Value-addition of Beef Meat By-products: Lipid Characterization by Chromatographic Techniques

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Abstract: The lipid characterization of storage fat, subcutaneous fat and bone marrow, originated from three different bovine categories, calf, young bull and cow, was carried out in order to develop a re-use of these by-products. After the lipid extraction, the compositions in fatty acids, triacylglycerols and cholesterol were determined by GC-FID. A total of 25 fatty acids were identified in all by-products; the oleic acid was the preponderant component, followed by palmitic, stearic, palmitoleic, linoleic and myristic acid. The long chain triacylglycerols (T48, T50 and T52) were the main products, especially in the cow’s by-products. The subcutaneous fat of all animals showed the highest cholesterol content, whereas it was present in low amount in the others. Phospholipids were analysed in bone marrow of all the animals and phosphatidylcholine was the most abundant compound. Because of the high fat content and essential fatty acids and the low cholesterol amount, storage fat and bone marrow could represent a valuable lipid resource in food and pharmaceutical industry.

Key words: beef by-products, cholesterol, fatty acids, phospholipids, triacylglycerols

1 Introduction

Most animal by-products originate during slaughter of cattle for human consumption, the production of dairy products, the disposal of carcasses and disease control. Animal by-products mean entire body or parts of it, products of animal origin or also products which are not for human consumption, including oocytes, embryos and semen (Regulation (EC) No 1069/2009). In order to reduce by-products generation, different industrial processes are used in the meat industry, but usually these by-products are discarded as waste or used for low-value purpose⁴. The large amounts of unused by-products cause a big loss of profit and an expense for their disposal, generating an increase in the cost of meat production. The use of these by-products not only increases profits, but also deals with the problems of environmental pollution associated with their disposal⁵. In particular, fat obtained from animal by-products can be melted fractionated and used in numerous cosmetic applications, such as the formulation of body lotions or other bath products. The fatty acids are used in numerous chemical processes including the polymerization of rubber and plastic, the production of fabric softeners, lubricants and plasticizers. Collagen, gelatin and glycerol are employed as ingredients to produce antifreeze substances, surfactants, paints, adhesives, detergents, as well as for pharmaceuticals⁶. The fat percentage in beef carcasses ranging from 30 to 40%, depending on endogenous and exogenous factors (ISMEA, 2015). The fat of the meat can be classified into different types: subcutaneous fat, which is formed by large adipocytes in the connective tissue and it is a reserve of energy on the surface of muscles; intermuscular fat, which is located in different muscles in the same anatomic part and it is mainly composed by triacylglycerols and saturated fatty acids as the subcutaneous fat. Intramuscular fat, the fat marbling, is located between the muscle fibres and the fibre bundles and it mainly influences the nutritional characteristics of meat; finally, intracellular fat is located within muscle fibres and it is particularly rich in phospholipids⁷. Half of the fatty acid total content of bovine meat is mainly composed by saturated fatty acids. The pattern of these compounds is partly affected by the animal species, breed, age and farmed technique. Unlike monogastric animals, the microflora in the rumen is able to hydrogenate most of the unsaturated fatty acids introduced by feeding, increasing saturated lipids⁸. Several studies report the positive bio-

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logical activities of food phospholipids on human health by virtue of both lipophilic and hydrophilic properties. The phospholipid content in beef muscle ranged between 0.5 and 0.6 g/100 g of fresh muscle.

The aim of this study is a screening of the lipid quality and composition of different by-products produced by bovine meat industry. The valorisation of these wastes is an important goal for the profit of the meat industry and for the environmental impact, avoiding their disposal. At the same time, literature is lacking in works on lipid quality of beef by-products, so these data can be of great importance in the promotion of their re-utilization, developing new technological applications.

2 Materials and Methods

2.1 Samples

A total of three by-products produced by a slaughterhouse in the north of Italy were analysed: storage fat, subcutaneous fat and bone marrow. These by-products were obtained from three Friesian bovine categories, differing in age and morphology, that is calf (CA), young bull (YB) and cow (CO). Calf was fed only with milk due to the young slaughter age. Young bull and cow were fed with a traditional diet, based on the use of concentrate and hay (dry feeding period) and fresh forage (fresh forage feeding period) without use of silages. The different categories were slaughtered according to slaughter legislation: calves were slaughtered at 5-6 months old, 200-250 kg; young bulls at 16-18 months old, 600-650 kg and cows at 14-18 months old, 420-480 kg. These samples are the result of all the waste obtained at the end of a working day; approximately thousand animals for each category. Before the lipid extraction, in order to obtain a homogeneous sampling and to facilitate the breaking of fat cells, all by-products were dissected and crumbled with knives and blade blender and then sonicated at room temperature.

