

Analytical approaches for discriminating native lard from other animal fats

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Abstract

Establishing the distinguishable characteristics of lard from other common animal fats might be helpful for authentication initiatives in foods and feeds. In this study, fatty acid and triacylglycerol compositions, thermal and spectroscopic characteristics of native lard (NL), respectively, were compared with those of beef tallow (BT), mutton tallow (MT), and chicken fat (CF) by using gas liquid chromatography (GLC), high-performance liquid chromatography (HPLC), and differential scanning calorimetry (DSC). GLC analysis showed that the comparison of the overall fatty acid data might not be suitable for the discrimination of different animal fats, but the use of the principal component analysis and the percent palmitic acid enrichment factor [PAEF (%)] calculations were useful. HPLC analysis showed that NL displayed a TAG profile, which was quite different from those of either BT or MT, but appeared to be closely similar to that of CF. Results of DSC thermal analysis showed that both melting and crystallization curves of NL were remarkably different from those of other animal fats.

Keywords: animal fats; fatty acid composition; food authentication; thermal analysis; triacylglycerol composition

Introduction

Meat has been the main source of protein for humankind from time immemorial. Beef, pork, chicken, and mutton have been the preferred sources of meat for mankind. According to the worldwide meat intake assessment, pork is the most widely consumed meat accounting for over 36%, followed by poultry (35%) and beef (22%) (FAO, 2012). The consumption patterns of different animal species across the world might vary according to religious beliefs and cultural preferences. For instance, consumption of pork is not desirable for some segments of the society due to their religious restrictions. Either deliberate or accidental mixing of pork with other meat species is a great concern for adherents of Islam and Orthodox Judaism. According to some reports, mixing of pork with other commonly consumed meat types is reported

to have occurred due to fraudulent reasons (Saeed *et al.*, 1989). Hence, a great deal of effort has been made to develop analytical methodologies to detect fraudulent practices related to the presence of pork in food systems. Researchers in the past frequently employed deoxyribonucleic-acid-based methodologies to differentiate between different meat species in meat products (Aida *et al.*, 2005). Lipid-based detection methods, on the other hand, are advantageous since lipid component in food matrices is minimally affected by the processing conditions (Che Man *et al.*, 2005; Marikkar *et al.*, 2005; Sawaya *et al.*, 1990b). As considerable amount of literature has been available on the detection of animal fats in food systems (Che Man *et al.*, 2011; Marikkar *et al.*, 2005; Nurjuliana *et al.*, 2011; Nina Naqiyah *et al.*, 2017; Rohman and Che Man, 2010; Rohman *et al.*, 2011), it would be highly beneficial to discuss the distinguishable

characteristics of native lard (NL) from other common animal fats. Hence, the purpose of this article is to explore the literature and experimental studies on NL and other animal fats in terms of compositional and physico-chemical characteristics using different analytical techniques, such as gas liquid chromatography (GLC), high performance liquid chromatography (HPLC), and differential scanning calorimetry (DSC).

Materials and Methods

Sampling and reagents

Samples of animal fats, namely, NL, beef tallow (BT), mutton tallow (MT), and chicken fat (CF) were extracted by rendering adipose tissues of animals collected in triplicate from three slaughter houses located in different locations of Sri Serdang, Malaysia. Thirty-six samples were collected and analyzed in this study. All chemicals used in this experiment were of either analytical or HPLC grade.

Fat extraction protocol

The rendered fat was squeezed out using double-folded muslin cloth. The removal of impurities was carried out by filtering the melted animal fats through cotton wool. A small proportion of anhydrous sodium sulfate was added to remove residual moisture, and the samples were then filtered through Whatman No. 2 filter papers (Marikkar *et al.*, 2001).

Analytical methodologies

Preparation of fatty acid methyl esters

The determination of fatty acid composition required a derivatization step with methyl ester for all the animal fats using sodium methoxide. Preparation of fatty acid methyl esters of animal fat was done according to the Association of Official Agricultural Chemists (AOAC) method 969.33. A 50-mg portion of animal fat was weighed into a 20-mL test tube (with screw cap). After adding 2 mL of 2N sodium hydroxide in methanol, the sample tube was closed and heated at 80°C for 1 h. After allowing the tube to cool down for a few minutes, 2 mL of 25% borontrifluoride solution in methanol was added. The tube was closed and heated again for 1 h at 80°C. Subsequently, 5 mL portions of water and hexane were added into this. The contents of the tube were shaken well and allowed to undergo phase separation. The clear supernatant of the solution was transferred into a 2-mL auto-sampler vial (AOAC, 2007).

