

Effects of fish oil replacement by vegetable oil blend on digestive enzymes and tissue histomorphology of European sea bass (*Dicentrarchus labrax*) juveniles

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Abstract The impact of replacing circa 70 % fish oil (FO) by a vegetable oil (VO) blend (rapeseed, linseed, palm oils; 20:50:30) in diets for European sea bass juveniles (IBW 96 ± 0.8 g) was evaluated in terms of activities of digestive enzymes (amylase, lipase, alkaline phosphatase, trypsin and total alkaline proteases) in the anterior (AI) and posterior (PI) intestine and tissue morphology (pyloric caeca—PC, AI, PI, distal intestine—DI and liver). For that purpose, fish were fed the experimental diets for 36 days and then liver and intestine were sampled at 2, 6 and 24 h after the last meal. Alkaline protease characterization was also done in AI and PI at 6 h post-feeding. Dietary VO promoted higher alkaline phosphatase activity at 2 h post-feeding

in the AI and at all sampling points in the PI. Total alkaline protease activity was higher at 6 h post-feeding in the PI of fish fed the FO diet. Identical number of bands was observed in zymograms of alkaline proteases of fish fed both diets. No alterations in the histomorphology of PC, AI, PI or DI were noticed in fish fed the VO diets, while in the liver a tendency towards increased hepatocyte vacuolization due to lipid accumulation was observed. Overall, and with the exception of a higher intestine alkaline phosphatase activity, 70 % FO replacement by a VO blend in diets for European sea bass resulted in no distinctive alterations on the postprandial pattern of digestive enzyme activities and intestine histomorphology.

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Introduction

In the aquaculture sector, constrains in fish oil (FO) availability and price dictate an urgent need to replace it by more readily available, cheaper and sustainable alternatives such as vegetable oils (VO). However, a drawback of using VO in aquafeeds is their fatty acid (FA) profile, which has a different n3:n6 ratio comparatively to FO, and lacks n-3 long-chain polyunsaturated fatty acids (LC-PUFA), particularly

eicosapentaenoic (EPA) and docosahexaenoic (DHA) acids, which are essential FA for marine fish. Also, changes in dietary FA composition has been reported to affect FA composition of fish storage lipids and of cell membranes, including that of hepatic (Caballero et al. 2002; Kjaer et al. 2008; Fountoulaki et al. 2009; Castro et al. 2015b) and intestinal (Caballero et al. 2003; Ruyter et al. 2006) cells, the histomorphology of the liver (Caballero et al. 2002; Kjaer et al. 2008; Fountoulaki et al. 2009) and digestive tract (Olsen et al. 1999; Moldal et al. 2014). Dietary FO replacement by a single source or by blends of VOs was shown to induce accumulation of lipid droplets in: hepatocytes of rainbow trout (Caballero et al. 2002), Atlantic salmon (Kjaer et al. 2008) and gilthead sea bream (Fountoulaki et al. 2009), and enterocytes of Arctic char (Olsen et al. 1999), rainbow trout (Caballero et al. 2002; Olsen et al. 2003), gilthead sea bream (Caballero et al. 2003; Santigosa et al. 2011) and pikeperch (Kowalska et al. 2010), and to affect gut integrity and induce tissue damage in Arctic char (Olsen et al. 1999). Additionally, reduction of mucosal fold height in the mid-intestine and reduction of wall thickness in the distal intestine were also reported in Atlantic salmon fed diets with olive, rapeseed or soybean oils (Moldal et al. 2014). These alterations may interfere with nutrient digestion, transport, diffusion and absorption efficiency or function and metabolism (Olsen et al. 1999; Caballero et al. 2002, 2003; Jutfelt et al. 2007; Wassef et al. 2007; Geurden et al. 2009; Santigosa et al. 2011).

Studies assessing the effect of dietary FO substitution by VOs on digestive enzyme activities in fish are scarce (Santigosa et al. 2011; Bowyer et al. 2012; Ribeiro et al. 2015). In yellowtail kingfish, dietary coconut oil promoted a decrease in trypsin and lipase activities but had no effect on amylase activity (Bowyer et al. 2012). In meagre, the dietary incorporation of a VO blend had no marked effects on the intestinal alkaline phosphatase activity (Ribeiro et al. 2015). In gilthead sea bream fed diets including a VO blend, lipase and amylase were unaffected, while total alkaline protease (TAP) activity increased in pyloric caeca and decreased in proximal intestine (Santigosa et al. 2011). Furthermore, zymogram determination of TAP composition revealed higher intensity of trypsin- and chymotrypsin-like proteolytic bands in the pyloric caeca of gilthead sea bream fed VO diet, concurring with the higher TAP activity observed in that tissue (Santigosa et al. 2011).

