, their authentication is of great interest nowadays. Many authentication methods use the measurement of oil fingerprints without any separation and sample pretreatment steps, e.g., Raman spectroscopy (4, 5), infrared spectroscopy (6, 7), nuclear magnetic resonance spectroscopy (8, 9), matrix-assisted laser desorption/ionization mass spectrometry (MS) (10, 11), electrospray ionization (ESI) MS (12, 13), atmospheric pressure photoionization MS (13), and so forth. Although the fingerprint methods are fast and simple, some plant oils have similar fingerprints differing only in low concentration components not detectable this way. TGs are compounds suitable for the authentication of plant oils because they are the main components of plant oils with several tens of different species occurring at different concentration levels. They are characterized by fatty acids esterified on the glycerol skeleton and their properties, i.e., carbon number (CN), double bond (DB) number, the configuration and position of DBs in acyl chains, and the stereochemical position of fatty acids on the glycerol skeleton. TG profiles differ for each type of plant oil which is used for authentication based on chromatographic separation, i.e., gas chromatography/isotopic ratio mass spectrometry (14), gas chromatography/flame ionization detection (GC/FID) (15, 16), high-performance liquid chromatography (HPLC)/refractive-index detection (17), HPLC/atmospheric pressure chemical ionization (APCI) MS (10, 18–20), and off-line two-dimensional HPLC/MS (21).

The highest number of identified TGs in plant oils have been reported using nonaqueous reversed-phase (NARP) HPLC with APCI-MS detection (22, 23). In NARP-HPLC mode, TGs are separated according to the equivalent carbon number (ECN)

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Table 1. Number of Identified Triacylglycerols (TGs) and Fatty Acids (FAs), Average Equivalent Carbon Number (aECN), Average Carbon Number (aCN), Average Double Bond (aDB) Number, the Relative Weight Concentration [%] of Essential Fatty Acids (Linoleic and Linolenic Acids), Fatty Acids with 18 (C18) and 16 (C16) Carbon Atoms, and Saturated (Sat), Monounsaturated (Mono), and Polyunsaturated (Poly) Fatty Acids in Analyzed Plant Oils Calculated from NARP-HPLC/APCI-MS of Triacylglycerols

