4-2009

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**Abstract**

Fiber type and nutrient compositions from *Biceps femoris* (Bf), *Longissimus dorsi* (Ld) and *Supraspinatus* (Ss) muscles of 10 Barrosã calves were determined and the respective relationships investigated by principal component analysis (PCA). The Ss muscle had higher proportion of type IIA fibers than Bf ($P < 0.001$) and Ld ($P < 0.01$), higher percentage of oxidative fibers than Ld muscle ($P < 0.01$) and lower proportion of IIB fibers than Bf ($P < 0.001$) and Ld ($P < 0.01$). The latter muscle had higher intramuscular total lipids (ITL) ($P < 0.01$), neutral lipids (NL) ($P < 0.01$), C18:2*c9t11*-CLA ($P < 0.001$), saturated fatty acids (SFA) ($P < 0.001$), *trans* fatty acids (TFA) ($P < 0.001$) and monounsaturated fatty acids (MUFA) ($P < 0.01$) contents than its counterparts. In contrast, Bf and Ss muscles had higher proportions of polyunsaturated fatty acids (PUFA) ($P < 0.001$), *n-6* PUFA ($P < 0.001$), *n-3* PUFA ($P < 0.01$), polyunsaturated/saturated fatty acids ratio (P/S) ($P < 0.001$) and hypocholesterolaemic/hypercholesterolaemic fatty acid ratio (h/H) ($P < 0.001$) than Ld muscle. The higher iron ($P < 0.001$) and phospholipids (PL) ($P < 0.05$) contents were found in Ss muscle. Cholesterol amount was unaffected by anatomical location and ranged from 0.50 to 0.54 mg/g. Barrosã muscles were not well distinguishable by PCA since, based on fiber and nutritional compositions, they did not clearly present different patterns. Barrosã-PDO veal is a low fat meat providing 367–558 mg of *n-6* PUFA and 151-172 mg of *n-3* PUFA of which 67–78 mg are EPA + DHA (based on 150 g consumption). The *n-6/n-3* ratio was close to the recommended values for human diet and the p/s index was below those values.

**Keywords:** Barrosã-PDO veal, fiber type, nutrient composition

## 1. Introduction

Fatty acid (FA) composition influences the nutritive value and sensory characteristics of meat. Since the dietary levels of fat and cholesterol have been related to an increased risk of several chronic diseases, health organizations recommend reductions in total fat as well as particularly saturated fatty acids (SFA), and the increase in ratios of polyunsaturated fatty acids (PUFA) to SFA and *n-3/n-6* intake (British Department of Health, 1994). In Western countries, meat has a negative public image and its contribution to the amount of fat, SFA and cholesterol in human diet is well established (Valsta et al., 2005).

However, meat is one of the few dietary sources of long chain (≥ C20) *n-3* FA (Ponnampalam, 2001), but this is frequently less widely recognized. Recent research has focused on the nutritional relevance of *trans* fatty acids (TFA) and conjugated linoleic acid (CLA) in the human diet. Some CLA isomers are considered beneficial to human health, due to its anticarcinogenic and antiatherogenic properties. In contrast, metabolic studies have shown that some TFA have adverse effects on blood lipid levels (Mulvihill, 2001).

Portuguese population is more diet conscious today, having special concerns about fat and cholesterol associated with meat consumption. For the modern consumer, taste and nu-

tritional value are important quality attributes of meat (Webb and O'Neill, 2008), with increased expectations in Protected Denomination of Origin (PDO) meat. The consumer's decision to purchase a more expensive PDO food is mainly guided by its high intrinsic quality and healthiness, which in the case of meat is largely related to its fat properties and to the absence of chemical residues. Barrosã-PDO meat is perceived as healthy due to the friendly extensive system used on animal production. Traditional Barrosã veal, produced according to PDO rules (Council Regulation n° 2081/92 of 14/7, EEC), has been based on small family herds and involves indoor management, maternal suckling, and complementary diet based on herd products. Being in the past (XIX century) an important exportation issue to the U.K., there are still more than 2000 Barrosã beef producers nowadays, generally owning only 2–5 dams in Minho and 10–15 dams in Barroso, located in the Norwest of Portugal, totalizing about 7500 animals.

Although the importance of Barrosã-PDO veal production on the regional economy is well recognized, being one of the two most important commercial Portuguese veal-PDO (266 carcass tons in 2003; Instituto do Desenvolvimento Rural e Hidráulica, 2003), no published information is available concerning the effect of muscle type on Barrosã-PDO meat composition. Therefore, the aim of this study was to report the nutrient content and fiber profile in *Biceps femoris* (Bf), *Longissimus dorsi* (Ld) and *Supraspinatus* (Ss) muscles and to identify possible interrelationships among nutrients and fiber characteristics in Barrosã PDO-meat.

## 2. Materials and methods

### 2.1. Animals and meat samples

Ten purebred Barrosã calves ($7.2 \pm 0.9$ months of age and $94.4 \pm 14.1$ kg of carcass weight), raised on a traditional production pasture-based system according to PDO specifications, were used in this study. Dams were maintained on pasture during most of the year and the calves were housed suckling mother's milk until weaning (6–7 months of age). Supplementation with cereal grains and forages (hay, straw, ryegrass and maize) obtained on the farms were provided to calves and dams, during periods of feed scarcity. The calves were slaughtered and dressed in an officially approved slaughterhouse. During the first hour *post-mortem*, samples were taken from the middle superficial layer of the *Longissimus dorsi* (L4–L6), and from the distal region of *Biceps femoris* and *Supraspinatus* muscles for determination of histological properties. These samples were immediately frozen by immersion into isopentane, cooled by liquid nitrogen, and kept at −80 °C until further analysis. For lipid, cholesterol and haem iron evaluations, samples with about 200 g were excised from Ld (L4–L6) and from the distal region of Bf and Ss muscles 1 day after slaughter and stored under vacuum at −20 °C, until analysis. Before chemical analyses, the samples were trimmed of visible adipose and connective tissues and minced. Investigated muscles were chosen due to their divergent growth patterns and functionalities *in vivo* and because they represent meat cuts of different expected eating quality and economical value (round, loin and chuck).

### 2.2. Histological traits

For determination of histological properties, serial transverse muscle sections (10 µm thick) were cut in a cryostat at − 24 °C and stained for myosin adenosine triphosphatase (ATPase) at pH 4.45 and for succinic dehydrogenase (SDH). Fibers identified according to ATPase were classified as type I, type IIA and type IIB, according to Brooke and Kaiser (1970). Type IIC were also identified but did not exceed 5% of the total fiber population in any section and were not included in the results. The succinic dehydrogenase (SDH) protocol described by Sheehan and Hrapchak (1987) was used to define fiber metabolic types as oxidative or glycolytic. The percentage of each fiber type and its mean cross-sectional area (CSA) was measured in two randomly selected areas on serial sections with an image analysis software program (Vector Works, 2003). On average, 200 fibers were analyzed on each serial section.