In fish, digestive enzyme activities were shown to be modulated by diet composition (Munilla-Morán and Saborido-Rey 1996a, b; Hidalgo et al. 1999; Fernández et al. 2001; Fountoulaki et al. 2005; Furné et al. 2005; Santigosa et al. 2008; Pérez-Jiménez et al. 2009; Castro et al. 2013) and by intestinal transit time, which determines the digesta arrival to each intestinal section (Einarsson et al. 1996; Fountoulaki et al. 2005; Caruso et al. 2008; Santigosa et al. 2008; Rodiles et al. 2012). The digesta transit time along the gastrointestinal tract is itself also affected by diet composition (Santigosa et al. 2008; Rodiles et al. 2012). All these conditions may influence the digestive and absorptive processes (Castro et al. 2015a).

European sea bass (*Dicentrarchus labrax*) is an economically important carnivorous fish species of aquaculture in Europe. In this species, data obtained so far demonstrate that FO can be replaced by a single source (soybean oil) or a blend (rapeseed, linseed palm and/or olive oils at different proportions) of VO up to 60 % without compromising growth performance and survival (Figueiredo-Silva et al. 2005; Mourente et al. 2005; Richard et al. 2006b; Castro et al. 2015b). While the impact of such substitutions at metabolic level, mainly in liver, has been assessed (Richard et al. 2006b; Castro et al. 2015b), little information is available on the potential effects in physiological functions of intestine (Castro et al. 2015a). Thus, this study aimed to evaluate the effect of replacing circa 70 % fish oil (FO) by a VO blend on the activities of digestive enzymes of pancreatic (total alkaline proteases, trypsin, amylase and lipase) and brush border membrane (alkaline phosphatase) origin and on the histomorphology of different sections of the digestive tract and liver of European sea bass juveniles.

Materials and methods

Experimental diets

Two isoproteic (45 % crude protein; CP) and isolipidic (18 % crude lipids; CL) diets were formulated to contain 20 % gelatinized maize starch and FO or VO blend as lipid sources (Table 1). The VO blend was composed of rapeseed, linseed and palm oils (20:50:30) and replaced circa 70 % of dietary lipids provided by cod liver oil and fish meal. The diets were similar in saturated fatty acids (SFA) content, but,

comparatively to the VO diet, the FO diet had higher levels of eicosapentaenoic acid (EPA, 20:5n-3), docosahexaenoic acid (DHA, 22:6n-3) and the monounsaturated fatty acids (MUFA) 20:1 and 22:1 and lower levels of linoleic acid (LA, 18:2n-6),

linolenic acid (LNA, 18:3n-3) and oleic acid (OA, 18:1n-9) (Table 2). Details on diet preparation and analysis are given in Castro et al. (2015a).

Animals, experimental conditions and sampling

The experiment was directed by accredited scientists (following FELASA category C recommendations) and conducted according to the European Union

Table 1 Proximate composition of the experimental diets

Diets	FO	VO
<i>Ingredients (% dry weight)</i>		
Fish meal ^a	63.6	63.6
Gelatinized maize starch ^b	20	20
Cod liver oil ^c	12.4	0.4
Vegetable oil blend ^d	–	12.0
Vitamin premix ^e	1.0	1.0
Mineral premix ^f	1.0	1.0
Choline chloride (50 %)	0.5	0.5
Binder ^g	1.0	1.0
Chromium	0.5	0.5
<i>Proximate analyses (% dry matter)</i>		
Dry matter (DM)	87.5	87.2
Crude protein (CP)	46.2	47.1
Crude lipid (CL)	17.9	18.2
Ash	13.9	13.7
Starch	20.4	19.8
Gross energy (MJ kg ⁻¹)	21.4	21.0

^a Steam-dried low-temperature fish meal, Pesquera Diamante, Perú (CP: 70.2 % DM; CL: 8.08 % DM)

^b C-Gel Instant-12018, Cerestar, Mechelen, Belgium

^c Fagron Iberica S.A.U., Spain

^d 30 % palm oil (Colmi, Malaysia), 50 % linseed oil (Sociedade Portuense de Drogas, S.A., Portugal) and 20 % rapeseed oil (Huilerie Emile Noël S.A.S., France)