| oil | no. | number of TGs/FAs | aECN | aCN | aDB | essential FAs [%] | C18 + C16 FAs [%] | Sat [%] | Mono [%] | Poly [%] |
|------------------|-----|-------------------|-------|-------|------|-------------------|-------------------|---------|----------|----------|
| Kiwi | 1 | 47/11 | 13.59 | 17.83 | 2.12 | 70.7 | 99.4 | 11.7 | 17.3 | 71.0 |
| Macadamia nut | 2 | 45/13 | 15.82 | 17.53 | 0.85 | 2.9 | 91.0 | 17.5 | 79.6 | 2.9 |
| Hemp | 3 | 70/18 | 14.03 | 17.87 | 1.89 | 70.7 | 98.0 | 12.9 | 11.8 | 75.3 |
| Brazil nut | 4 | 26/7 | 15.38 | 17.66 | 1.14 | 39.9 | 99.9 | 24.9 | 35.2 | 39.9 |
| Mango | 6 | 53/13 | 16.57 | 17.86 | 0.65 | 8.2 | 96.5 | 44.2 | 47.6 | 8.2 |
| Dog rose | 7 | 51/14 | 14.27 | 17.93 | 1.83 | 69.7 | 98.4 | 8.2 | 22.0 | 69.8 |
| Hazelnut | 9 | 30/10 | 15.62 | 17.83 | 1.10 | 21.3 | 99.2 | 11.1 | 67.6 | 21.3 |
| | 10 | 30/10 | 15.72 | 17.84 | 1.06 | 17.6 | 99.1 | 11.4 | 71.0 | 17.6 |
| Sweet chestnut | 11 | 49/16 | 15.04 | 17.72 | 1.34 | 44.2 | 98.7 | 16.4 | 39.3 | 44.3 |
| Pumpkin | 12 | 31/9 | 15.01 | 17.71 | 1.35 | 56.0 | 99.0 | 19.9 | 24.1 | 56.0 |
| Lemon | 13 | 58/12 | 14.90 | 17.62 | 1.36 | 46.9 | 99.3 | 23.0 | 30.1 | 46.9 |
| Bell pepper | 14 | 44/16 | 14.61 | 17.77 | 1.58 | 74.5 | 98.4 | 15.9 | 9.6 | 74.6 |
| Grapefruit | 15 | 51/14 | 15.07 | 17.46 | 1.19 | 44.7 | 99.2 | 29.9 | 25.5 | 44.7 |
| Cucumber | 16 | 45/13 | 14.54 | 17.62 | 1.54 | 72.1 | 99.1 | 20.0 | 7.9 | 72.1 |
| Blackcurrant | 18 | 80/14 | 13.76 | 17.85 | 2.05 | 57.5 | 99.3 | 8.9 | 15.2 | 75.9 |
| Mandarin orange | 19 | 56/14 | 15.03 | 17.55 | 1.26 | 48.1 | 98.8 | 26.4 | 25.5 | 48.1 |
| Blueberry | 20 | 37/9 | 13.94 | 17.86 | 1.96 | 70.0 | 99.7 | 7.9 | 22.0 | 70.0 |
| Melon cantaloupe | 21 | 37/11 | 14.70 | 17.76 | 1.53 | 68.6 | 99.5 | 15.6 | 15.8 | 68.6 |
| Papaya | 22 | 55/17 | 15.93 | 17.66 | 0.87 | 9.4 | 97.9 | 22.4 | 68.2 | 9.4 |
| Buckwheat | 23 | 59/16 | 15.51 | 17.92 | 1.20 | 39.4 | 92.0 | 21.5 | 39.0 | 39.5 |
| Pistachio | 24 | 40/11 | 15.28 | 17.81 | 1.27 | 38.5 | 99.1 | 11.5 | 50.0 | 38.5 |
| Peanut | 26 | 60/16 | 15.60 | 17.94 | 1.17 | 37.9 | 92.9 | 21.2 | 40.8 | 37.9 |
| Camellia | 27 | 26/12 | 15.87 | 17.80 | 0.97 | 9.5 | 99.1 | 12.5 | 78.0 | 9.5 |
| | 28 | 26/12 | 15.88 | 17.82 | 0.97 | 8.8 | 99.2 | 11.6 | 79.6 | 8.8 |
| Rice | 29 | 48/12 | 15.30 | 17.66 | 1.18 | 38.1 | 97.9 | 20.6 | 41.3 | 38.1 |
| Coffee butter | 30 | 68/14 | 14.05 | 15.01 | 0.48 | 20.5 | 57.6 | 72.3 | 7.2 | 20.5 |
| Apricot kernel | 31 | 27/10 | 15.40 | 17.86 | 1.23 | 31.1 | 99.8 | 7.3 | 61.7 | 31.1 |
| Raspberry | 32 | 51/13 | 13.91 | 17.94 | 2.02 | 79.7 | 99.4 | 4.8 | 15.5 | 79.7 |
| Argan | 33 | 60/16 | 15.45 | 17.71 | 1.13 | 33.3 | 98.8 | 19.5 | 47.2 | 33.3 |
| Black cumin | 34 | 35/9 | 14.89 | 17.80 | 1.45 | 58.8 | 96.6 | 15.0 | 23.5 | 61.5 |
| Moringa | 35 | 33/16 | 16.65 | 18.24 | 0.80 | 0.9 | 86.4 | 22.1 | 77.0 | 0.9 |
| Tamanu | 36 | 43/12 | 15.40 | 17.79 | 1.19 | 41.7 | 98.6 | 21.7 | 36.6 | 41.7 |
| Soya | 38 | 66/14 | 14.86 | 17.79 | 1.47 | 56.9 | 98.5 | 16.8 | 26.3 | 56.9 |
| Rapeseed | 41 | 55/13 | 15.29 | 17.90 | 1.31 | 30.8 | 97.9 | 9.6 | 59.6 | 30.8 |
| Sunflower | 44 | 50/16 | 14.91 | 17.88 | 1.49 | 61.9 | 97.9 | 13.3 | 24.8 | 61.9 |
| Olive | 53 | 37/15 | 15.90 | 17.75 | 0.92 | 7.5 | 98.5 | 15.8 | 76.7 | 7.5 |

from 17.46 to 18.24, aDB from 0.48 to 2.12, and the sum of C18 and C16 fatty acids from 86.4% to 99.8%, showing that plant oils are composed almost exclusively from TGs containing fatty acids with 16 and 18 carbon atoms and 0 to 4 DBs, i.e., palmitic (ECN; CN; DB-16; 16; 0), stearic (18; 18; 0), oleic (16; 18; 1), linoleic (14; 18; 2), and linolenic (12; 18; 3) acids (Tables 2 and S3 (Supporting Information)). Remaining fatty acids with low or usually trace concentrations represent long or short-chain acids, odd-number acids, and acids with unusual DB positions. Higher differences are found among the sums of essential (from 0.9% to 79.7%), saturated (from 4.8% to 72.3%), monounsaturated (from 7.2% to 79.6%), and polyunsaturated (from 0.9% to 75.9%) fatty acids differing significantly for individual oils, and therefore, these parameters can be used for fast consideration of nutritional values or possible industrial applications.