Determination of fatty acid composition

The top hexane layer of the FAME solution was injected on an Agilent 6890N gas chromatograph (Agilent Technologies, Singapore) equipped with a polar capillary column RTX-5 (0.32 mm internal diameter, 30 m length, and 0.25 mm film thickness; Restex Corp., Bellefonte, PA) and a Flame Ionization Detector (FID). Split injection was conducted with a split ratio of 58:1 using nitrogen as a carrier gas at a flow-rate of 1.00 mL/min. The temperature of the column was 50°C (for 1 min), and was programmed to increase to 200°C at 8°C/min. The temperatures of the injector and detector were maintained at 200°C (Nina Naqiyah *et al.*, 2013).

HPLC analysis of TAG composition

The separation of triacylglyceride (TAG) components of animal fats was performed on a Waters Model 510 liquid chromatograph equipped with a Merck Lichrosphere RP-18 column (5 µm) (12.5 cm × 4 mm i.d.; Merck, Darmstadt, Germany) and a differential refractometer Model 410 as a detector (Waters Associates, Milford, MA). Each fat sample was loaded in the sample vials after dissolving in chloroform and filtering through a Whatman no. 40 filter paper. The isocratic mobile phase was a mixture of acetone:acetonitrile (63.5:36.5 v/v) and the flow rate was set at 1.5 mL/min. Each sample was chromatographed three times, and the data were reported as peak area percentages. The identification of the TAG peak of the samples was done in accordance with the TAG profiles of animal fats reported previously by Marikkar *et al.* (2002).

Thermal analysis by DSC

Thermal analyses of all animal fats were carried out on a Mettler Toledo differential scanning calorimeter (DSC 823 Model, Columbus, Ohio, USA). Data elaboration was performed with a thermal analysis software (STARe software, Version 9.0x, Schwerzenbach, Switzerland). Nitrogen (99.99% purity) was used as the purge gas at a rate of ~20 mL/min. Approximately 4–8 mg of molten sample was placed in a standard DSC aluminum pan and then hermetically sealed. An empty, hermetically sealed DSC aluminum pan was used as the control. The samples were held at 70°C isotherm for 1 min to eliminate the thermal history of the samples, then cooled at 5°C/min to –70°C and held for 1 min. The sample was then brought from –70°C to 70°C at the same rate (Marikkar *et al.*, 2001; Nina Naqiyah *et al.*, 2017).

Statistical analysis

All the analyses were performed in triplicate and the results were expressed as mean value ± standard deviation. Data were statistically analyzed by one-way analysis of variance (ANOVA) and Tukey's HSD post hoc test using MINITAB™ Statistical Software (MINITAB®

Release 14.12.0, New York, USA) at $\alpha = 0.05$. Principal component analysis (PCA), which is a multivariate statistical method that entails data reconstruction and reduction, was carried out using Unscrambler 9.7 (Camo, USA) software.

Results and Discussions

Fatty acid distribution

Fatty acid composition is one of the important attributes that determine the nature of most natural lipids. In this study, the fatty acid composition of four common animal fats is compared as shown in Table 1. In NL and other animal fats, oleic acid ($C_{18:1}$) has been the most dominant fatty acid. Palmitic acid ($C_{16:0}$), linoleic acid ($C_{18:2}$), and myristic acid ($C_{14:0}$) are other fatty acid types that occur in appreciable amounts in NL. Other than these, occurrence of fatty acids such as $C_{14:1}$ and $C_{15:0}$ was rarely detected in NL, but they were detected in minute amounts in other animal fats. According to Haas (2005), the prevalent fatty acids of animal depot lipids are either 16 or 18 carbons in their chain length and are either fully saturated or contain one or two double bonds. Investigations by Marikkar and Yanty (2014) showed that NL was generally composed of higher amounts of unsaturated fatty acids (50.2–74.72%) than saturated fatty acids (33.76–46.08%). The relative variations of individual fatty acids in NL has been mainly attributed to differences in sex and the breed types, climatic conditions due to the diet, and the system of animal rearing (Enser, 1995; Muriel *et al.*, 2002). According to Che Man *et al.* (2005), experimental diets composed by soybean or canola oil could cause an increase in the proportions of unsaturated fatty acids in swine fat. Among samples extracted from the different body parts of the animals, the fatty acid composition of NL can also slightly vary. However, in the majority of the cases, oleic acid was always found to be the most predominant fatty acid of NL (Enser, 1995). Similar to NL, CF was also found to possess high oleic acid (43.94%), followed by palmitic acid (25.39%) (Table 1). These values were closely comparable to those reported previously by Lee and Foglia (2000a). In contrast to both NL and CF, the distribution of fatty acids in BT and MT would be remarkably different. Particularly, the stearic acid content of BT and MT might differ from that of NL, which is low in saturated fatty acids. Hence, this feature was thought to be helpful to distinguish NL from BT/MT. Even though the presence of high stearic acid content is a common characteristic feature of both BT and MT, it may be varied slightly due to the influence of various other factor factors, namely, breed, sex, and nutritional conditions (Grompone, 1989; Grompone and Moyna, 1983; Holia and Press, 1987; Yilmaz and Karakaya, 2009).