### 2.3. Fatty acid analysis

Intramuscular total lipids (ITL) from duplicate 20 g muscle samples, trimmed of visible adipose and connective tissues were extracted as described by Folch et al. (1957). Lipid extracts were esterified with KOH (2 N) in methanol (ISO 5509, 2000) and resulting fatty acid (FA) methyl esters were analyzed by gas-liquid chromatography, using a HRGC 5160, Mega series from Carlo Erba instruments, equipped with a flame ionization detector and a 60 m long DB 23 capillary column. The oven temperature was raised from 70 to 195 °C (10 min), and then to 220 °C (60 min) at a rate of 5 °C/min for FA analysis. Injector and detector temperatures were 220 °C and 280 °C, respectively.

Identification of FA was based on comparison of retention times of the individual FA with standard mixtures (Supelco and Nuchek GLC reference standard FAME mixture) and confirmed using a GC-MS (Saturn 2200, Varian, Walnut Creek, CA, USA) equipped with a fused silica column (BPX-70 120 m × 0.25 mm × 0.25 µm, Scientific Instrument Services, Inc.). The FA were expressed as weight percentage.

### 2.4. Cholesterol analysis

Cholesterol was evaluated according to the procedure described by Costa et al. (2006b), using a HPLC apparatus (Spectra-Physics Model Spectra 100) equipped with variable wavelength UV detector set at 206 nm and a Spherisorb S5W silica cartridge, 5 µm, 4.0 × 125 mm (Waters PSS 845549). The mobile phase was hexane/isopropanol (97:3) at a flow rate of 1.0 mL/min. The cholesterol content was expressed as g kg$^{-1}$ of muscle.

### 2.5. Haem iron content

Haem iron was quantified following the analytical procedure of Lombardi-Boccia et al. (2002). Briefly, samples were analyzed for haem pigment content according to Hornsey (1956) and the haem iron concentration was calculated from the standard curve as follows: haem iron (mg/100 g) = hematin content (mg/100 g) × AW/MW, where AW is the atomic weight of iron and MW the molecular weight of hematin.

**Table 1.** Muscle fiber composition in *Biceps femoris* (Bf), *Longissimus dorsi* (Ld) and *Supra spinatus* (Ss) muscles from Barrosã calves (*n* = 10).

| Trait | Muscle | | | SE | Effect |
|---|---|---|---|---|---|
| | Bf | Ld | Ss | | |
| *Fiber type distribution (%)* | | | | | |
| I | 22.35 | 22.08 | 22.79 | 1.44 | ns |
| IIA | 17.77[c] | 22.27[b] | 28.48[a] | 1.49 | *** |
| IIB | 59.87[a] | 54.58[b] | 48.73[c] | 1.90 | *** |
| Oxidative | 51.96[ab] | 48.69[b] | 54.13[a] | 1.31 | * |
| *Fiber cross-sectional area-CSA ($\mu m^2$)* | | | | | |
| I | 997.00[b] | 1281.28[a] | 1400.63[a] | 106.29 | ** |
| IIA | 1592.20 | 1845.81 | 1607.93 | 120.14 | ns |
| IIB | 2227.40[b] | 2521.21[a] | 1789.74[c] | 125.28 | *** |
| Glycolytic | 2679.03[a] | 2697.35[a] | 1912.86[b] | 153.82 | *** |
| Oxidative | 1347.17 | 1450.61 | 1465.99 | 95.02 | ns |
| *Relative area of the muscle-RA (%)* | | | | | |
| I | 12.03[b] | 13.69[b] | 19.40[a] | 1.39 | *** |
| IIA | 16.04[c] | 20.50[b] | 27.68[a] | 1.95 | *** |
| IIB | 71.93[a] | 65.92[b] | 52.91[c] | 2.41 | *** |
| Oxidative | 35.53[b] | 33.69[b] | 47.66[a] | 1.82 | *** |

ns = not statistically significant; * $P < 0.05$ ; ** $P < 0.01$ ; *** $P < 0.001$.
Means in the same row with different superscripts are different ($P < 0.05$).
SE = standard error.

## 2.6. Statistical analysis

The data were analyzed using the MIXED procedure of SAS 9.1.3 (2004), considering the animal within the group as subject and the muscle as repeated measures. Analysis of means was performed by the LSD test for 95% probability. Pearson correlation coefficients were generated between the variables studied. A principal component analysis (PCA) of meat composition was carried out using STATISTICA software (Stat-Soft Inc., 2004), with the variables for PCA being standardized to a mean of zero and variance of one. The PCA has been used before with sensory and nutritional data to describe meat quality (Galbraith et al., 2006). Most applications of chemical data analysis are by nature multivariate and the most suitable method for such cases is PCA (Sharaf et al., 1986). PCA is based on the correlation among variables mapping samples through scores and variables by the loadings in a new space defined by the principal components (PC). The PC are simple linear combination of original variables. Score plots allow sample identification, checking if they are similar or dissimilar, typical or outlier. The most important variables are identified from the loadings. An important point in such evaluation is that maximum variability in the data can be explained in a reduced variable set. The first principal component, PC1, is defined in the direction of maximum variance in the data set, and the subsequent components are orthogonal (uncorrelated) to one another and maximize the remaining variance.

## 3. Results

### 3.1. Fiber type characteristics

Fiber type distribution, their cross-sectional area (CSA) and the relative area of the muscle occupied by each fiber type (RA) in Bf, Ld and Ss muscles are summarized in Table 1. The Ss muscle had higher proportion of type IIA fibers than Bf

**Table 2.** Intramuscular total lipids (ITL, g kg$^{-1}$), neutral lipids (NL, g kg$^{-1}$), phospholipids (PL, g kg$^{-1}$), cholesterol (g kg$^{-1}$) and haemic iron (mg kg$^{-1}$) and main fatty acid contents (w/w %) in *Biceps femoris* (Bf), *Longissimus dorsi* (Ld) and *Supra spinatus* (Ss) muscles from Barrosã calves (*n* = 10).

| | Muscle | | | SE | Effect |
|---|---|---|---|---|---|
| | Bf | Ld | Ss | | |
| *Muscle trait* | | | | | |
| ITL | 16.3[c] | 27.8[a] | 21.6[b] | 1.7 | *** |
| NL | 9.8[c] | 22.3[a] | 13.5[b] | 1.7 | *** |
| PL | 6.7[b] | 6.6[b] | 7.2[a] | 0.2 | * |
| Cholesterol | 0.54 | 0.50 | 0.54 | 0.03 | ns |
| Haem iron | 13.0[b] | 12.9[b] | 15.7[a] | 0.8 | *** |
| *ITL composition* | | | | | |
| CLA | 0.66[b] | 0.79[a] | 0.66[b] | 0.04 | *** |
| SFA | 38.91[b] | 45.22[a] | 39.17[b] | 0.80 | *** |
| TFA | 1.41[b] | 1.98[a] | 1.48[b] | 0.10 | *** |
| MUFA | 38.92[b] | 42.53[a] | 36.48[c] | 0.85 | *** |
| PUFA | 21.87[a] | 15.13[b] | 20.33[a] | 0.93 | *** |
| *n-6* PUFA | 15.02[a] | 10.42[b] | 14.32[a] | 0.67 | *** |
| *n-3* PUFA | 6.19[a] | 3.94[b] | 5.32[a] | 0.34 | *** |
| P/S | 0.37[a] | 0.26[b] | 0.35[a] | 0.02 | *** |
| *n-6/n-3* | 3.09 | 3.55 | 3.13 | 0.21 | ns |
| h/H | 2.27[a] | 1.93[b] | 2.28[a] | 0.08 | *** |