PCA of TG Composition. The evaluation of TG profiles is an important step in the quality control of plant oils. The concentration of individual TG species can be used for simple comparison of various plant oils, but such comparison is not practical due to a high number of detected TGs. For detailed characterization, the comparison of all TG species in all analyzed samples is necessary, which leads to the complex multidimensional data set. Multivariate data analysis using PCA is used for the evaluation of TG composition in all analyzed samples. First, PCA analysis using TG concentrations based on APCI-MS response

approach and TG relative peak areas are compared. No significant differences in resulting PCA plots are found, and therefore, relative peak areas are used for further PCA analysis of all samples. The final data set contains 93 plant oils (i.e., objects) of 60 different types characterized by relative peak areas of 355 identified TG species (i.e., variables). Thirteen variables are excluded from the data set because of their zero variability corresponding to the content of this variable in all plant oils lower than the limit of detection (0.01%). Data values of other 342 variables range between 0.01% and 49.32%, i.e., in the range of 3.5 orders of magnitude. Hence, no scaling, normalization, or centering is applied, and the data set without any modification is taken for the direct PCA analysis. Multivariate data set of 342 nonredundant variables is visualized as a set of coordinates in a multidimensional data space with $N = 342$ (one axis per variable) dimensions.

Figure 5 shows the score plots of the first (t[1]) and second (t[2]) PCs of the general PCA model with a good resolution of analyzed samples. These two variables describe 82% of the total variability in the data set, where the first PC t[1] describes 52% and second PC t[2] 30% of the total variability. Other PCs describe significantly lower variability, e.g., t[3] has 4% and t[4] 3% of the total variability. The projection of PCs t[3] and t[4] (Figure S6 (Supporting Information)) shows only a small variance among analyzed samples, and most of samples are grouped around the

Table 2. Relative Weight Concentrations [%] of Individual Fatty Acids in Analyzed Plant Oils Calculated from HPLC/APCI-MS of Triacylglycerols