2.2 Lipid extraction

The lipid fraction of the samples was extracted using the procedure described by Folch et al., which has been slightly modified as reported elsewhere. Each extraction was carried out three times for each by-product (n = 9 for animal sample).

2.3 Fatty acid analysis

The fatty acid composition was determined as fatty acid methyl esters (FAMEs) by capillary gas chromatography analysis after alkaline treatment. Methyl tridecanoate (C13:0, 2 mg/mL) was used as internal standard and FAMEs were measured on a GC 2010 Plus gas chromatograph (Shimadzu Corporation, Kyoto, Japan) equipped with a flame ionisation detector (FID) and an AOC-20s auto sampler (Shimadzu Corporation), according to Verardo et al., with slight modifications. The FAMEs were analysed using a BPX70 fused silica capillary column (10 m × 0.1 mm i.d. 0.2 μm film thickness; SGE Analytical Science, Ringwood, VIC, Australia). The injector and flame ionization detector temperatures were set at 250°C. Hydrogen was used as carrier gas at a flow rate of 0.8 mL/min. The oven temperature was held at 50°C for 0.2 min, increased to 175°C at 120°C/min, held at 175°C for 2 min and finally increased from 175 to 220°C at 20°C/min. Samples were injected in split mode (0.3 μL) with a split ratio set at 1:100. Peak identification was accomplished by comparing peak retention time with GLC-463 standard mixture from Nu-Check (Elysian, MN, USA) and FAME 189-19 standard mixtures from Sigma-Aldrich Chemicals (St. Louis, MO, USA) and expressed as weight percentage of total FAMEs. FAMEs composition was measured in 2 replicates for each lipid extract (n = 6) and each analysis lasted 7 minutes.

2.4 Triacylglycerols analysis

Triacylglycerols (TGs) analysis was carried out with the injection of 1.0 μL of solution (10 mg/mL of fat in n-hexane) into a GC-2010 Plus gas chromatograph (Shimadzu Corporation, Kyoto, Japan) equipped with a flame ionisation detector (FID) and an AOC-20s auto sampler (Shimadzu Corporation) according to Guerra et al., with slight modification. TGs separation was performed with an Rtx-65 TG fused silica capillary column (30 m × 0.25 mm × 0.1 μm film thickness) with 35% dimethyl, 65% diphenyl polysiloxane (Restek, Chromatography Products, Super- chrom Milano, Italy). The initial oven temperature of 140°C was raised to 360°C at a rate of 25°C/min and was held at 360°C for 5 min. The injector and detector temperatures were set at 360°C. The hydrogen flow rate was 3.84 mL/min. The split ratio was 1:30. TGs were identified based on retention time with GLC-463 standard mixture from Nu-Check (Elysian, MN, USA) and Elysian. MN, USA and each analysis lasted 26 minutes.

2.5 Cholesterol determination

Cholesterol was collected by cold saponification at room temperature after addition of 500 μL of internal standard (dihydrocholesterol, 2 mg/mL) to 250 μg of fat. The unsaponifiable fraction was evaporated by vacuum evaporator, silylated and dried again under gentle nitrogen flow. After redisolution in 500 μL of n-hexane, 1 μL was injected into a GC 2010 Plus gas chromatograph (Shimadzu, Kyoto, Japan), equipped with a flame ionisation detector (FID).
and an AOC-20i autosampler, according to the method reported by Guerra et al.\textsuperscript{18}. A Rxi-5ms fused silica capillary column (10 m × 0.10 mm i.d. × 0.10 μm film thickness) from Restek (Restek Corporation, Bellefonte, USA) was used. Oven temperature was set at 240°C and the injector and detector temperatures were 325°C. The column flow of hydrogen was 0.4 mL/min and the injection volume was 0.30 μL, with a split ratio of 1:50. Cholesterol content was measured in 2 replicates for each lipid extract (n = 6) and was expressed in mg cholesterol/kg of fat; and each analysis lasted 15 minutes.