The degree of unsaturation could also be considered a parameter for the discrimination of NL from other common animal fats. As shown in Table 1, unsaturated/saturated (US/S) fatty acid ratio was found to be useful in discriminating NL from BT and MT, but not NL and CF. This finding was largely in conformity with the results previously reported by Rashood *et al.* (1996). If either overall fatty acids (FA) distribution or degree of unsaturation was considered singularly, differentiating NL from CF might not be feasible. In such situations, applying multivariate data analysis techniques such as PCA on the overall fatty acid data would be beneficial. In essence, PCA helps to reduce the number of observed variables into a smaller number of variables that accounts for most of the variance in a particular dataset. An outcome of PCA would possibly group samples with dissimilar characteristics into different groups. As shown in Table 1, fatty acids such as lauric acid ($C_{12:0}$), myristic acid ($C_{14:0}$), palmitic acid ($C_{16:0}$), palmitoleic acid ($C_{16:1}$), stearic acid ($C_{18:0}$), oleic acid ($C_{18:1}$) and linoleic acid ($C_{18:2}$) were present in all four types of animal fats. Henceforth, they can be taken as independent variables to develop a PCA model.

The score plot shown in Figure 1 represents the projection of samples defined by the principal component 1 (PC1, which explains 86% of the dataset variance) and the principal component 2 (PC2 which explains 9% of the dataset variance). According to the score plot, a good group separation was achieved with the PCA model, in such a way that NL samples were located in the lower left quadrant, CF in the upper left quadrant, MT in the upper right quadrant, and BT in the lower right quadrant. According to the loading plot shown in Figure 2, stearic acid ($C_{18:0}$), oleic acid ($C_{18:1}$), and linoleic acid ($C_{18:2}$) were the most discriminating parameters that influence separation along the PC 1 axis, while palmitic acid ($C_{16:0}$) and palmitoleic acid ($C_{16:1}$) were the most discriminating parameter along the PC 2 axis.

Determination of *sn*-2 positional distribution of fatty acids within the glycerol chain using the pancreatic lipase hydrolysis method was also adopted as an alternative option to find effective discrimination between NL and other animal fats. This procedure is laborious, as it involves isolation of triacylglycerol using column chromatography followed by preparation of 2-MAG by pancreatic lipolysis. Calculation of the fatty acid enrichment factor (FAEF) corresponding to individual fatty acid could be possible based on the FA distribution of 2-MAG. As shown in Figure 3, FAEF corresponding to the palmitic acid was quite high for NL (283%) when compared to that of any other animal fat [BT (69%); MT (70%); CF (84%)]. This was because more than 74% of the fatty acids present in 2-MAG of NL were saturated fatty acids, of which about 68% was palmitic acid (Marikkar

Table 1. Fatty acid compositions (% peak area) of native lard, chicken fat, beef fat, and mutton fat.¹

Fatty acids	Native lard	Chicken fat	Beef fat	Mutton fat
C10:0	0.08 ^b ± 0.01	—	—	0.29 ^a ± 0.06
C12:0	0.19 ^a ± 0.11	0.64 ^a ± 0.93	0.11 ^a ± 0.05	0.46 ^a ± 0.06
C14:0	2.28 ^b ± 1.10	1.62 ^b ± 0.65	6.15 ^a ± 0.31	6.40 ^a ± 0.49
C15:0	0.05 ^c ± 0.04	—	0.46 ^b ± 0.24	0.76 ^a ± 0.02
C16:0	24.64 ^b ± 1.90	25.39 ^b ± 1.01	31.07 ^a ± 0.78	27.38 ^{ab} ± 1.22
C16:1	1.07 ^c ± 0.46	5.32 ^a ± 0.48	2.56 ^b ± 0.07	0.52 ^c ± 0.18
C17:0	0.25 ^b ± 0.23	—	0.82 ^b ± 0.62	1.85 ^a ± 0.13
C18:0	11.53 ^c ± 1.67	4.84 ^d ± 0.18	16.53 ^b ± 1.25	30.90 ^a ± 0.50
C18:1	42.62 ^a ± 0.74	43.94 ^a ± 1.77	35.70 ^b ± 1.71	29.82 ^c ± 1.04
C18:2	17.29 ^a ± 3.11	18.26 ^a ± 1.64	6.59 ^b ± 0.61	1.61 ^b ± 0.06
∑SFA	39.02	32.48	55.15	68.05
∑USFA	60.98	67.52	44.85	31.95
US/S	1.56	2.08	0.81	0.47