| oil | no. | Cy | C | La | M | M | P | Po | Ma | Mo | S | O | L | Ln | γ Ln | St | A | G | B | Lg | C26:0 | | | | | | | | | |
|------------------|-----|------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|------|
| | | C8:0 | C10:0 | C12:0 | C14:0 | C15:0 | C16:0 | C16:1 | C17:0 | C17:1 | C18:0 | C18:1 | C18:2 | C18:3 | C18:3 | C18:4 | C19:0 | C20:0 | C20:1 | C20:2 | C21:0 | C22:0 | C22:1 | C23:0 | C24:0 | C24:1 | C25:0 | C26:0 | | |
| Kiwi | 1 | | | | | | 8.5 | 0.01 | | | 3.1 | 17.1 | 17.1 | 53.6 | | | 0.04 | 0.2 | 0.3 | 0.02 | | | | | | | | | | |
| Macadamia nut | 2 | | | 0.2 | 1.4 | | 8.0 | | | 0.1 | 3.3 | 57.9 | 2.9 | | | | 2.8 | 2.6 | | 0.02 | | 1.3 | 0.1 | | 0.5 | | | | | |
| Hemp | 3 | | | | | | 8.6 | | | 0.02 | 2.6 | 11.5 | 49.5 | 21.2 | 3.2 | 1.4 | 1.0 | 0.3 | 0.03 | | <0.01 | 0.5 | 0.01 | 0.03 | 0.1 | 0.01 | | <0.01 | | |
| Brazil nut | 4 | | | | | | 16.3 | 0.08 | | | 8.5 | 35.1 | 39.9 | | | | 0.1 | 0.02 | | | | | | | | | | | | |
| Mango | 6 | | | | | | 10.7 | | | | 30.3 | 47.3 | 7.4 | 0.8 | | 0.03 | 1.7 | 0.3 | 0.07 | 0.5 | 0.01 | 0.1 | 0.04 | 0.6 | 0.6 | 0.03 | | | | |
| Dog rose | 7 | | | | | | 4.6 | | | 0.04 | 2.4 | 21.7 | 47.9 | 21.8 | | | 1.0 | 0.3 | 0.03 | | | 0.1 | 0.03 | 0.04 | | | | | | |
| Hazelnut | 9 | | | | | | 7.8 | 0.6 | | 0.1 | 0.2 | 3.0 | 66.5 | 20.8 | 0.5 | | 0.2 | 0.3 | | | | | | | | | | | | |
| | 10 | | | | | | 7.5 | 0.3 | | 0.1 | 0.2 | 3.5 | 70.2 | 17.3 | 0.3 | | 0.3 | 0.3 | | | | | | | | | | | | |
| Sweet chestnut | 11 | | | | | 0.04 | 14.3 | 0.08 | | 0.1 | 0.04 | 1.5 | 38.6 | 37.1 | 7.1 | | 0.3 | 0.6 | 0.06 | 0.01 | 0.1 | 0.1 | 0.04 | 0.03 | | | | | | |
| Pumpkin | 12 | | | | | | 14.5 | | | 0.1 | 4.5 | 24.0 | 56.0 | | | | 0.6 | 0.06 | | | | 0.2 | 0.04 | | | | | | | |
| Lemon | 13 | | | | | | 18.8 | | | 0.08 | 3.5 | 30.1 | 33.4 | 13.5 | | | 0.3 | 0.03 | | | 0.08 | 0.2 | <0.01 | 0.2 | | | 0.01 | 0.01 | | |
| Bell pepper | 14 | | | | | 0.06 | 12.5 | | | 0.1 | 0.2 | 2.1 | 9.3 | 73.7 | 0.8 | | 0.4 | 0.07 | 0.08 | 0.01 | 0.3 | 0.3 | 0.03 | 0.3 | | | 0.05 | 0.05 | | |
| Grapefruit | 15 | | | | | | 25.9 | 0.8 | | 0.2 | 3.2 | 24.6 | 38.6 | 6.1 | | | 0.3 | 0.05 | | | 0.02 | 0.02 | 0.01 | 0.2 | | | 0.01 | 0.01 | | |
| Cucumber | 16 | | | | 0.4 | | 17.5 | | | 0.1 | 1.7 | 7.8 | 68.3 | 3.8 | | | 0.2 | 0.09 | | | 0.04 | 0.04 | 0.01 | 0.05 | | | 0.01 | 0.01 | | |
| Blackcurrant | 18 | | | | | | 7.7 | | | 0.02 | 1.1 | 14.7 | 40.9 | 16.6 | 14.6 | 3.7 | 0.1 | 0.5 | 0.07 | | 0.01 | 0.1 | <0.01 | <0.01 | | | 0.01 | 0.01 | | |
| Mandarin orange | 19 | | | | | | 22.1 | 0.4 | | 0.2 | 3.3 | 24.9 | 42.6 | 5.5 | | | 0.3 | 0.2 | 0.02 | | 0.1 | 0.1 | 0.02 | 0.3 | | | 0.03 | 0.05 | | |
| Blueberry | 20 | | | | | | 6.6 | | | 0.02 | 1.1 | 22.0 | 35.7 | 34.3 | | | 0.2 | 0.06 | | | | 0.1 | 0.02 | | | | | | | |