2.6 Phospholipids determination

The phospholipids (PLs) extraction was made according to Avalli and Contarini\textsuperscript{19} by a purification of the lipid extracts with solid-phase extraction (SPE) cartridges. The identification and quantification of PL classes was performed using an Agilent liquid chromatography HP 1200 Series (HPLC; Agilent Technologies, Palo Alto, CA, USA) combined with an evaporative light-scattering detector (ELS; PL-ELS1000, Polymer laboratories, Church Stretton, Shropshire, UK). The separation of PLs was achieved using a silica column, 150 mm × 3 mm with 3 μm particle diameter (Phenomenex, Torrance, CA, USA) and applying the method by Verardo et al.\textsuperscript{20} with some modifications. The HPLC system was controlled by Agilent Chem-Station software (Agilent Technologies), whilst chromatogram and data processing were assessed by ClarityLite (ver. 2.4.0.190, Data- Apex, Praha, Czech Republic). PLs were identified by comparison with pure standards and quantified with external calibration curves, prepared separately for each phospholipid identified (from 1 to 500 mg/mL of PE, PI, PS, PC and SM). Each analysis lasted 36 minutes.

2.7 Statistical analysis

Relative standard deviation was obtained, where appropriate, for all data collected. One-way analysis of variance (ANOVA) was evaluated using Statistica 8 software (2006, StatSoft, Tulsa, OK, USA). The differences between the means of data for the three different by-products (storage fat, subcutaneous fat and bone marrow) and for the three different animals (calf, young bull and cow) were compared at the 5% level of significance (p < 0.05) using Tukey honest significant difference (HSD) test.

3 Results and Discussion

3.1 Total lipid content of by-products

The fat content of the different by-products is reported in Table 1 and it is expressed as percentage (%) on the total fresh weight of the sample. The calf’s storage fat and bone marrow showed a significant (p < 0.05) lower fat content compared to the other two animals. This result reflects the different characteristics of the animals in regard to their age, morphology and feeding\textsuperscript{16} in addition to the cut and degree of trimming\textsuperscript{21}. On the other hand, young bull (YB) and cow (CO) did not show significant differences (p > 0.05) between them with more than 80% of storage fat content and 85% of bone marrow.

Differently, the subcutaneous fat was present in not different amounts (p > 0.05) among the animal samples. For this by-product, the repeatability of the data was poor probably due to the non-homogeneous matrix. In fact, the standard deviations of these data were very high, causing a flattening of the differences between samples.

3.2 Fatty acid composition of by-products

A total of 25 individual fatty acids were identified and quantified in all samples by fast GC-FID analysis, within a run time less than 7 min, due to a fast GC-FID system. As shown in Table 2, the predominant fatty acid in all by-products of the different animals was oleic acid (C18:1 cis9), ranging from about 32 to 46%. Palmitic acid (C16:0) constituted the second major fatty acid detected (~20-28%), followed by stearic acid (C18:0, ~10-19%), palmitoleic acid (C16:1 cis, ~2-5%), myristic acid (C14:0, ~2-4%) and linoleic acid (C18:2 n6, ~2-10%).

In the storage fat, the saturated fatty acids (SFA) were present in significantly (p < 0.05) greater amounts in calf and young bull (46.76% and 47.62%), whereas mono-unsaturated fatty acids (MUFA) were more abundant in cow, in the amount of 55.70%. Polyunsaturated fatty acids (PUFA) were present in low concentrations in all animal samples, accounting for 5.21%, 3.81% and 3.50% for CA, YB and CO, respectively. These trends reflect those of the main individual fatty acids: CO showed the highest content of oleic acid (44.78%) and palmitoleic acid (5.13%), whereas YB and CA presented the main content of stearic acid (16.64%) and palmitic acid (26.44%), respectively. Besides, CA sample had the highest concentration in the C18:2 n6. According to De Smet et al.\textsuperscript{22} and Bruguiapaglia et al.\textsuperscript{21} about the investigation on the fatty acid profile of Longissimus thoracis muscle from different Belgian and Italian young bulls, SFA was the predominant class followed by MUFA and PUFA. For SFA class our results (range