¹Each fatty acid value in the table represents the mean ± standard deviation of three replicates. Means within each row with different superscripts are significantly ($P < 0.05$) different.

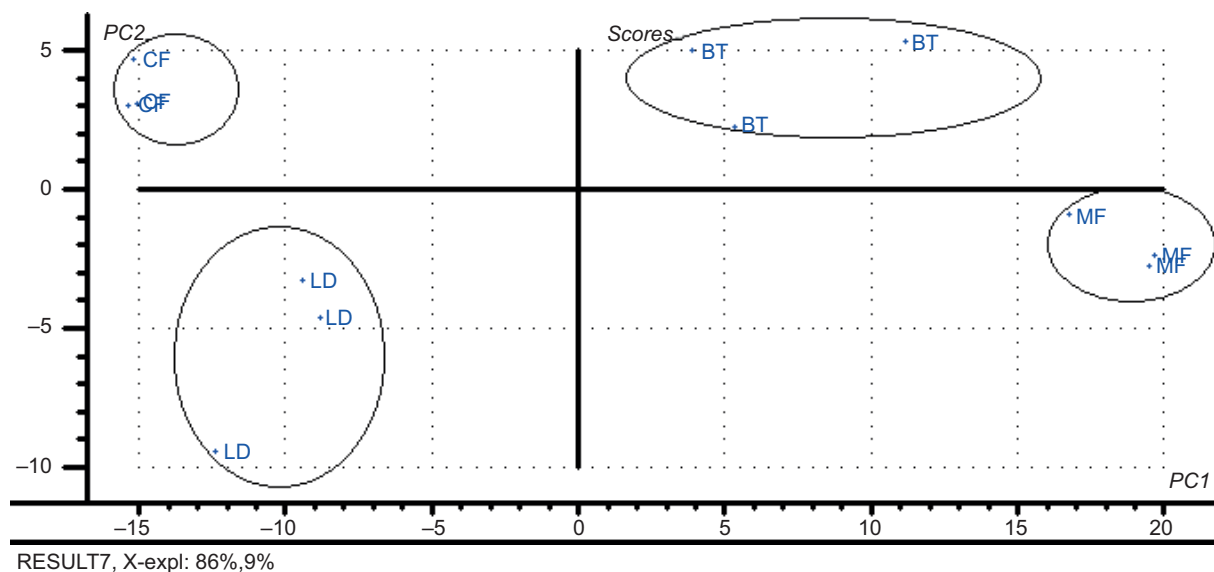


Figure 1. Score plot of PCA based on fatty acid composition for the discrimination of animal fats. Abbreviations: LD, lard; CF, chicken fat; BT, beef tallow; MT, mutton tallow.

et al., 2005; Rashood *et al.*, 1996). This characteristic feature could be effectively used to differentiate NL from other animal fats.

TAG distributional pattern

Comparative TAG distributional ratios of different triacylglycerol classes of animal fats are presented in Table 2. The TAG molecules present in most natural oils and fats could be classified into four subgroupings: trisaturates (SSS), monounsaturates (SUS/SSU), diunsaturates (SUU), and triunsaturates (UUU) by considering their degree of unsaturation. According to Table 2, there was

hardly any significant ($P > 0.05$) difference among animal fats with respect to the amount of triunsaturates (UUU), but the amount of trisaturates (SSS) was remarkably higher for both BT and MT than those for either NL or CF. This was due to the fact that the major TAG molecular species of NL were PLL, OOL, LPO, OPO, PPO, and SPO, of which SPO, LPO, and OPO had become the most dominant TAG molecular species (Marikkar *et al.*, 2002; Yanty *et al.*, 2011).