| Melon cantaloupe | 21 | | | | | 0.05 | 11.8 | | | 0.08 | 3.4 | 15.7 | 67.8 | 0.8 | | | 0.2 | 0.09 | | | 0.06 | 0.2 | 0.02 | 0.02 | 0.02 | | 0.01 | 0.01 | 0.01 | |
| Papaya | 22 | | | | 0.2 | | 16.6 | 0.9 | | 0.2 | 4.5 | 66.5 | 8.9 | 0.5 | | | 0.6 | 0.7 | | | 0.02 | 0.2 | 0.03 | 0.01 | 0.07 | | 0.01 | 0.01 | 0.01 | |
| Buckwheat | 23 | | | | | | 14.7 | 0.2 | | 0.06 | 2.0 | 35.7 | 36.9 | 2.5 | | | 1.1 | 3.0 | 0.08 | 0.04 | 2.0 | 0.1 | 0.2 | 1.4 | | | 0.02 | 0.02 | 0.02 | |
| Pistachio | 24 | | | | | | 9.7 | | | 0.05 | 1.5 | 49.4 | 38.2 | 0.3 | | | 0.1 | 0.5 | | | 0.1 | 0.1 | 0.1 | 0.5 | | | 0.05 | 0.05 | 0.1 | |
| Peanut | 26 | | | | | | 12.6 | | | 0.09 | 0.08 | 2.7 | 39.7 | 37.6 | 0.3 | | 1.1 | 1.0 | 0.04 | 0.01 | 3.1 | 0.05 | 0.03 | 0.07 | | | 0.03 | 0.03 | 0.03 | |
| Camellia | 27 | | | | | | 10.0 | 0.08 | | 0.08 | 2.3 | 77.2 | 9.2 | 0.3 | | | 0.06 | 0.6 | | | | 0.03 | 0.03 | 0.03 | | | 0.03 | 0.03 | 0.03 | |
| | 28 | | | | | | 9.1 | 0.07 | | 0.06 | 2.4 | 78.8 | 8.5 | 0.3 | | | 0.04 | 0.6 | | | | 0.2 | 0.03 | 0.03 | | | 0.03 | 0.03 | 0.03 | |
| Rice | 29 | | | | | | 16.5 | 0.7 | | | 2.5 | 40.1 | 36.1 | 2.0 | | | 0.5 | 0.5 | | | | 0.2 | 0.2 | 0.4 | | | 0.1 | 0.1 | 0.1 | |
| Coffee butter | 30 | 5.1 | 1.5 | 24.4 | 10.1 | | 17.6 | | | 0.01 | 12.3 | 7.2 | 19.7 | 0.8 | | | 0.9 | 0.03 | | | 0.3 | 0.3 | 0.06 | 0.06 | | | 0.02 | 0.02 | 0.02 | |
| Apricot kernel | 31 | | | | | | 6.2 | 0.3 | | 0.03 | 1.0 | 61.2 | 31.0 | 0.05 | | | 0.06 | 0.06 | | | 0.01 | 0.2 | 0.01 | 0.01 | 0.02 | | | <0.01 | <0.01 | |
| Raspberry | 32 | | | | | | 3.4 | | | 0.02 | 0.8 | 15.5 | 52.9 | 26.8 | | | 0.3 | 0.04 | | | | 0.2 | 0.01 | 0.01 | 0.07 | | | 0.01 | 0.01 | 0.01 |
| Argan | 33 | | | | | | 14.0 | 0.4 | | 0.05 | 4.7 | 46.4 | 33.1 | 0.2 | | | 0.3 | 0.4 | | | 0.01 | 0.2 | 0.01 | 0.01 | 0.07 | | | 0.01 | 0.01 | |
| Black cumin | 34 | | | | | | 12.4 | | | | 2.3 | 23.1 | 58.8 | | | | 0.2 | 0.4 | | | | 0.06 | 0.06 | 0.04 | | | | | | |
| Moringa | 35 | | | | | | 6.4 | 1.2 | | 0.05 | 4.7 | 73.2 | 0.9 | | | 0.02 | 3.4 | 2.5 | | | 0.06 | 5.8 | 0.07 | 0.05 | 1.6 | | | 0.01 | 0.01 | 0.01 |
| Tamanu | 36 | | | | | | 11.2 | 0.03 | | 0.08 | 0.03 | 9.3 | 36.4 | 41.7 | | | 0.8 | 0.1 | | | 0.02 | 0.3 | 0.04 | 0.04 | | | | | | |
| Soya | 38 | | | | | | 12.1 | | | 0.04 | 0.03 | 3.3 | 26.2 | 49.7 | 7.2 | | 0.4 | 0.1 | | | 0.6 | 0.6 | 0.1 | 0.2 | | | 0.01 | 0.01 | 0.01 | |
| Rapeseed | 41 | | | | | | 7.1 | 0.2 | | 0.1 | 1.6 | 58.2 | 21.2 | 9.6 | | | 0.4 | 1.1 | | | 0.3 | 0.3 | 0.01 | 0.1 | 0.09 | | | 0.01 | 0.01 | 0.01 |
| Sunflower | 44 | | | | | | 7.6 | | | 0.05 | 0.08 | 3.8 | 24.6 | 61.5 | 0.4 | | <0.01 | 0.4 | 0.1 | | 0.03 | 0.8 | 0.06 | 0.3 | | | 0.01 | 0.01 | 0.01 | |
| Olive | 53 | | | | | | 12.5 | 1.3 | | 0.1 | 2.3 | 74.9 | 6.5 | 1.0 | | | 0.5 | 0.4 | | | 0.03 | 0.3 | 0.05 | 0.1 | | | 0.01 | 0.01 | 0.01 | |