CLA = C18:2*c9t11*; SFA = saturated fatty acids; TFA = *trans* fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids.
*n-3* PUFA = C18:3*n-3* + C20:3*n-3* + C20:5*n-3* + C22:5*n-3* + C22:6*n-3*.
*n-6* PUFA = C18:2*n-6* + C18:3*n-6* + C20:3*n-6* + C20:3*n-6* + C20:4*n-6* + C22:2*n-6* + C22:4*n-6*.
P/S = (C18:2*n-6* + C18:3*n-3*)/(C14:0 + C16:0 + C18:0).
h/H = hypocholesterolaemic/hypercholesterolaemic ratio = (C18:1*c9* + C18:2*n-6* + C18:3*n-6* + C18:3*n-3* + C20:2*n-6* + C20:3*n-6* + C20:3*n-3* + C20:4*n-6* + C20:5*n-3* + C22:2*n-6* + C22:4*n-6* + C22:5*n-3* + C22:6*n-3*)/(C12:0 + C14:0 + C16:0)].
ns = not statistically significant, * $P < 0.05$ ; ** $P < 0.01$ ; *** $P < 0.001$
Means in the same row with different superscripts are different ($P < 0.05$).
SE = standard error.

($P < 0.001$) and Ld ($P < 0.01$), higher proportion of oxidative fibers than Ld muscle ($P < 0.01$) and lower proportion of IIB fibers than Bf ($P < 0.001$) and Ld ($P < 0.01$). The CSA of type I fibers was higher ($P < 0.01$) in Ld and Ss muscles than in Bf, while the CSA of type IIB fibers was higher in the Ld muscle than in Bf ($P < 0.05$) and Ss muscles ($P < 0.001$). The latter had glycolytic fibers with lower CSA ($P < 0.001$) and higher RA occupied by type I ($P < 0.01$–$0.001$), IIA ($P < 0.001$) and oxidative fibers ($P < 0.001$) than Bf and Ld muscles.

### 3.2. Nutritional composition

Intramuscular total lipids (ITL), neutral lipids (NL), phospholipids (PL), cholesterol and haemic iron contents and FA composition and nutritional relevant FA ratios in Bf, Ld and Ss muscles from Barrosã calves are presented in Table 2.

The Ld muscle had higher ITL ($P < 0.01$) and NL ($P < 0.01$) contents than Bf and Ss muscles. The latter had higher ITL ($P < 0.01$) and NL ($P < 0.05$) amounts than Bf muscle and higher haemic iron ($P < 0.001$) and PL ($P < 0.05$) contents than Bf and Ld muscles. Cholesterol levels were unaffected by anatomical location and ranged from 0.50 to 0.54 g kg$^{-1}$.

The Ld muscle had higher levels of C18:2*c9t11*-CLA ($P < 0.001$), SFA ($P < 0.001$), TFA ($P < 0.001$) and MUFA ($P < 0.01$) than its counterparts (Table 2). Conversely, Bf and
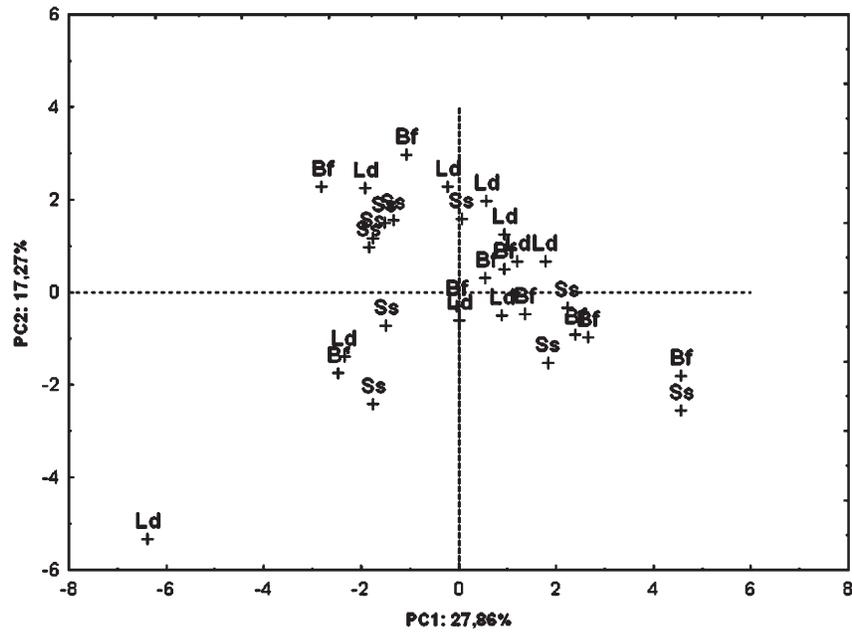
**Figure 1.** Projection of the data of the *Biceps femoris* (Bf), *Longissimus dorsi* (Ld) and *Supra spinatus* (Ss) muscles defined by the two first principal components.

Ss muscles had higher proportions of PUFA ($P < 0.001$), *n-6* PUFA ($P < 0.001$), *n-3* PUFA ($P < 0.01$), polyunsaturated/saturated ratio (P/S) ($P < 0.001$) and hypocholesterolaemic/hyper-cholesterolaemic ratio (h/H) ($P < 0.001$) than Ld muscle. Despite those differences, the *n-6/n-3* ratio was similar among muscles.
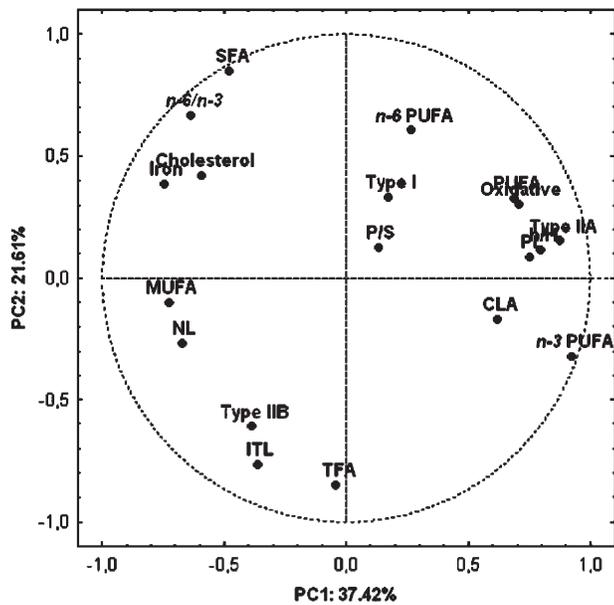


**Figure 2.** Projection of the variables in the plane defined by the two first principal components for *Biceps femoris* (Bf) muscle. Abbreviations: ITL = intramuscular total lipids; NL = neutral lipids; PL = polar lipids; SFA = saturated fatty acids; TFA = trans fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids; CLA = C18:2*c9t11*; *n-3* PUFA = C18:3*n-3* + C20:3*n-3* + C20:5*n-3* + C22:5*n-3* + C22:6*n-3*; *n-6* PUFA = C18:2*n-6* + C18:3*n-6* + C20:2*n-6* + C20:3*n-6* + C20:4*n-6* + C22:2*n-6* + C22:4*n-6*; P/S = (C18:2*n-6* + C18:3*n-3*)/(C14:0 + C16:0 + C18:0); h/H = hypocholesterolaemic/hypercholesterolaemic ratio = [(sum of C18:1c9, C18:2*n-6*, C18:3*n-6*, C18:3*n-3*, C20:2*n-6*, C20:3*n-6*, C20:3*n-3*, C20:4*n-6*, C20:5*n-3*, C22:2*n-6*, C22:4*n-6*, C22:5*n-3* and C22:6*n-3*)/(sum of C12:0, C14:0 and C16:0)].