On the other hand, the most commonly found TAG molecular species of BT and MT were SSP, SSS POO,

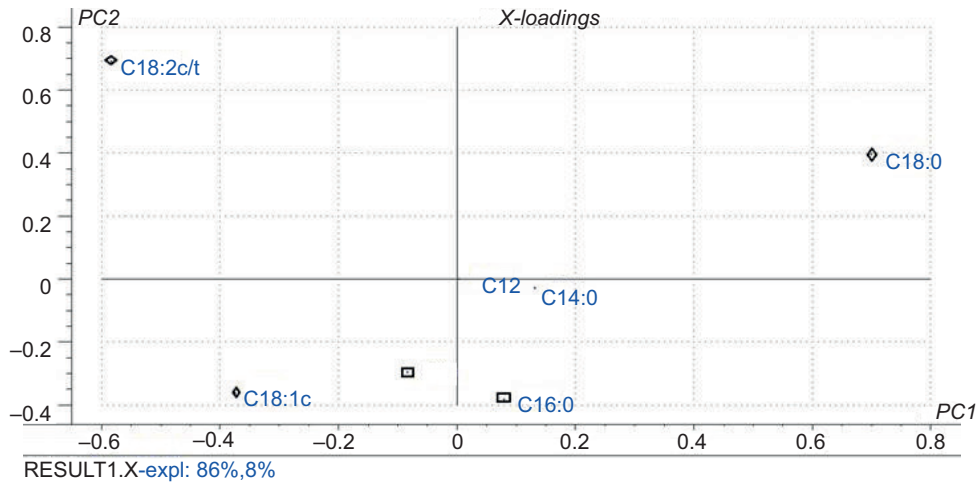


Figure 2. Loading plot of PCA of the animal fats based on the fatty acid composition. (PC1: \diamond , PC2: \square).

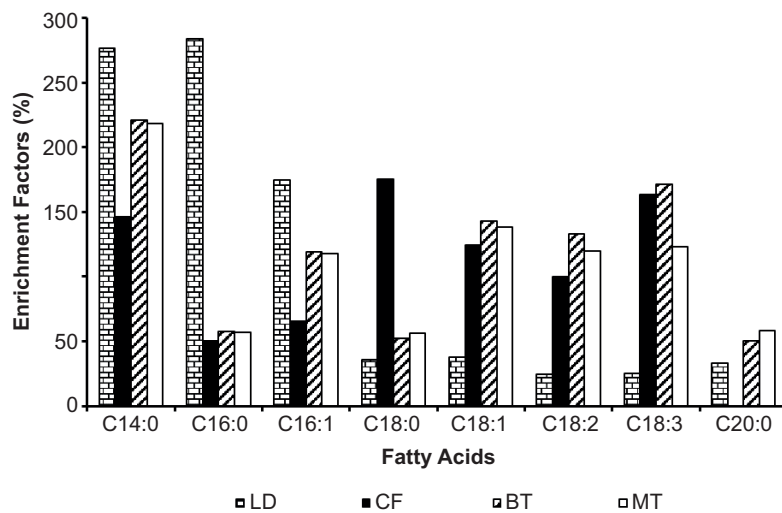


Figure 3. Comparative fatty acid enrichment factors (%) of lard and other common animal fats. Abbreviations: LD, native lard; BT, beef tallow; MT, mutton tallow; CF, chicken fat.

Table 2. Comparison of trisaturated (SSS), monounsaturated (SSU), diunsaturated (SUU), and triunsaturated (UUU) triacylglycerol distribution among different animal fats.¹

Parameter	Native lard	Chicken fat	Beef fat	Mutton fat
% Trisaturated	36.15 ^c ± 0.30	33.99 ^d ± 0.44	63.07 ^a ± 0.65	55.11 ^b ± 0.58
% Monounsaturated	44.86 ^b ± 0.26	51.52 ^a ± 0.72	34.62 ^d ± 0.35	41.86 ^c ± 0.51
% Diunsaturated	18.22 ^a ± 0.28	13.76 ^b ± 0.55	1.84 ^c ± 0.33	2.11 ^c ± 0.45
% Triunsaturated	0.69 ^a ± 0.10	0.75 ^a ± 0.14	0.51 ^a ± 0.07	0.95 ^a ± 0.17
US/S ratio	1.76:1	1.94:1	1:1.71	1:1.23

¹Each value in the table represents the mean of three replicates. Means within each row with different superscripts are significantly (P < 0.05) different.