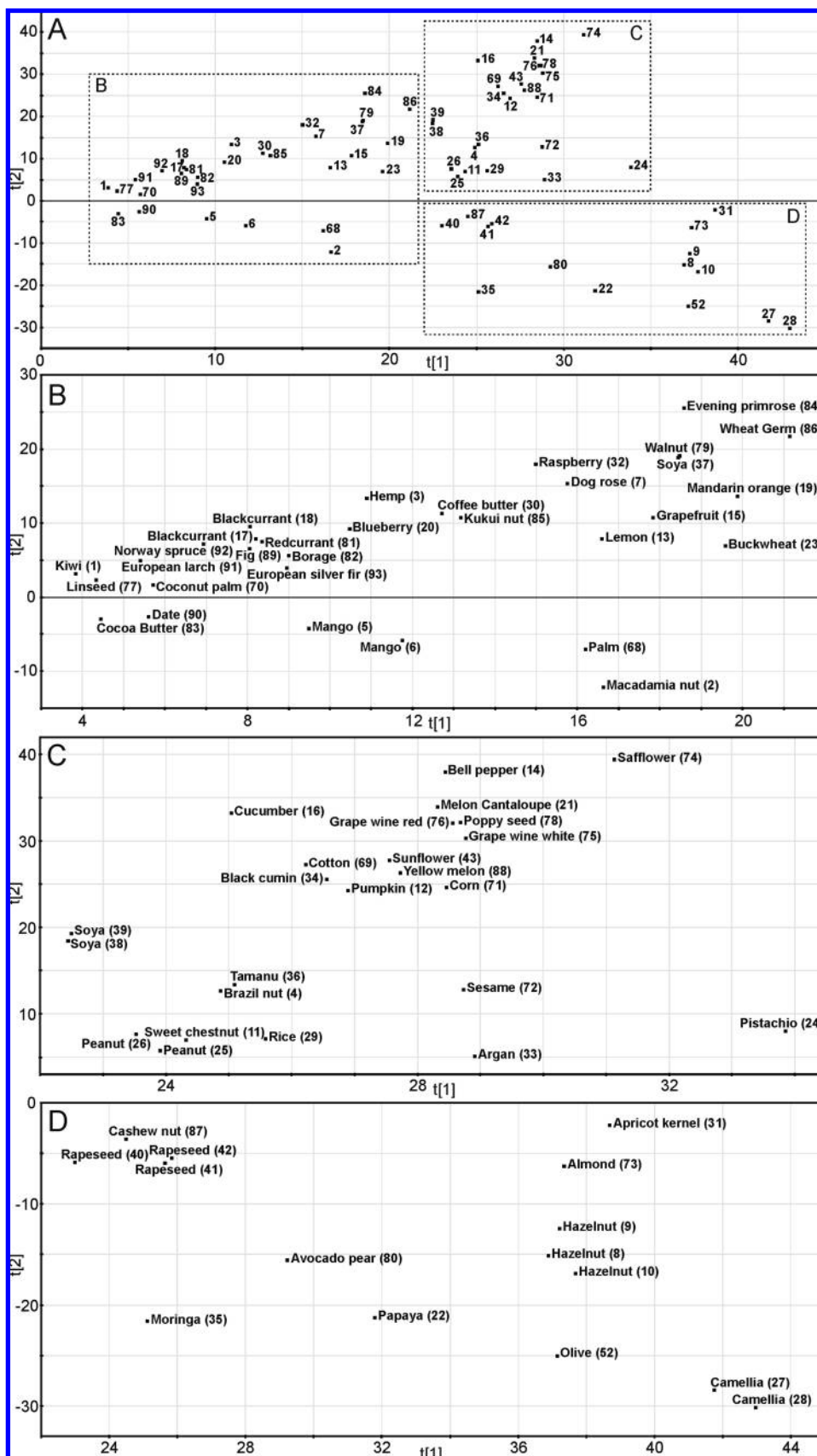


Figure 5. Projection of principal components $t[1]$ and $t[2]$ in two-dimensional scatter plot for all measured samples (A) and zooms of individual regions (B, C, and D).

zero of both PCs. Only samples with significantly different composition containing high concentrations of TGs with highly unsaturated (linseed, kiwi and blueberry oils) or saturated (cacao

butter and mango oils) fatty acids are clearly distinguished from other samples in Figure S6 (Supporting Information). The projection of analyzed samples using $t[1]$ and $t[2]$ PCs provides