| Table 1: Fat content of the different by-product samples. |
|----------|---------|---------|---------|
|          | CA      | YB      | CO      |
| Storage fat | 61.4 ± 4.1\textsuperscript{a} | 81.6 ± 4.6\textsuperscript{b} | 83.6 ± 0.5\textsuperscript{a} |
| Subcutaneous fat | 8.2 ± 0.7\textsuperscript{a} | 22.3 ± 8.8\textsuperscript{b} | 14.8 ± 7.0\textsuperscript{b} |
| Bone marrow | 70.5 ± 1.1\textsuperscript{a} | 86.6 ± 3.4\textsuperscript{a} | 89.8 ± 0.9\textsuperscript{a} |

Abbreviations: CA, calf; YB, young bull; CO, cow. Data (means ± SD) are expressed in percentage (%); results of the analysis of variance by Tukey’s test are shown: p < 0.05, lowercase letters on the same row show significantly different values within each by-product for the three animals.
between 40.80% in cow and 47.62% in young bull) were slightly lower than their data (43.62-48.51% \(^{22}\) and 46.05-49.25% \(^{23}\) ); instead for MUFA and PUFA we registered higher and lower concentrations respectively rather than these two works. For MUFA we had a range from 48.03%, in calf, to 55.70%, in cow (33.94-38.42% \(^{22}\) and 32.98-42.08% \(^{23}\) ); and for PUFA we had a range between 3.50, in cow, and 5.21%, in calf (15.04-20.48% \(^{22}\) and 10.69-21.87% \(^{23}\) ).

Also in subcutaneous fat, the principal classes were SFA and MUFA, showing not significant differences between the three bovine categories analysed and with a range between 45.52 and 49.59%, and between 45.37 and 49.73%, respectively. PUFA were present in considerably smaller amounts and with a significantly higher content in calf than the other two animals (\(p<0.05\)), as it was observed for storage fat. In fact, CA subcutaneous fat reported more linoleic acid (C18:2 n6) than YB and CO, in the amount of 4.69%. These results are in agreement with the study of Indurain et al. \(^{24}\) on the fatty acid profile of subcutaneous adipose tissue in young bulls. Other investigations were carried out on the fatty acid compositions of bulls and heifers. Schiavon and co-workers\(^{25}\) showed a similar MUFA content in beef subcutaneous fat, whereas SFA and PUFA were present in higher and lower amount compared to our results. Noci et al.\(^{25}\) have observed low SFA concentration in heifers and MUFA and PUFA with higher values than our samples.

In the bone marrow, animals showed more variability in SFA and MUFA contents. SFA were present in higher amount in YB sample, with an average content of 53.42%, followed with significant lower values by CO (45.29%) and CA (35.08%). On the other hand, MUFA were more abun-

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**Table 2** Fatty acids composition of by-products.