**Figure 3.** Projection of the variables in the plane defined by the two first principal components for *Longissimus dorsi* (Ld) muscle. Abbreviations: ITL = intramuscular total lipids; NL = neutral lipids; PL = polar lipids SFA = saturated fatty acids; TFA = trans fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids; CLA = C18:2*c9t11*; *n-3* PUFA = C18:3*n-3* + C20:3*n-3* + C20:5*n-3* + C22:5*n-3* + C22:6*n-3*; *n-6* PUFA = C18:2*n-6* + C18:3*n-6* + C20:2*n-6* + C20:3*n-6* + C20:4*n-6* + C22:2*n-6* + C22:4*n-6*; P/S = (C18:2*n-6* + C18:3*n-3*)/(C14:0 + C16:0 + C18:0); h/H = hypocholesterolaemic/hypercholesterolaemic ratio = [(sum of C18:1c9, C18:2*n-6*, C18:3*n-6*, C18:3*n-3*, C20:2*n-6*, C20:3*n-6*, C20:3*n-3*, C20:4*n-6*, C20:5*n-3*, C22:2*n-6*, C22:4*n-6*, C22:5*n-3* and C22:6*n-3*)/(sum of C12:0, C14:0 and C16:0)].
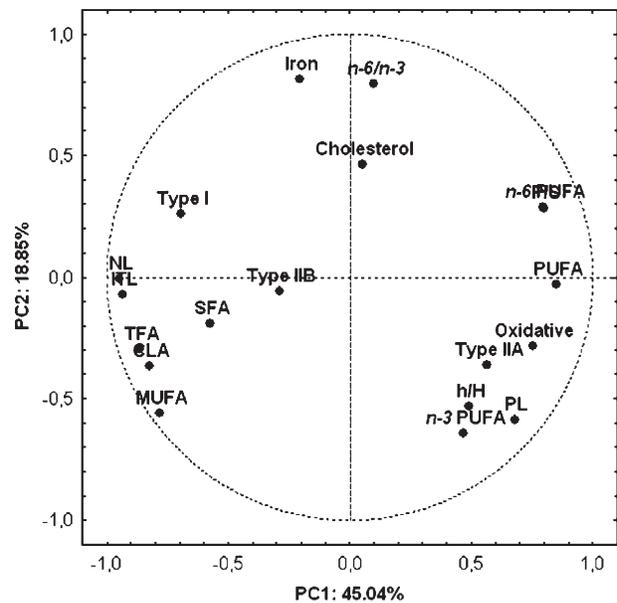
*3.3. Principal component analysis and correlations between traits*

The results of the PCA are presented in Figures 1–4 and Table 3. Figure 1 shows the projected data of the Bf, Ld and Ss muscles in the plane defined by the first two PC and does not exhibit separated groups of points. The PC1 in Bf muscle explained 37% of the total variation (Figure 2) and was mainly characterized by PUFA, *n-3* PUFA, h/H, PL, oxidative and IIA fibers on the right side and haemic iron and MUFA on the left side. The PC2 explained more than 21% of the variation and was defined by SFA and in the opposite direction by ITL and TFA.

Regarding Ld muscle (Figure 3), PC1 and PC2 explained 45% and 19% of the variance, respectively. Two groups of variables were clearly distinguished lying on the PC1, far from the origin. The first group included PUFA, *n-6* PUFA, P/S and oxidative fibers in the right side of the plot. The other group included ITL, NL, CLA, TFA and MUFA components, lying in the left hand side of the plot. The PC2 was mainly affected by iron and *n-6/n-3* and by *n-3* located in the opposite direction.

The PC1 accounted for 33% of the total variation of the nutrient and fiber compositions in Ss muscle. This component was mainly influenced by polyunsaturated fatty acids (PUFA, *n-6* PUFA and *n-3* PUFA), CLA and h/H on the right side and MUFA and *n-6/n-3* on the left side of the plot. The PC2 in Ss muscle explained 24% of the variance and was influenced by NL and iron and by a group of components (PL, cholesterol
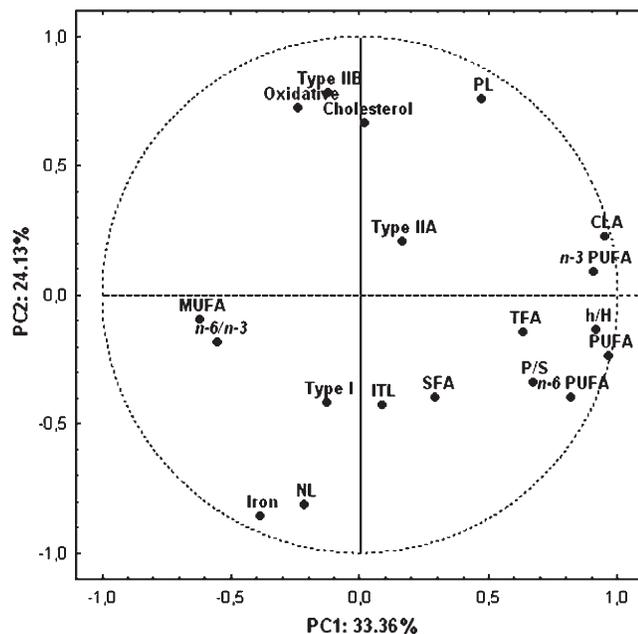
and type IIB and oxidative fibers) located in the opposite extreme of the plot.

Pearson correlation coefficients between nutrients and fiber profiles in muscles Bf, Ld and Ss are shown in Tables 4–6, respectively. Regarding Bf muscle, type I fibers were negatively correlated with NL ($r = -0.63$, $P < 0.05$). The proportion of type IIA fibers was positively correlated with PL ($r = 0.64$, $P < 0.05$), oxidative fibers ($r = 0.63$, $P < 0.05$), C18:2*c9t11*-CLA ($r = 0.65$, $P < 0.05$), *n-3* PUFA ($r = 0.83$, $P < 0.01$) and h/H ($r = 0.68$, $P < 0.05$). Negative correlations were observed between IIA fibers and the proportions of NL ($r = -0.63$, $P < 0.05$) and MUFA ($r = -0.63$, $P < 0.05$). The ITL ($r = 0.64$, $P < 0.05$) was positively correlated with type IIB fibers. The proportion of oxidative fibers was significantly correlated with the amounts of NL ($r = -0.70$, $P < 0.05$) and PL ($r = 0.63$, $P < 0.05$).