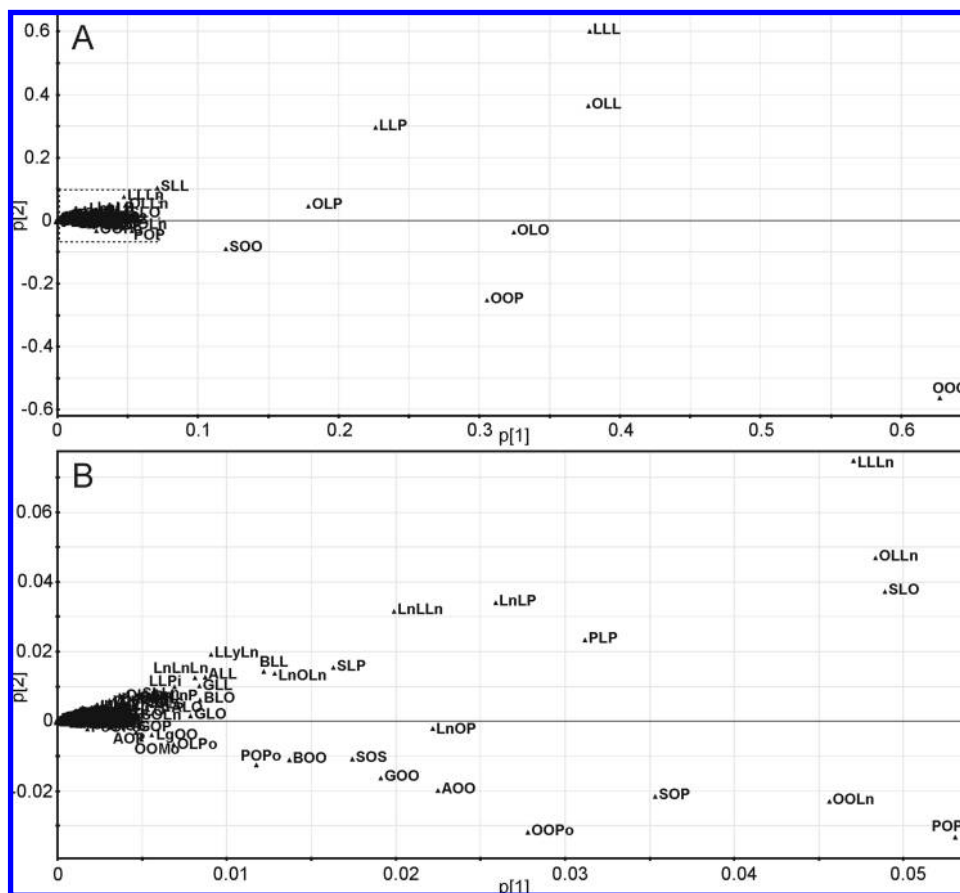


Figure 6. Projection of variables $p[1]$ and $p[2]$ in two-dimensional loadings plot for all measured samples showing the major variables representing TG concentrations (A) and zoomed area (B).

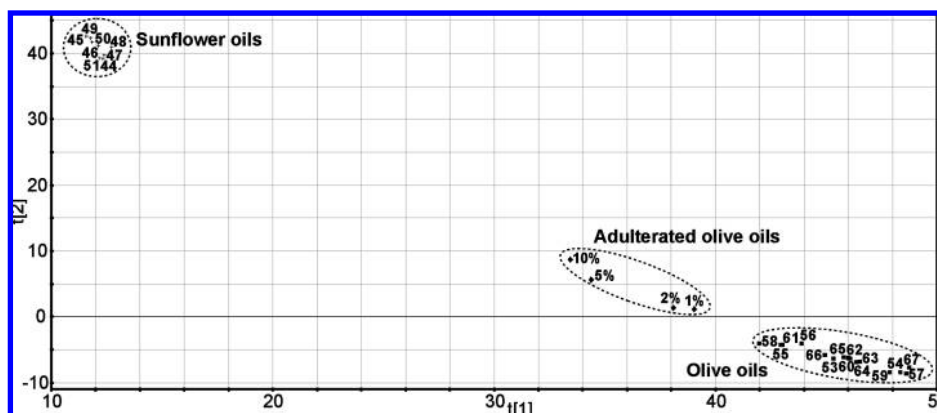


Figure 7. Projection of principal components $t[1]$ and $t[2]$ in two-dimensional scatter plot for analyzed sunflower (44–51) and olive (53–67) cooking oils and four samples of adulterated olive oil by 1%, 2%, 5%, or 10% of sunflower oil.

significantly better resolution of analyzed samples for the comparison of their properties. Samples of one type of plant oil having similar TG composition form narrow clusters, e.g., different samples of hazelnut and camellia oils (Figure 5D). Samples with similar TG profiles are grouped in small regions in the PCA plot, e.g., samples of blackcurrant and redcurrant oils (Figure 5B). Similar positions of various samples in the PCA plot indicate their similar properties, e.g., Brazil nut and tamanu oils in Figure 5C. Their similar properties can be confirmed by the comparison of their average parameters and sums of fatty acids for individual plant oils calculated from TG composition (Table 1), i.e., aECN of Brazil nut oil/tamanu oil = 15.38/15.40, aCN = 17.66/17.79, aDB = 1.14/1.19, C18 + C16 fatty acids = 99.9%/98.6%,

polyunsaturated fatty acids = 39.9%/41.7%, etc. Figure 6 shows variables (TG concentrations) in our PCA, and their variance model mostly affects the variability of samples. The most significant variable is the concentration of OOO with more than 60% positive effect on $t[1]$, while –55% effect on $t[2]$. Other important variables are LLL, OLL, OLO, OOP, LLP, OLP, and SOO. These eight variables are the most significant parameters for the statistical differentiation among plant oils.