^e Vitamins (mg kg⁻¹ diet): retinol acetate, 18,000 (IU kg⁻¹ diet); cholecalciferol, 2000 (IU kg⁻¹ diet); alpha-tocopherol acetate, 35; sodium menadione bisulphate, 10; thiamine-HCl, 15; riboflavin, 25; calcium pantothenate, 50; nicotinic acid, 200; pyridoxine HCl, 5; folic acid 10; cyanocobalamin, 0.02; biotin, 1.5; ascorbic acid, 50; inositol, 400. Premix, Viana do Castelo, Portugal

^f Minerals (mg kg⁻¹ diet): cobalt sulphate, 1.91; copper sulphate, 19.6; iron sulphate, 200; sodium fluoride, 2.21; potassium iodide, 0.78; magnesium oxide, 830; manganese oxide, 26; sodium selenite, 0.66; zinc oxide, 37.5; dibasic calcium phosphate, 5.93 (g kg⁻¹ diet); potassium chloride, 1.15 (g kg⁻¹ diet); sodium chloride, 0.40 (g kg⁻¹ diet). Premix, Viana do Castelo, Portugal

^g Aquacube (guar gum, polymethyl carbamide, manioc starch blend, hydrate calcium sulphate). Agil, England

Table 2 Fatty acid composition (% of total fatty acids) of the experimental diets

Diets	FO	VO
14:0	5.3	1.4
15:0	0.4	0.1
16:0	13.7	14.1
17:0	0.3	0.2
18:0	3.0	3.7
20:0	0.1	0.3
Sum saturates (Sat)	22.9	19.9
16:1	7.6	2.1
18:1	18.7	27.0
20:1	7.1	0.8
22:1	5.4	0.2
Sum monoenes	39.0	30.2
16:2 n-4	0.5	0.2
16:3 n-4	0.9	0.3
16:4 n-1	0.8	0.4
Sum other PUFA	2.4	1.0
18:2 n-6	2.1	10.6
18:3 n-6	0.2	0.1
20:2 n-6	0.2	0.1
20:4 n-6	0.7	0.5
Sum n-6 PUFA	3.3	11.4
18:3 n-3	0.8	21.1
18:4 n-3	2.0	0.6
20:4 n-3	0.7	0.2
20:5 n-3	10.2	5.3
21:5 n-3	–	0.2
22:5 n-3	0.8	0.7
22:6 n-3	12.3	7.6
Sum n-3 PUFA	26.9	35.6
Sat/PUFA	0.7	0.4
n3/n6	8.2	3.1

PUFA polyunsaturated fatty acids

Directive (2010/63/EU) on the protection of animals for scientific purposes, at the Marine Zoological Station, University of Porto, Portugal. European sea bass (*D. labrax*) juveniles with a mean initial body mass of 96 ± 0.8 g were placed in a thermoregulated recirculation water system equipped with a battery of twelve fibreglass tanks of 60 L capacity and supplied with continuous flow of filtered seawater. After 2 weeks of adaptation to the experimental conditions, 16 fish were randomly distributed to each tank and each diet was randomly assigned to triplicate groups of fish. The trial lasted 36 days, and fish were hand-fed to satiation twice a day. During the trial, water temperature averaged 24.1 ± 0.9 °C, salinity averaged 34.7 ± 0.8 g L⁻¹ and dissolved oxygen was kept near saturation. The photoperiod was the natural one for May to June. This study is part of a bigger study, and further information on nutrient digestibility, serum metabolites and hepatic and intestinal gene expression analysis of key genes involved in lipid transport and lipoprotein metabolism is presented elsewhere (Castro et al. 2015a).

At the end of the trial, two fish from each tank were randomly sampled at 2, 6 and 24 h after the last meal and euthanized by a sharp blow to the head. Fish were dissected on chilled trays, and the liver and digestive tract were excised and freed from adherent adipose and connective tissues. For digestive enzyme analyses, the intestine was divided in two sections: anterior intestine (AI, section between the last pyloric caeca and the mid-line of the intestinal length) and posterior intestine (PI, section from the mid-line of the intestinal to distal intestine). Each section with intestinal content was immediately frozen in liquid nitrogen and stored at -80 °C until analysis.

For histological analyses, liver and intestine samples were collected from the fish euthanized at 6 h post-feeding. For intestinal evaluation, two pyloric caeca, and sections with approximately 0.5 cm of AI, PI and distal intestine (DI, collected from the last part of the posterior intestine, visually distinguished by a darker and thicker mucosa) were collected. Tissue samples were rinsed in phosphate-buffered saline (PBS), blotted dry with a paper towel, immediately fixed in phosphate-buffered formalin (4 %, pH 7.4) for 24 h and then transferred to ethanol (70 %) until further processing. An additional section of the liver was collected to cryotubes containing 10 % formal + 30 % saccharose + 1 % CaCl₂, left at 4 °C

for 48 h, then transferred to frozen embedding medium (Thermo Scientific™ Shandon Cryomatrix™) and stored at -20 °C until processing.