Significant correlations were observed between type I fibers in Ld muscle and the levels of ITL ($r = 0.69$, $P < 0.05$), NL ($r = 0.69$, $P < 0.05$) and PL ($r = -0.63$, $P < 0.05$) and the proportion of IIA fibers ($r = -0.93$, $P < 0.001$). The percentage of the latter fibers was correlated with PL ($r = 0.63$, $P < 0.05$). The oxidative fibers in Ld muscle were positively correlated with PL ($r = 0.70$, $P < 0.05$) and negatively correlated with the levels of ITL ($r = -0.82$, $P < 0.01$) and NL ($r = -0.85$, $P < 0.01$).

Regarding Ss muscle, the PL content was negatively correlated with type I fibers ($r = -0.66$, $P < 0.05$) and positively correlated with oxidative fibers ($r = 0.66$, $P < 0.05$) and IIB



**Figure 4.** Projection of the variables in the plane defined by the two first principal components for Supraspinatus (Ss) muscle. Abbreviations: ITL = intramuscular total lipids; NL = neutral lipids; PL = polar lipids SFA = saturated fatty acids; TFA = trans fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids; CLA = C18:2c9t11 n-3 PUFA = C18:3*n-3* + C20:3*n-3* + C20:5*n-3* + C22:5*n-3* + C22:6*n-3*; *n-6* PUFA = C18:2*n-6* + C18:3*n-6* + C20:2*n-6* + C20:3*n-6* + C20:4*n-6* + C22:2*n-6* + C22:4*n-6*; P/S = (C18:2*n-6* + C18:3*n-3*)/(C14:0 + C16:0 + C18:0); h/H = hypocholesterolaemic/hypercholesterolaemic ratio = [(sum of C18:1c9, C18:2*n-6*, C18:3*n-6*, C18:3*n-3*, C20:2*n-6*, C20:3*n-6*, C20:3*n-3*, C20:4*n-6*, C20:5*n-3*, C22:2*n-6*, C22:4*n-6*, C22:5*n-3* and C22:6*n-3*)/(sum of C12:0, C14:0 and C16:0)].

**Table 3.** Coefficients of the loadings (eigenvectors) for the first two principal components (PC).

| Component | Bf ($n = 10$) | | Ld ($n = 10$) | | Ss ($n = 10$) | |
|---|---|---|---|---|---|---|
| | PC1 (%) | PC2 (%) | PC1 (%) | PC2 (%) | PC1 (%) | PC2 (%) |
| ITL | − 0.36 | − **0.77** | − **0.93** | − 0.07 | 0.09 | − 0.43 |
| NL | − **0.67** | − 0.27 | − **0.95** | − 0.01 | − 0.21 | − **0.82** |
| PL | **0.75** | 0.08 | **0.68** | − 0.59 | 0.47 | **0.76** |
| Cholesterol | − **0.59** | 0.42 | 0.05 | 0.47 | 0.02 | **0.66** |
| Iron | − **0.75** | 0.38 | − 0.21 | **0.81** | − 0.39 | − **0.86** |
| Type I | 0.17 | 0.33 | − **0.69** | 0.26 | − 0.13 | − 0.42 |
| Type IIA | **0.88** | 0.15 | **0.58** | − 0.36 | 0.16 | 0.20 |
| Tipe IIB | − 0.39 | − **0.61** | − 0.29 | − 0.06 | − 0.12 | **0.78** |
| Oxidative | **0.71** | 0.30 | **0.75** | − 0.28 | − 0.24 | 0.72 |
| CLA | **0.62** | − 0.17 | − **0.83** | − 0.37 | **0.95** | 0.22 |
| SFA | − 0.48 | **0.85** | − **0.57** | − 0.19 | 0.29 | − 0.40 |
| TFA | − 0.04 | − **0.85** | − **0.86** | − 0.30 | **0.64** | − 0.15 |
| MUFA | − **0.72** | − 0.10 | − **0.78** | − 0.56 | − **0.62** | − 0.10 |
| PUFA | **0.69** | 0.32 | **0.85** | − 0.03 | **0.96** | − 0.24 |
| *n-6* PUFA | 0.27 | **0.60** | **0.80** | 0.29 | **0.82** | − 0.40 |
| *n-3* PUFA | **0.92** | − 0.32 | 0.47 | − **0.64** | **0.91** | 0.09 |
| P/S | 0.13 | 0.12 | **0.80** | 0.29 | **0.67** | − 0.34 |
| *n-6/n-3* | − 0.64 | **0.67** | 0.10 | **0.80** | − 0.55 | − 0.19 |
| h/H | **0.80** | 0.12 | 0.49 | − 0.53 | **0.92** | − 0.13 |

CLA = C18:2*c9t11*; SFA = saturated fatty acids; TFA = trans fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids.
*n-3* PUFA = C18:3*n-3* + C20:3*n-3* + C20:5*n-3* + C22:5*n-3* + C22:6*n-3*.
*n-6* PUFA = C18:2*n-6* + C18:3*n-6* + C20:2*n-6* + C20:3*n-6* + C20:4*n-6* + C22:2*n-6* + C22:4*n-6*; P/S = (C18:2*n-6* + C18:3*n-3*)/(C14:0 + C16:0 + C18:0).
P/S = (C18:2*n-6* + C18:3*n-3*)/(C14:0 + C16:0 + C18:0).
h/H = hypocholesterolaemic/hypercholesterolaemic ratio = (C18:1*c9* + C18:2*n-6* + C18:3*n-6* + C18:3*n-3* + C20:2*n-6* + C20:3*n-6* + C20:3*n-3* + C20:4*n-6* + C20:5*n-3* + C22:2*n-6* + C22:4*n-6* + C22:5*n-3* + C22:6*n-3*)/(C12:0 + C14:0 + C16:0)].
The most significant loadings are highlighted in **boldface**.

**Table 4.** Pearson correlations among fiber type profile, intramuscular total lipids (ITL), neutral lipids (NL), phospholipids (PL), cholesterol, haemic iron and main fatty acid in *Biceps femoris* (Bf) muscle.