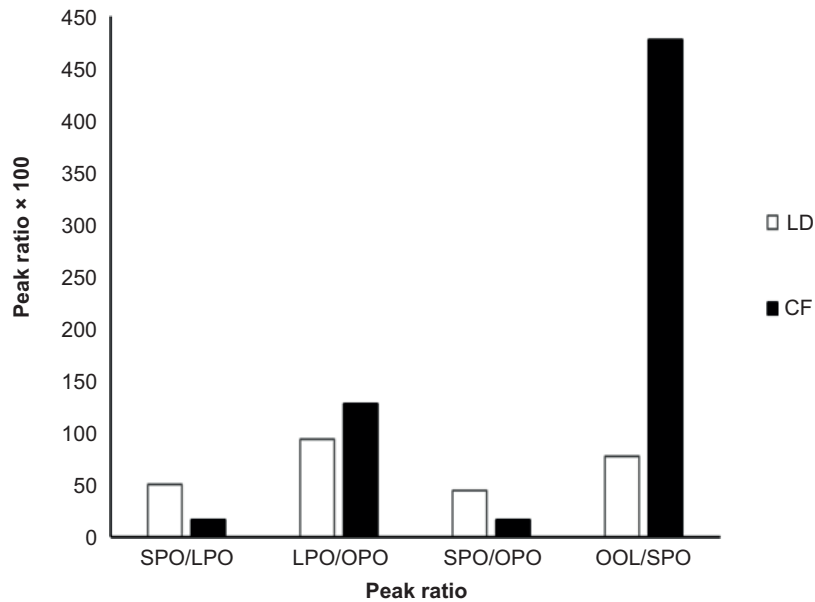


Figure 4. Selected TAG peak ratio of lard and chicken fat. Abbreviations: LD, native lard; CF, chicken fat.

PPO, PPP, SOO, PSO, SPP, SSO, and OOO (Marikkar *et al.*, 2002). Some early investigations of the 1960s suggested that NL possessed 38% of SSU, 41% of USU, and very low percentages of SUS (1%) and UUS (7%) (Chacko and Perkin, 1965). According to the literature, UUU and SUU were the most abundant TAG species present in CF (Lee and Foglia, 2000b). Among these, LOO, OOO, OPO, and PSO were found to be predominant in CF (Marikkar *et al.*, 2002). Some major differences in the TAG distribution were found between NL and BT/MT, and those differences can make them easily distinguishable. Hence, the US/S ratio for NL (1.76:1) was remarkably different from those of either MT (1:1.23) or BT (1:1.71). On the contrary, the differences in the US/S ratio between NL (1.76:1) and CF (1.94:1) were minor, which could make their discrimination more difficult. In order to discriminate between NL and CF, TAG peak ratios such as SPO/LPO, LPO/OPO, SPO/OPO, and OOL/SPO can be calculated. According to Figure 4, NL and CF displayed only slight differences among ratios of SPO/LPO, LPO/OPO, and SPO/OPO, but showed a strong difference with respect to the OOL/SPO ratio. In this case, CF was found to show higher OOL/SPO ratio than that of NL.

DSC thermal profiles

The overlay presented in Figure 5 shows DSC cooling curves of NL (Curve A), BT (Curve B), MT (Curve C), and CF (Curve D). DSC cooling curves are recognized worldwide to provide thermal characteristics of plant oils (Tan and Che Man, 2000), animal body fats (Yilmaz and Karakaya, 2009), lard (Yanty *et al.*, 2011), milk fat (Campos *et al.*, 2002), etc. According to the cooling curve of NL (Figure 5; Curve A), there were two well-separated

and defined sharp exothermic thermal transitions, indicating two steps of crystallization; transition above 0°C was defined as the high-melting group (HMG) while the transition below 0 °C was identified as the low-melting group (LMG). These thermal features are largely in conformity with the results mentioned in previous reports (Marikkar *et al.*, 2001; Rashood *et al.*, 1996; Yanty *et al.*, 2011). However, a few differences observed in the onset temperatures and heat flux profiles could be possible due to either sample to sample variation or variation in the DSC scanning rate. In an early report, Campos *et al.* (2002) demonstrated the effect of different cooling rates on the DSC curve of NL. As illustrated in Figure 5, the cooling profiles of other animal fats were more complex heat flux curves involving multiple thermal transitions as compared to that of NL. Particularly, peak broadening is a characteristic feature seen in Curves B, C, and D, indicating a wider range of melting points of the constituent TAG molecules. Both BT and MT showed sharp thermal transitions at 29.8 and 26.5°C, and broad crystallization transitions at 5.9 and 4.9°C, respectively. The cooling profile of BT was comparable to that of buffalo body fat reported previously by Lambelet and Ghanguli (1983). The occurrence of broader melting peaks in the low-temperature range in BT and MT could be due to the difference in the nature of TAG distributional pattern in these animal fats as compared to that of NL. The cooling profile of CF represented by curve D, displayed a single sharp peak at 8.9°C and a broad peak at -1.9°C. Overall, the thermal events recorded by CF in this study were roughly similar to the one reported previously by Lee and Foglia (2000b). As these features were quite dissimilar from those of NL, the thermal profile obtained