Digestive enzyme activities

Each section of the digestive tract was homogenized (dilution 1:9, w/v) in ice-cold buffer (100 mM Tris-HCl, 0.1 mM EDTA and 0.1 % Triton-X-100 (v/v), pH 7.8) and centrifuged at 30,000g for 30 min at 4 °C. The resultant supernatants were collected, and aliquots were stored at -80 °C until digestive enzyme analysis.

For each enzyme activity, assay dilution tests were previously done to ensure optimum ratio between enzyme and substrate. All enzyme activities were measured at 37 °C in a microplate reader (ELx808; Bio-Tek Instruments). The specific assay conditions for each enzyme were as follows:

α -Amylase (EC 3.2.1.1) activity was determined with a commercial kit (ref. 41201, Spinreact, Girona, Spain), with modification in the proportion of supernatants and assay buffer (200 μ L of the assay buffer with 10 μ L of supernatants). The rate of product formation (2-chloro-4-nitrophenol) was quantified at 405 nm.

Lipase (EC 3.1.1.3) activity was determined using a commercial kit (ref. 1001275, Spinreact, Girona, Spain) with modification in the proportion of supernatants and assay reactives (200 μ L of the assay buffer, 40 μ L substrate with 10 μ L of supernatants). 1-2-O-dilauryl-rac-glycero-3-glutaric acid-(6'-methylresorufin)-ester was used as substrate, and the formation rate of methylresorufin was followed at 580 nm.

Trypsin (EC 3.4.21.4) activity was determined according to Faulk et al. (2007), using 1 mM N α -Benzoyl-L-arginine 4-nitroanilide hydrochloride (BAPNA) as substrate combined with 50 mM Tris-HCl, 20 mM CaCl₂, pH 8.2 buffer. Production of p-nitroaniline was monitored at 410 nm.

Alkaline phosphatase activity was determined using a commercial kit from Sigma (Sigma procedure no. 104) as previously described by Krogdahl et al. (2003). The reaction was measured at 420 nm using p-nitrophenyl phosphate as substrate and p-nitrophenol as standard.

Total alkaline protease activity was determined by the casein-hydrolysis method described by Walter (1984) and adapted by Hidalgo et al. (1999). A reaction mixture containing 0.25 mL casein at 1 % (w/

v), 0.25 mL buffer (0.1 M Tris HCl, pH 9) and supernatant from the homogenates (0.1 mL) was incubated for 1 h at 37 °C. A control blank for each sample was assayed adding the supernatant from the homogenates after the incubation time. The reaction was stopped by the addition of 0.6 mL 8 % (w/v) trichloroacetic acid solution in blanks and reaction samples. After being kept for 1 h at 4 °C, blanks and reaction samples were centrifuged at 1800g for 10 min and the absorbance of supernatants measured at 280 nm. Tyrosine solution was used as standard.

Unit (U) of enzyme activity was defined as μmol of product generated per minute under the measurement conditions described above and expressed per mg soluble protein (specific activity). Protein concentration was determined according to Bradford (1976) using Sigma protein assay kit and bovine serum albumin as standard.

Protease inhibition, SDS-PAGE and casein zymography

Alkaline protease zymograms were obtained after resolving by SDS-PAGE the homogenates of AI and PI of fish sampled at 6 h post-feeding (Moyano et al. 1996). Prior to electrophoresis, a range of specific inhibition solutions were used and selected according to Alarcón et al. (1998). Thus, 10 mM TLCK (N-alpha-tosyl-L-lysiny-chloromethylketone) was used to inhibit trypsin-like serine proteases; 10 mM ZPCK (N-carbobenzoxy-L-phenylalanine chloromethyl ketone) was used to inhibit chymotrypsin-like serine proteases, and 100 mM PMSF (phenylmethanesulfonyl fluoride) was used to inhibit serine proteases in general. Inhibition reactions were prepared by combining 40 μL of the intestinal homogenate containing 35 mg of soluble protein with 10 μL of each inhibitor (TLCK, ZPCK or PMSF) or with 10 μL of H_2O (control). Following 45-min incubation at room temperature, each sample mixture (with or without inhibitor) was mixed with sample loading buffer (2 % SDS, 0.125 M Tris, 5 % glycerol, 0.025 % bromophenol blue) and loaded on 12.5 % polyacrylamide gel for protein resolution by SDS-PAGE, according to Laemmli (1970) and adapted by García-Carreño et al. (1993). The full-range Amersham Rainbow Marker (GE Healthcare Bio-Sciences AB, Uppsala, Sweden) was used to estimate the molecular weight of proteins. Electrophoresis was performed on a Mini-PROTEAN® Tetra Cell (Bio-