<table>
<thead>
<tr>
<th>FA</th>
<th>CA</th>
<th>YB</th>
<th>CO</th>
<th>Subcutaneous fat</th>
<th>Bone marrow</th>
</tr>
</thead>
<tbody>
<tr>
<td>C12:0</td>
<td>0.18±0.14^a</td>
<td>0.11±0.04^a</td>
<td>0.09±0.02^a</td>
<td>0.11±0.01^a</td>
<td>0.07±0.01^a</td>
</tr>
<tr>
<td>C14:0</td>
<td>3.57±0.52^a</td>
<td>3.73±0.61^a</td>
<td>3.04±0.03^a</td>
<td>2.81±0.05^a</td>
<td>2.36±0.10^a</td>
</tr>
<tr>
<td>C14:1c</td>
<td>0.81±0.06^a</td>
<td>0.94±0.06^a</td>
<td>1.87±0.06^a</td>
<td>0.81±0.05^a</td>
<td>1.12±0.19^a</td>
</tr>
<tr>
<td>C15:0</td>
<td>0.48±0.03^a</td>
<td>0.66±0.03^a</td>
<td>0.38±0.01^a</td>
<td>0.41±0.02^a</td>
<td>0.55±0.13^a</td>
</tr>
<tr>
<td>C15:1c</td>
<td>0.10±0.01^a</td>
<td>0.14±0.01^a</td>
<td>0.15±0.03^a</td>
<td>0.12±0.01^a</td>
<td>0.28±0.03^a</td>
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<tr>
<td>C16:0</td>
<td>26.44±0.66^a</td>
<td>29.42±0.65^a</td>
<td>25.63±0.19^a</td>
<td>25.03±0.46^a</td>
<td>27.06±0.58^a</td>
</tr>
<tr>
<td>C16:1 trans</td>
<td>0.29±0.05^a</td>
<td>0.24±0.02^a</td>
<td>0.20±0.06^a</td>
<td>0.32±0.03^a</td>
<td>0.25±0.05^a</td>
</tr>
<tr>
<td>C16:1 cis</td>
<td>3.62±0.36^a</td>
<td>3.71±0.17^a</td>
<td>5.13±0.10^a</td>
<td>3.31±0.08^a</td>
<td>3.90±0.17^a</td>
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<tr>
<td>C17:0</td>
<td>1.06±0.06^a</td>
<td>1.57±0.04^a</td>
<td>0.85±0.03^a</td>
<td>0.94±0.04^a</td>
<td>1.15±0.26^a</td>
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<td>C17:1</td>
<td>0.68±0.07^a</td>
<td>1.11±0.02^a</td>
<td>0.95±0.04^a</td>
<td>0.60±0.02^a</td>
<td>0.81±0.07^a</td>
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<tr>
<td>C18:0</td>
<td>15.03±0.79^a</td>
<td>16.64±0.52^a</td>
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<td>17.21±0.39^a</td>
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<td>C18:1 trans</td>
<td>8.52±1.19^a</td>
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<td>C18:1 cis 9</td>
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<td>36.81±0.61^a</td>
<td>44.78±0.63^a</td>
<td>37.13±0.54^a</td>
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<tr>
<td>C18:2 trans</td>
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<tr>
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<td>3.81±0.87^a</td>
<td>1.83±0.12^a</td>
<td>1.41±0.09^a</td>
<td>4.69±0.26^a</td>
<td>2.31±0.35^a</td>
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<td>C18:3 n6 +</td>
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<td>0.27±0.02^a</td>
<td>0.35±0.09^a</td>
<td>0.23±0.06^a</td>
<td>0.33±0.03^a</td>
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<tr>
<td>C18:3 n9</td>
<td>0.24±0.09^a</td>
<td>0.18±0.01^c</td>
<td>0.18±0.03^a</td>
<td>0.17±0.02^a</td>
<td>0.41±0.06^a</td>
</tr>
<tr>
<td>CLA</td>
<td>0.37±0.05^a</td>
<td>0.58±0.03^a</td>
<td>0.48±0.06^a</td>
<td>0.60±0.02^a</td>
<td>0.59±0.04^a</td>
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<tr>
<td>C20:1</td>
<td>0.27±0.05^a</td>
<td>0.21±0.02^a</td>
<td>0.47±0.04^b</td>
<td>0.33±0.02^a</td>
<td>0.43±0.06^a</td>
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<tr>
<td>C20:2 n6</td>
<td>0.08±0.02^a</td>
<td>0.09±0.02^a</td>
<td>0.05±0.01^a</td>
<td>0.09±0.02^a</td>
<td>0.05±0.01^a</td>
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<tr>
<td>C20:3 n6</td>
<td>0.07±0.01^a</td>
<td>0.07±0.01^a</td>
<td>0.16±0.06^a</td>
<td>0.16±0.05^a</td>
<td>0.16±0.02^a</td>
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<tr>
<td>C20:4 n6</td>
<td>0.07±0.01^a</td>
<td>0.06±0.01^a</td>
<td>0.06±0.01^a</td>
<td>0.45±0.06^a</td>
<td>0.20±0.05^a</td>
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<tr>
<td>C22:3 +5</td>
<td>0.07±0.01^a</td>
<td>0.07±0.01^a</td>
<td>0.07±0.01^a</td>
<td>0.10±0.04^a</td>
<td>0.18±0.01^a</td>
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<tr>
<td>C22:5 n3</td>
<td>0.07±0.01^a</td>
<td>0.09±0.02^a</td>
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<td>0.30±0.06^a</td>
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<td>SFA</td>
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<td>47.62±0.53^a</td>
<td>40.80±0.28^a</td>
<td>46.51±0.47^a</td>
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<td>MUFA</td>
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<td>48.57±0.64^a</td>
<td>55.70±0.50^a</td>
<td>46.29±0.69^a</td>
<td>49.73±4.92^a</td>
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<tr>
<td>PUFA</td>
<td>5.21±0.93^a</td>
<td>3.81±0.13^a</td>
<td>3.50±0.43^a</td>
<td>7.20±0.26^a</td>
<td>4.70±0.49^a</td>
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<tr>
<td>CHOL</td>
<td>867±2.1054^a</td>
<td>732.4±14.94^a</td>
<td>739.7±31.06^a</td>
<td>5192.4±899.6^a</td>
<td>3431±2.1926^a</td>
</tr>
</tbody>
</table>