| | ITL | NL | PL | Chole-sterol | Iron | Type I | Type IIA | Type IIB | Oxida-tive | CLA | SFA | TFA | MUFA | PUFA | n-6 PUFA | n-3 PUFA | P/S | n-6/n-3 | h/H |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| ITL | 1.00 | 0.73* | − 0.24 | − 0.38 | 0.19 | − 0.45 | − 0.45 | 0.64* | − 0.52 | − 0.23 | − 0.41 | 0.52 | 0.36 | − 0.29 | − 0.25 | − 0.30 | − 0.09 | − 0.37 | − 0.19 |
| NL | | 1.00 | − 0.68* | 0.14 | 0.70* | − 0.63* | − 0.63* | 0.49 | − 0.70* | − 0.35 | 0.19 | 0.19 | 0.42 | − 0.15 | 0.00 | 0.11 | − 0.47 | 0.13 | − 0.22 |
| PL | | | 1.00 | − 0.67* | − 0.55 | 0.39 | 0.64* | − 0.46 | 0.63* | 0.31 | − 0.26 | − 0.31 | − 0.25 | 0.35 | − 0.21 | 0.06 | 0.64* | − 0.47 | 0.52 |
| Cholesterol | | | | 1.00 | 0.51 | − 0.12 | − 0.37 | − 0.12 | − 0.23 | − 0.07 | 0.58 | − 0.09 | 0.36 | − 0.41 | 0.35 | − 0.12 | − 0.63* | 0.72* | − 0.51 |
| Iron | | | | | 1.00 | − 0.37 | − 0.51 | 0.12 | − 0.41 | − 0.43 | 0.76* | − 0.40 | 0.53 | − 0.18 | − 0.00 | 0.24 | − 0.75* | 0.70* | − 0.31 |
| Type I | | | | | | 1.00 | − 0.13 | − 0.29 | 0.37 | − 0.28 | 0.10 | − 0.16 | − 0.09 | 0.03 | − 0.52 | 0.11 | − 0.11 | 0.15 | − 0.05 |
| Type IIA | | | | | | | 1.00 | − 0.42 | 0.63* | 0.65* | − 0.27 | − 0.26 | − 0.66* | 0.59 | 0.42 | 0.21 | 0.83** | − 0.42 | 0.68* |
| Type IIB | | | | | | | | 1.00 | − 0.46 | − 0.14 | − 0.29 | 0.50 | 0.17 | − 0.29 | − 0.33 | − 0.28 | − 0.12 | − 0.01 | − 0.23 |
| Oxidative | | | | | | | | | 1.00 | 0.39 | − 0.17 | − 0.14 | − 0.51 | 0.43 | − 0.08 | 0.21 | 0.52 | − 0.09 | 0.48 |
| CLA | | | | | | | | | | 1.00 | − 0.37 | 0.21 | − 0.20 | 0.32 | 0.49 | − 0.07 | 0.74* | − 0.48 | 0.58 |
| SFA | | | | | | | | | | | 1.00 | − 0.78** | 0.33 | 0.02 | 0.07 | 0.46 | − 0.69* | 0.82** | − 0.18 |
| TFA | | | | | | | | | | | | 1.00 | 0.01 | − 0.31 | − 0.04 | − 0.54 | 0.23 | − 0.47 | − 0.18 |
| MUFA | | | | | | | | | | | | | 1.00 | − 0.72* | − 0.15 | − 0.47 | − 0.65* | 0.31 | − 0.58 |
| PUFA | | | | | | | | | | | | | | 1.00 | 0.12 | 0.88** | 0.58 | − 0.30 | 0.91*** |
| n-6 PUFA | | | | | | | | | | | | | | | 1.00 | − 0.63* | 0.11 | − 0.05 | − 0.21 |
| n-3 PUFA | | | | | | | | | | | | | | | | 1.00 | 0.08 | 0.14 | 0.64* |
| P/S | | | | | | | | | | | | | | | | | 1.00 | − 0.80** | 0.76* |
| n-6/n-3 | | | | | | | | | | | | | | | | | | 1.00 | − 0.44 |
| h/H | | | | | | | | | | | | | | | | | | | 1.00 |

CLA = C18:2c9t11;SFA = saturated fatty acids;TFA = *trans* fatty acids;MUFA = monounsaturated fatty acids;PUFA = polyunsaturated fatty acids;n-3 PUFA = C18:3n-3 + C20:3n-3 + C20:5n-3 + C22:5n-3 + C22:6n-3; n-6 PUFA = C18:2n-6 + C18:3n-6 + C20:2n-6 + C20:3n-6 + C20:4n-6 + C22:2n-6 + C22:4n-6; P/S = (C18:2n-6 + C18:3n-3)/(C14:0 + C16:0 + C18:0); h/H = hypocholesterolaemic/hypercholesterolaemic ratio = [(sum of C18:1c9, C18:2n-6, C18:3n-6, C18:3n-3, C20:3n-3, C20:4n-6, C20:5n-3, C22:2n-6, C22:4n-6, C22:5n-3 and C22:6n-3)/(sum of C12:0, C14:0 and C16:0)]; ns = not statistically significant ; * *P* < 0.05 ; ** *P* < 0.01 ; *** *P* < 0.001.

**Table 5.** Pearson correlations among fiber type profile, intramuscular total lipids (ITL), neutral lipids (NL), phospholipids (PL), cholesterol, haemic iron and main fatty acid in *Longissimus dorsi* (Ld) muscle.

| | ITL | NL | PL | Cholesterol | Iron | Type I | Type IIA | Type IIB | Oxidative | CLA | SFA | TFA | MUFA | PUFA | n-6 PUFA | n-3 PUFA | P/S | n-6/n-3 | h/H |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| ITL | 1.00 | 1.00*** | – 0.55 | 0.02 | 0.17 | 0.69* | – 0.59 | 0.39 | – 0.82** | 0.78** | 0.58 | 0.76* | 0.77* | – 0.67* | – 0.67* | – 0.66* | – 0.30 | – 0.23 | – 0.29 |
| NL | | 1.00 | – 0.63* | 0.06 | 0.23 | 0.69* | – 0.59 | 0.41 | – 0.85** | 0.78** | 0.58 | 0.77* | 0.74* | – 0.68* | – 0.68* | – 0.66* | – 0.35 | – 0.17 | – 0.34 |
| PL | | | 1.00 | – 0.27 | – 0.56 | – 0.63* | 0.63* | – 0.12 | 0.70* | – 0.24 | – 0.02 | – 0.42 | – 0.13 | 0.66* | 0.52 | 0.49 | 0.58 | – 0.30 | 0.63* |
| Cholesterol | | | | 1.00 | 0.76* | – 0.01 | 0.06 | 0.47 | – 0.31 | – 0.25 | – 0.06 | – 0.11 | – 0.32 | 0.25 | 0.16 | 0.22 | 0.16 | 0.03 | – 0.09 |
| Iron | | | | | 1.00 | 0.29 | – 0.25 | 0.23 | – 0.40 | – 0.04 | 0.16 | 0.00 | – 0.23 | – 0.07 | 0.15 | 0.15 | – 0.50 | 0.51 | – 0.47 |
| Type I | | | | | | 1.00 | – 0.93*** | – 0.26 | – 0.55 | 0.40 | 0.26 | 0.44 | 0.45 | – 0.53 | – 0.39 | – 0.42 | – 0.43 | 0.06 | – 0.18 |
| Type IIA | | | | | | | 1.00 | 0.34 | 0.59 | – 0.21 | – 0.11 | – 0.29 | – 0.26 | 0.42 | 0.21 | 0.27 | 0.46 | – 0.12 | 0.14 |
| Type IIB | | | | | | | | 1.00 | – 0.54 | 0.30 | 0.34 | 0.32 | 0.15 | – 0.05 | – 0.29 | – 0.14 | 0.15 | – 0.12 | – 0.21 |
| Oxidative | | | | | | | | | 1.00 | – 0.45 | – 0.42 | – 0.59 | – 0.35 | 0.44 | 0.43 | 0.37 | 0.34 | – 0.04 | 0.33 |
| CLA | | | | | | | | | | 1.00 | 0.84** | 0.91*** | 0.93*** | – 0.62 | – 0.65* | – 0.63 | – 0.30 | – 0.21 | – 0.19 |
| SFA | | | | | | | | | | | 1.00 | 0.73* | 0.69* | – 0.24 | – 0.21 | – 0.18 | – 0.32 | 0.03 | – 0.05 |
| TFA | | | | | | | | | | | | 1.00 | 0.86** | – 0.66* | – 0.74* | – 0.70* | – 0.23 | – 0.25 | – 0.22 |
| MUFA | | | | | | | | | | | | | 1.00 | – 0.61 | – 0.69* | – 0.72* | – 0.08 | – 0.49 | – 0.03 |
| PUFA | | | | | | | | | | | | | | 1.00 | 0.86** | 0.92*** | 0.54 | 0.03 | 0.73* |
| n-6 PUFA | | | | | | | | | | | | | | | 1.00 | 0.96*** | 0.14 | 0.34 | 0.12 |
| n-3 PUFA | | | | | | | | | | | | | | | | 1.00 | 0.17 | 0.39 | 0.50 |
| P/S | | | | | | | | | | | | | | | | | 1.00 | – 0.74* | 0.72* |
| n-6/n-3 | | | | | | | | | | | | | | | | | | 1.00 | – 0.59 |
| h/H | | | | | | | | | | | | | | | | | | | 1.00 |