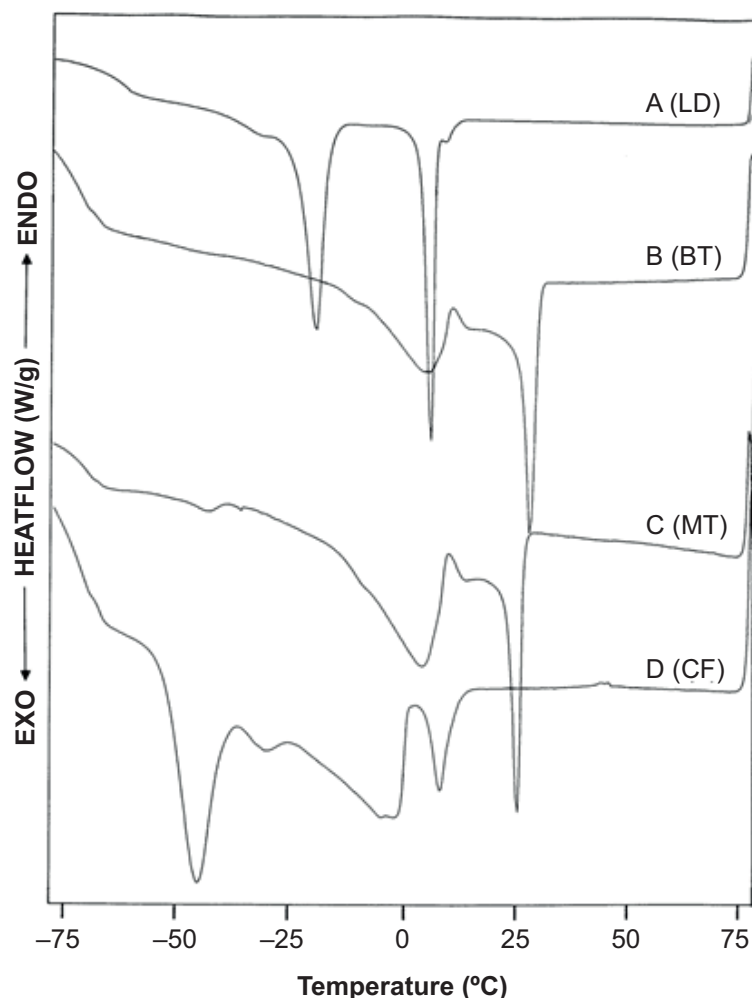


Figure 5. DSC Cooling thermograms of lard and other common animal fats. Abbreviations: LD, native lard; BT, beef tallow; MT, mutton tallow; CF, chicken fat.

from this study could provide a basis for differentiating CF from NL.

A comparison between the heating profiles of NL and other animal fats is shown in Figure 6. As previously seen with cooling curves, the heating curves of NL also had two well-separated endothermic transitions, identified as the low- and high-melting regions (Curve A). The low-melting region consisted of a major sharp peak at 1.10°C with a shoulder-peak at -17.2°C, while the high-melting transition was represented by a sharp peak with the maxima at 28.25°C. In the melting profile of CF (Curve B), there was substantial overlap between the low- and high-melting transitions. The low-melting region consists of a major peak at 1.85°C with two shoulder peaks at -8.0 and -23.0°C, respectively, while the high-melting region is represented by a broad peak resulting from the overlap of two adjacent peaks appearing at 17.2 and 35.6°C, respectively. In the case of BT and MT, their melting curves were comparably similar to each other, but they were remarkably different from

those of NL and CF. In the melting curve of BT, the major peak was located at 12.1°C, with two smaller shoulder peaks at 6.5 and -15.0°C, respectively, representing the low-melting region, while the peak appearing at 47.5°C with a shoulder at 34.1°C represented the high-melting region (Figure 6). This curve was comparably similar to the DSC melting profile of BT reported previously by Rodriguez *et al* (2001) and Aktas and Kaya (2001). The melting profile of MT (Figure 6, Curve D) also showed four steps of melting transitions. The low-melting region of MT is represented by a major peak at 11.4°C with two smaller shoulder peaks at 4.05 and -15.00°C, respectively, while the HMG fraction had a major peak at 45.10°C with a shoulder peak at 32.65°C.