Abbreviations: CA, calf YB, young bull; CO, cow; CLA, conjugated linoleic acid; SFA, saturated fatty acid; MUFA, monounsaturated fatty acid; PUFA, polyunsaturated fatty acid; CHOL, cholesterol. Data (means ± SD) are expressed in mg/FA/100 mg FAME (%).
Characterization of Lipid Fraction of By-products Originated from Bovine Meat Industry

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The conjugated linoleic acid (CLA) was present in all samples in a range between 0.23% of the CA’s bone marrow and 0.70% of the YB’s bone marrow.

The low PUFAs content of the analysed cattle’s by-products confirm what it is already reported in literature. Bovines are ruminants and the rumen microflora conduct a hydro-generation process of unsaturated fatty acids making them saturated, increasing the total SFA amount.

Compared to previous works on fatty acid profile of bovine samples, this study, obtained with a fast GC-FID, allows to get a large number of fatty acids in just 7 minutes of run time.

3.3 Determination of triacylglycerols

Because of fatty acids make up the bulk of a triacylglycerol molecule, this study also focused on triacylglycerols (TGs) composition of samples, using a fast GC-FID analysis with a time of less than 13 min. As reported in Table 3, six classes of triacylglycerols were identified and quantified in all by-products, corresponding to TGs with 42 to 52 carbon number (CN).

All the by-products showed a high content of long-chain TGs (CN48, CN50 and CN52), with CN50 as the most abundant class, expressing about the 50% of the total TGs in samples. Following the TGs with CN48 were present with a content of about 18-26% and in decreasing amount the CN52 (~12-19%), CN46 (~5-13%), CN44 (~1-4%) and CN42 (~0.2-0.9%) classes. These results reflect a close relationship between the fatty acids and TGs content of samples. Indeed, a high content in TGs CN50 and CN48 was coincident with high levels of C18:1, C18:0, C16:0 and C14:0.

In storage fat, the three animals did not show significant differences in TGs content from CN42 to CN48. Whereas for the long-chain TGs, CN50 and CN52, CA samples presented a significantly lower and higher content, respectively, than the other two animals. More variability was observed for the TG content in subcutaneous fat and mainly for the long-chain classes, CN50 and CN52. In the first case, CO showed the highest content followed by YB and CA, whereas the CN52 class was more concentrated in CA than YB and CO. Finally, the TG distribution in the bone marrow samples were significantly different among the three bovine categories. In particular, TGs content from CN42 to CN48 was significantly high in YB, followed by CA and CO; on the contrary, TGs with CN50 were present in high amount in CO and TGs with CN52 in CA (Table 3).

3.4 Cholesterol content of by-products

Cholesterol is a sterol of animal origin which belong to lipid unsaponifiable fraction. It is predominantly localized in the membrane of blood cells and it can affect their fluidity.

As shown in Table 2, no differences (p<0.05) were found among the three animals for each by-product, whereas each animal showed a significantly higher cholesterol content in subcutaneous fat compared to the storage fat and bone marrow. This result was plausible being cholesterol a leading and fundamental component of cell membranes. In the other two by-products, storage fat and bone marrow, no significant differences were found ranging from 732.4 mg/kg in the storage fat of YB to 1160.3 mg/kg in the bone marrow of CO. Despite the lack of literature about cholesterol content in beef by-products, our results are higher than other studies. Several authors investigated the cholesterol amount on beef meat and all of them had lower cholesterol content than our by-products (460.8-490.7 mg/kg, 511.0-630.0 mg/kg, 30.0-340.4 mg/kg and 360.3-410.5 mg cholesterol/kg).