CLA = C18:2c9t11; SFA = saturated fatty acids; TFA = *trans* fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids; n-3 PUFA = C18:3n-3 + C20:3n-3 + C20:5n-3 + C22:5n-3 + C22:6n-3; n-6 PUFA = C18:2n-6 + C18:3n-6 + C20:2n-6 + C20:3n-6 + C20:4n-6 + C22:2n-6 + C22:4n-6; P/S = (C18:2n-6 + C18:3n-3)/(C14:0 + C16:0 + C18:0); h/H = hypocholesterolaemic/hypercholesterolaemic ratio = [(sum of C18:1c9.C18:2n-6.C18:3n-6.C18:3n-3.C20:2n-6.C20:3n-6.C20:3n-3.C20:4n-6.C20:5n-3.C22:2n-6.C22:4n-6.C22:5n-3 and C22:6n-3)/(sum of C12:0.C14:0 and C16:0)]; ns = not statistically significant ; * $P < 0.05$ ; ** $P < 0.01$ ; *** $P < 0.001$

**Table 6.** Pearson correlations among fiber type profile, intramuscular total lipids (ITL), neutral lipids (NL), phospholipids (PL), cholesterol, haemic iron and main fatty acid in *Supraspinatus* (Ss) muscle.

| | ITL | NL | PL | Cholesterol | Iron | Type I | Type IIA | Type IIB | Oxidative | CLA | SFA | TFA | MUFA | PUFA | n-6 PUFA | n-3 PUFA | P/S | n-6/n-3 | h/H |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| ITL | 1.00 | 0.78** | − 0.04 | − 0.18 | 0.38 | − 0.17 | 0.30 | 0.02 | 0.00 | − 0.02 | 0.53 | − 0.19 | 0.03 | 0.22 | 0.24 | 0.37 | − 0.05 | 0.30 | 0.28 |
| NL | | 1.00 | − 0.55 | − 0.51 | 0.86** | 0.15 | − 0.08 | − 0.38 | − 0.37 | − 0.37 | 0.45 | − 0.24 | 0.25 | 0.01 | 0.09 | 0.21 | − 0.28 | 0.36 | 0.02 |
| PL | | | 1.00 | 0.43 | − 0.75* | − 0.66* | 0.38 | 0.70* | 0.66* | 0.58 | − 0.03 | − 0.04 | − 0.54 | 0.31 | 0.17 | 0.21 | 0.39 | − 0.22 | 0.31 |
| Cholesterol | | | | 1.00 | − 0.75* | 0.00 | − 0.32 | 0.55 | 0.40 | 0.21 | 0.08 | 0.09 | 0.29 | − 0.13 | − 0.34 | − 0.29 | 0.16 | − 0.05 | 0.08 |
| Iron | | | | | 1.00 | 0.19 | − 0.15 | − 0.54 | − 0.52 | − 0.56 | 0.20 | − 0.31 | 0.19 | − 0.16 | 0.05 | 0.06 | − 0.45 | 0.34 | − 0.26 |
| Type I | | | | | | 1.00 | − 0.46 | − 0.78*** | − 0.32 | − 0.15 | − 0.11 | 0.66* | 0.47 | − 0.08 | − 0.25 | − 0.18 | 0.07 | − 0.17 | − 0.05 |
| Type IIA | | | | | | | 1.00 | 0.20 | 0.32 | 0.11 | − 0.34 | − 0.20 | − 0.57 | 0.11 | 0.34 | 0.18 | − 0.03 | 0.09 | 0.01 |
| Type IIB | | | | | | | | 1.00 | 0.68* | 0.03 | 0.02 | − 0.55 | − 0.04 | − 0.25 | − 0.20 | − 0.24 | − 0.17 | 0.20 | − 0.14 |
| Oxidative | | | | | | | | | 1.00 | − 0.12 | − 0.13 | − 0.23 | 0.07 | − 0.36 | − 0.33 | − 0.37 | − 0.24 | 0.18 | − 0.35 |
| CLA | | | | | | | | | | 1.00 | 0.14 | 0.67* | − 0.50 | 0.85** | 0.43 | 0.66* | 0.97*** | − 0.71* | 0.90*** |
| SFA | | | | | | | | | | | 1.00 | 0.12 | 0.03 | 0.46 | 0.47 | 0.66* | 0.10 | 0.30 | 0.39 |
| TFA | | | | | | | | | | | | 1.00 | − 0.03 | 0.60 | 0.25 | 0.38 | 0.74* | − 0.59 | 0.60 |
| MUFA | | | | | | | | | | | | | 1.00 | − 0.67* | − 0.75* | − 0.66* | − 0.34 | 0.09 | − 0.34 |
| PUFA | | | | | | | | | | | | | | 1.00 | 0.78*** | 0.93*** | 0.82** | 0.41 | 0.91*** |
| n-6 PUFA | | | | | | | | | | | | | | | 1.00 | 0.92*** | 0.31 | 0.16 | 0.52 |
| n-3 PUFA | | | | | | | | | | | | | | | | 1.00 | 0.54 | − 0.07 | 0.75* |
| P/S | | | | | | | | | | | | | | | | | 1.00 | − 0.82** | 0.90*** |
| n-6/n-3 | | | | | | | | | | | | | | | | | | 1.00 | − 0.56 |
| h/H | | | | | | | | | | | | | | | | | | | 1.00 |

CLA = C18:2c9t11; SFA = saturated fatty acids; TFA = *trans* fatty acids; MUFA = monounsaturated fatty acids; PUFA = polyunsaturated fatty acids; n-3 PUFA = C18:3n-3 + C20:3n-3 + C20:5n-3 + C22:5n-3 + C22:6n-3; n-6 PUFA = C18:2n-6 + C18:3n-6 + C20:2n-6 + C20:3n-6 + C20:4n-6 + C22:2n-6 + C22:4n-6; P/S = (C18:2n-6 + C18:3n-3)/(C14:0 + C16:0 + C18:0); h/H = hypocholesterolaemic/hypercholesterolaemic ratio = [(sum of C18:1c9,C18:2n-6,C18:3n-6,C18:3n-3,C20:2n-6,C20:3n-6,C20:3n-3,C20:4n-6,C20:5n-3,C22:2n-6,C22:4n-6,C22:5n-3 and C22:6n-3)/(sum of C12:0,C14:0 and C16:0)]; ns = not statistically significant ; * $P < 0.05$ ; ** $P < 0.01$ ; *** $P < 0.001$

($r$ = 0.70, $P$ < 0.05). The latter was positively correlated with oxidative fibers ($r$ = 0.68, $P$ < 0.05) and negatively correlated with type I fibers ($r$ = − 0.78, $P$ < 0.01). The proportion of type I fibers was correlated with TFA ($r$ = 0.68, $P$ < 0.05) in Ss muscle.