Authentication of Olive Oil. Olive oil is one of the most expensive plant oils used in dietetics. For its healthy properties, it is an important ingredient in the so-called Mediterranean diet of southern nations. High prices of olive oils can lead some merchants to illegal falsification by cheaper plant oils, which decrease

their nutritional value. Most favorable oils for adulteration are plant oils with similar TG composition, which are difficult to distinguish using common analytical techniques. The TG composition of hazelnut, camellia, or papaya oils is relatively close to olive oil composition, but they are still clearly distinguished using our HPLC/MS method and PCA analysis (Figure 5D). Moreover, their prices and small quantity production in comparison to those of the most common plant oils are not favorable for falsification. The utilization of low-price plant oils produced in large quantities in the same geographical region is more favorable, e.g., sunflower oil. The set of 8 sunflower and 15 olive cooking oils, and 4 model samples of adulterated olive oil by 1%, 2%, 5%, or 10% of sunflower oil (Table S2, Supporting Information) is tested to develop an unambiguous method to identify adulteration even at very low amounts of adulterant. Figure 7 shows the scores plot of the first (t[1]) and second (t[2]) PCs of all cooking and model adulterated oil samples. This data set is represented by 27 objects (oil samples) and 62 variables (TG concentrations) with significant variability. PCs t[1] and t[2] account for 99.6% of total variability, where t[1] represents 73.5% and t[2] represents 26.1% of total variability. Samples of sunflower oil have small differences in TG composition and form a small cluster clearly distinguished from other samples in the PCA plot. Samples of olive oil have a wider distribution in comparison to the cluster of sunflower oils because of slightly different TG composition of different types (virgin oil, pomace oil, etc.) and different origin of samples, which are not differentiated in this study. Anyway, a clear resolution of sunflower and olive oil samples and their grouping into small clusters enable the resolution of model samples of adulterated olive oil by sunflower oil (Figure 7). Samples of adulterated olive oil with increasing concentrations of sunflower oil have an increasing distance from the olive oil cluster in the PCA plot. Even the adulteration of olive oil by 1% of sunflower oil can be clearly visualized in a PCA plot regardless of the fact that different types and origin of olive oils are neglected. This approach is well suitable for the detection of possible adulteration in tested samples.

The presented results demonstrate the utilization of HPLC/MS analysis and statistical evaluation in the quality control of plant oils. A carefully optimized HPLC/MS method is used for detailed characterization of TG profiles of plant oils. PCA evaluation of multidimensional data matrix of TG profiles enables the comparison of all analyzed samples and the resolution of samples with similar properties. PCA analysis is used for the authentication of expensive olive oil. PCA enables the detection of adulterated olive oil starting from 1% of added sunflower oil as an adulterant.

ABBREVIATIONS USED

CN, carbon number; DB, double bond; ECN, equivalent carbon number; ESI, electrospray ionization; MS, mass spectrometry; NARP-HPLC, nonaqueous reversed-phase high-performance liquid chromatography; PC, principal component; PCA, principal component analysis; TG, triacylglycerol; fatty acid abbreviations, Cy, caprylic (CN:DB, C8:0); C, capric (C10:0); La, lauric (C12:0); M, myristic (C14:0); C15:0, pentadecanoic; P, palmitic (C16:0); Po, palmitoleic (Δ^9 -C16:1); Ma, margaric (C17:0); Mo, margaroleic (Δ^9 -C17:1); S, stearic (C18:0); O, oleic (Δ^9 -C18:1); L, linoleic ($\Delta^9,12$ -C18:2); Ln, α -linolenic ($\Delta^9,12,15$ -C18:3); γ Ln, γ -linolenic ($\Delta^6,9,12$ -C18:3); St, stearidonic ($\Delta^6,9,12,15$ -C18:4); C19:0, nonadecanoic (C19:0); A, arachidic (C20:0); G, gadoleic (Δ^9 -C20:1); C20:2, eicosadienoic ($\Delta^{11,14}$ -C20:2); C21:0, heneicosanoic (C21:0); B, behenic (C22:0); C22:1, erucic (Δ^{13} -C22:1); C23:0, tricosanoic (C23:0); 24:1, nervonic (Δ^{15} -C24:1); Lg, lignoceric (C24:0); C25:0, pentacosanoic (C25:0); C26:0, hexacosanoic (C26:0).

Supporting Information Available: HPLC/MS chromatograms of analyzed plant oils (Figures S1–S5), projection of PCs t[3] and t[4] (Figure S6), relative weight concentrations of triacylglycerols (Tables S1 and S2) and fatty acids (Tables S3), average parameters (Table S4) of analyzed plant oils. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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