3.5 Phospholipids content of bone marrow

The phospholipids content and the distribution of individual phospholipid species were determined only in the

<table>
<thead>
<tr>
<th>TGs</th>
<th>Storage fat</th>
<th>Subcutaneous fat</th>
<th>Bone marrow</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>CA</td>
<td>YB</td>
<td>CO</td>
</tr>
<tr>
<td>CN42</td>
<td>0.5 ± 0.2&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0.4 ± 0.0&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0.5 ± 0.0&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>CN44</td>
<td>2.9 ± 0.5&lt;sup&gt;a&lt;/sup&gt;</td>
<td>2.9 ± 0.1&lt;sup&gt;b&lt;/sup&gt;</td>
<td>3.2 ± 0.1&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>CN46</td>
<td>11.3 ± 0.5&lt;sup&gt;a&lt;/sup&gt;</td>
<td>11.4 ± 0.3&lt;sup&gt;a&lt;/sup&gt;</td>
<td>11.7 ± 0.3&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>CN48</td>
<td>26.0 ± 0.7&lt;sup&gt;a&lt;/sup&gt;</td>
<td>26.0 ± 0.6&lt;sup&gt;a&lt;/sup&gt;</td>
<td>25.5 ± 0.2&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>CN50</td>
<td>42.1 ± 1.3&lt;sup&gt;a&lt;/sup&gt;</td>
<td>43.0 ± 0.6&lt;sup&gt;a&lt;/sup&gt;</td>
<td>43.5 ± 0.5&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>CN52</td>
<td>17.2 ± 0.4&lt;sup&gt;a&lt;/sup&gt;</td>
<td>16.3 ± 0.9&lt;sup&gt;a&lt;/sup&gt;</td>
<td>15.6 ± 0.2&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

Abbreviations: CA, calf; YB, young bull; CO, cow. Data (means ± SD) are expressed in g TGs/100 g of fat (%); results of the analysis of variance by Tukey’s test are shown: p < 0.05, lowercase letters on the same row show significantly different values within each by-product for the three animals; capital letters in the same row show significantly different mean values within each individual animal for the by-products.
bone marrow of the three different bovine categories (Table 4). Two phospholipids, phosphatidylcholine (PC) and sphingomyelin (SM), were detected in all the samples. According to literature \(^{15,36}\), PC was the major phospholipid in all the three animals, accounting about the 60% of the total phospholipids content, because it is the key building blocks of membrane bilayers. In all samples, SM was present in lower amount compared to PC but it plays an important role in animals because it is a substitute for PC as a building block of membranes.

Besides, CA’s bone marrow showed the highest content both in total and individual phospholipids compared to YB and CO samples. This result can be expected since CA is a young growing animal and its cell membranes need more phospholipids, critical to cells’ ability to function and grow \(^{36}\).

4 Conclusion

Because of the lack of literature on the lipid quality of meat by-products, this study represents a first screening of bioactive lipids in different wastes produced by bovine meat industry. The obtained results underline that subcutaneous fat is a non-homogeneous waste with a highly variable composition. Nevertheless, as expected, subcutaneous fat showed the highest cholesterol content compared to the other by-products, because cholesterol is involved in the structure of the membranes that are present where the storage fat is low. On the other hand, the storage fat and especially the bone marrow presented a high fat amount and they were characterized by a low cholesterol content and important concentration of some essential fatty acids (linoleic acid, \(\alpha\) and \(\gamma\) linolenic acid, eicosadienoic acid and arachidonic acid). In particular, calf showed the maximum content of oleic acid, linoleic acid and phospholipids, but also the highest content of cholesterol in the subcutaneous fat rather than young bull and cow. Conversely, young bull presented the lowest cholesterol amount and cow the highest TG content, especially CN48 class.

According to this first evaluation, it could be assumed a future application of these bovine by-products for other purposes. Appropriate research and development activity can help to convert these animal by-products in bioactive components for nutritional properties and other non-food applications like pharmaceuticals, cosmetics or energy.

In particular, a next step of this study will be a crystallization of the lipid fraction of by-products in order to separate and increase the percentage of some interesting bioactive compounds.

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The authors declare no conflict of interest.

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