## 4. Discussion

### 4.1. Nutritional composition and muscle fiber profile

According to fiber proportions, Bf, Ld and Ss can be considered as muscles of intermediate metabolism with the latter presenting a more oxidative prone than Bf, which is more evident when the relative area occupied by each fiber type is considered (Table 1). These results are in good agreement with PCA. In fact, the projection of Bf, Ld and Ss muscles in a plane defined by the first two PC did not show a clear separation of muscles (Figure 1), which points out that, based on the fiber and fat compositions, they did not clearly present different patterns. Comparatively, a study from Costa et al. (2006a) on Ld, Bf and Ss muscles from Maronesa calves of similar age, weight at slaughter and handling, showed higher oxidative metabolism, lower percentages of type IIB and higher proportion of type IIA fibers than Barrosã breed.

Results obtained in pigs suggested that muscle oxidative capacity is primarily related to the presence of type I and IIA muscle fibers but some IIB fibers also exhibit that characteristic (Ruusunen and Puolanne, 1997). The oxidative fibers were positively correlated with IIA fibers in Bf muscle ($P$ < 0.05) and with IIB fibers in Ss muscle ($P$ < 0.05) in Barrosã calves. Interestingly, type I fibers were negatively correlated with oxidative fibers ($P$ > 0.05) in Ld and Ss muscles, suggesting that fibers classified according to mATPase activity could have distinct oxidative enzyme activities among muscles. Muscle fibers location and function are undoubtedly the most important factors that influence fiber type composition within an animal (Lefaucheur and Gerrard, 2000). Contractile and metabolic properties of the same fiber type can vary with anatomical location (Nakatani et al., 2003; Rosser et al., 1992) and within muscle (Armstrong and Laughlin, 1985). Usually, smaller-sized fibers have higher oxidative enzyme activities than larger-sized fibers, even within the same fiber type (Nakatani et al., 2003). The Ss muscle from Barrosã calves had IIB with lower ($P$ < 0.01) CSA than Bf and Ld muscles. In contrast, the lower ($P$ < 0.05) CSA of type I fibers was found in Bf muscle, suggesting that type IIB and I fibers could be more oxidative in Ss and Bf muscles (respectively) than in their counterparts.

It is generally accepted that high proportion of oxidative fibers implies a greater amount of ITL and NL, due to their ability to use FA as an energy source. For this reason they usually contain higher levels of intra-fiber lipids than the glycolytic counterparts (Alasnier et al. 1996). In apparent contradiction, the proportion of oxidative fibers was negatively correlated with NL content in Bf ($P$ < 0.05), Ld ($P$ < 0.01) and Ss ($P$ > 0.05) muscles in the current study. Moreover, the latter was more oxidative ($P$ < 0.01) and had lower ($P$ < 0.01) ITL and NL contents than Ld muscle. Other studies obtained similar results, and concluded that total lipid content has no relationship with the predominant metabolism of the muscle (Andrés et al., 2001; Leseigneur-Meynier and Gandemer, 1991). The lack of positive correlation between the proportion of oxidative fibers and ITL and NL contents found here could be due to the fact

that muscle lipid extract contain not only intramyocellular but also extramyocellular triglycerides (Guo, 2001) and the latter, richer in NL, are more important quantitatively and unrelated to fiber type composition (Essen-Gustavsson et al., 1994). On the contrary, PL content was positively correlated with oxidative fibers ($P$ < 0.05) in Bf, Ld and Ss muscles which could be associated with a higher relative total membrane surface per volume unit, due to the increased number of mitochondria in oxidative fibers.

The PCA showed weak associations between metabolic/contractile traits and nutrient composition, suggesting that some other factors could have more biological relevance on lipid composition in Bf, Ld and Ss muscles than fiber composition. As told before, the myosin ATPase histochemical reaction does not necessarily reflect the metabolic capacity of the fibers. Previous studies showed a spectrum of metabolic capacities within a given fiber type, suggesting a specialization within fiber types of a given muscle (Reichmann and Pette, 1982; Rosser et al., 1992). This lack of relationship between mATPase activity and metabolism and the possible different biochemical characteristics within a given fiber type could have contributed to the results obtained.

### 4.2. Barrosã-PDO meat composition and dietary recommendations for humans

On the basis of a daily consumption of a 150 g steak, trimmed of all visible fat, except for intramuscular fat, Barrosã-PDO meat provides 75–81 mg of cholesterol which represent 25–27% of the maximum daily cholesterol recommendations ( < 300 mg/day) (USDA, 2005) and 22–28% and 13–16% of daily iron recommended requirements for male and female human adults, respectively (Martin et al., 2001).

Optimal dietary intake of CLA remains to be established. However, these are the only natural FA accepted by the National Academy of Sciences of USA as exhibiting consistent antitumor properties at levels as low as 0.25–1.0% of fat (Eynard and Lopez, 2003). The CLA content in Barrosã meat was more than 0.60% of fat, with a 150 g steak providing 16–33 mg of C18:2*c9t11*-CLA.

Based on 11.3 MJ/day for human adults (25–49 years old), the daily recommended intakes include a minimum of 9000 mg of *n-6* PUFA and 1500 mg of *n-3* PUFA (Scientific Review Committee, 1990). Within *n-3* PUFA, current recommendations for intake of EPA (C20:5 *n-3*) + DHA (C22:6 *n-3*) range from 270 to 700 mg/day (Givens and Gibbs, 2008). Barrosã-PDO meat (150 g) provides 367–558 mg of *n-6* PUFA and 151–172 mg of *n-3* PUFA of which 67–78 mg are EPA + DHA.

The P/S ranged from 0.26 to 0.37 and was below the nutrition recommendations (British Department of Health, 1994). However, the use of P/S ratio for nutritional characterization of ruminant fats may be questioned based on the fact that C18:0 might not be a health deleterious FA and ruminant fats have high C18:0 content. An alternative approach might be grouping the FA using a functional (i.e. h/H) rather than the structural criteria (P/S), as proposed by Santos-Silva et al. (2002). Within the PUFA, the *n-6*/*n-3* ratio in the current research was in agreement with previous studies (Dias et al., 2008; Alfaia et al., 2007) (around 3), which was a slightly higher than dietary guidelines (< 2) (Simopoulos, 1999). However, the usefulness of the *n-6*/*n-3* ratio to estimate cardiovascular disease risk of diet fats should be taken with caution (Stanley et al., 2007).

## 5. Conclusions

This research suggests that fiber type and nutrition compositions vary across *Biceps femoris*, *Longissimus dorsi* and *Supraspinatus* muscles but these parameters were not closely related. Based on 150 g consumption, the low fat Barrosã-PDO veal provides 367–558 mg of *n-6* PUFA and 151–172 mg of *n-3* PUFA of which 67–78 mg are EPA + DHA. The *n-6/n-3* ratio was close to the recommended values for human diet and the P/S index was below those values.

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