Conclusions

A comparison between lard and other common animal fats has been made on the basis of fatty acid, TAG composition, and DSC thermal profiles. The different nature

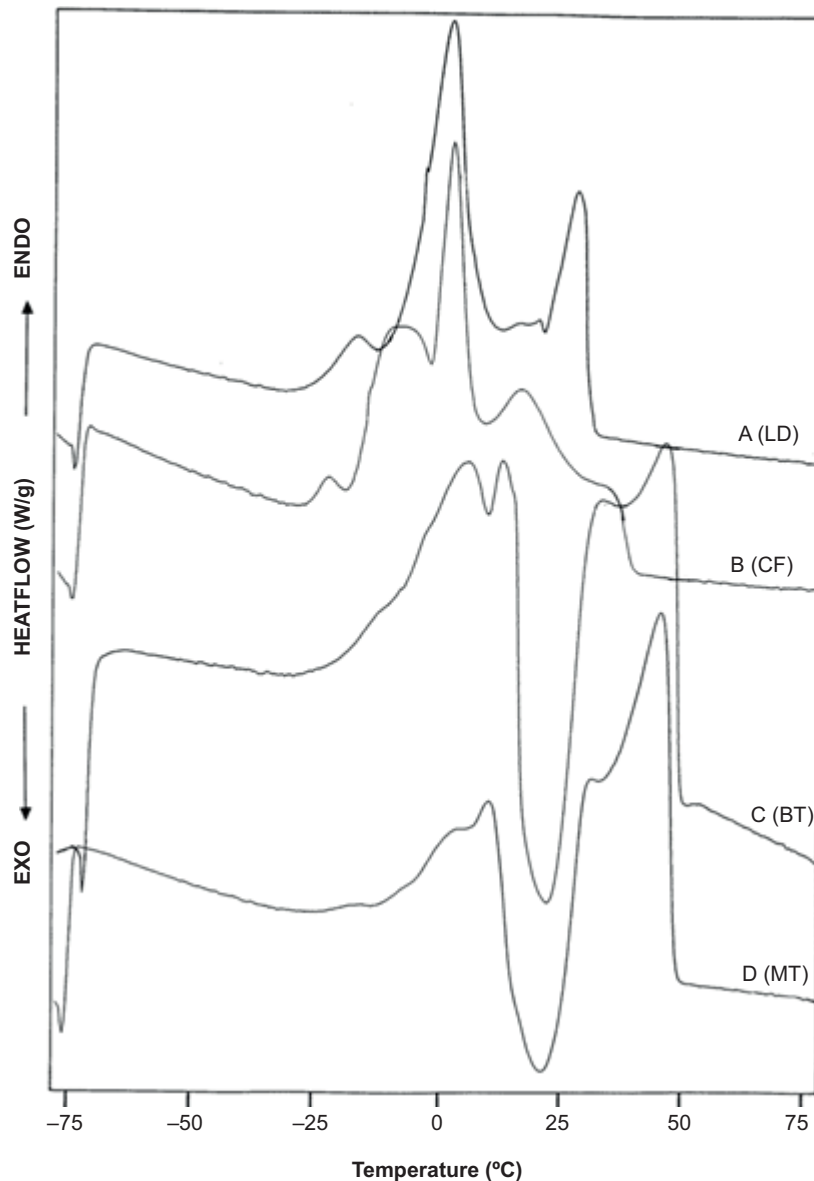


Figure 6. DSC heating thermograms of lard and other common animal fats. Abbreviations: LD, native lard; BT, beef tallow; MT, mutton tallow; CF, chicken fat.

of esterification of different FA in TAG molecules is the primary factor that makes animal fats to differ in their chemical and physical characteristics. Fatty acid compositional analysis showed that PAEF of lard was much higher than those of other common animal fats. Major differences between NL and BT/MT were also found on the basis of TAG composition. The degree of unsaturation of the TAG species in animal fats had a strong influence on the TAG separation in the reverse-phase silica column and could help differentiate different animal fats. Lard displayed a DSC cooling curve with two distinct exothermic sharp peaks at 4.9 and -16.9°C, respectively, while cooling curves of other animal fats were more complex, involving multiple thermal transitions, resulting in broader peaks.

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