Rad Laboratories, Inc.) at 25 mA per gel during 120 min. After electrophoresis, the gels were washed 3×10 min with 2.5 % Triton-X-100 to remove SDS and then incubated for 30 min at 4 °C, under agitation, with a 2 % casein substrate solution prepared in 50 mM TrisHCl, pH 8.2. The gels were then kept for additional 90 min at room temperature with shaking. Following substrate incubation, the gels were stained for 50 min with staining solution (0.3 % Coomassie blue R-250, 50 % ethanol, 10 % acetic acid) and de-stained using a mixture of 30 % ethanol and 10 % acetic acid, for protein visualization. The effect of each inhibitor on digestive proteases was visualized by partial or total disappearance of one or more hydrolysis bands, when compared to the profile of enzyme extract pre-incubated with distilled water.

Histology

Samples were processed and sectioned using standard histological techniques and stained with haematoxylin and eosin (H–E). Cryopreserved liver samples were sectioned with cryostat (CM3050 S, Leica, Deerfield, IL), placed on slides, air-dried for 30 min and stored at -20 °C until Sudan black staining (SB). SB was performed as follows: slides were washed in water, passed through ethanol 70 %, stained in SB (Merck 1387, Darmstadt, Germany; 3 % w/v in ethanol 70 %), washed in ethanol 70 %, passed through water and mounted with aqueous mounting media (S3025-Dako).

Liver and intestinal sections were subjected to a double-blinded evaluation using a semi-quantitative scoring system ranging from 1 to 5. Score 1 was considered the normal tissue appearance, and subsequent scores were accounted for increasing alterations of normal tissue histomorphology. Intestinal samples were evaluated according to the criteria suggested by Krogdahl et al. (2003): widening and shortening of the intestinal folds, loss of the supranuclear vacuolization in the absorptive cells (enterocytes) in the intestinal epithelium, widening of the lamina propria in the intestinal folds and infiltration of mixed leucocyte population in the lamina propria and submucosa.

Liver samples were evaluated for general histomorphology, giving particular attention to cytoplasm vacuolization of the hepatocytes and any signs of inflammation. The SB sections were used to confirm the nature of hepatocellular vacuolization observed in H–E-stained preparations.

Statistical analysis

Digestive enzyme activity data were checked for normal distribution and homogeneity of variances (Shapiro–Wilk and Levene tests, respectively) and normalized when appropriate. The effects of dietary lipid source (LS) sampling time (TIME) and intestinal section (SECTION) on digestive enzyme activities were analysed by 3×3 factorial analysis (three-way ANOVA). Significant differences between means were evaluated by the Tukey multiple range test. In case of significant interaction, a one-way ANOVA was performed to evaluate differences between time and section. Histological data were neither normal nor homogeneous; thus, the Kruskal–Wallis nonparametric test was used. The significance level of 0.05 was used for rejection of the null hypothesis. All statistical analyses were done using the IBM SPSS 21 software package (SPSS® Inc, Chicago, IL, USA) for Windows.

Results

Dietary VO inclusion had no effect on amylase, lipase, total alkaline proteases and trypsin activities ($\text{mU mg protein}^{-1}$) in the AI and PI. Contrarily, alkaline phosphatase activity was higher in both intestinal sections in fish fed the VO diet (Table 3).

Differences in digestive enzyme activities among AI and PI sections were mostly time-related (time \times section interaction) (Table 3). At 2 h post-feeding, amylase, lipase, alkaline phosphatase and trypsin activities were higher in AI than in PI, while at 6 and 24 h after feeding, lipase and total alkaline protease activities were lower in AI than in PI (Fig. 1). At 6 and 24 h post-feeding, alkaline phosphatase and trypsin activities were still higher in AI than in PI, respectively (Fig. 1). Differences among AI and PI sections on amylase (at 6 and 24 h post-feeding), total alkaline proteases (at 2 h post-feeding), alkaline phosphatase (at 24 h post-feeding) and trypsin (at 6 h post-feeding) activities were not observed (Fig. 1). Time \times LS confirmed that there were no significant differences in these enzymes among AI and PI.

In AI, digestive enzyme activities were similar at 2 and 6 h after feeding, and lowest values were observed at 24 h (Fig. 1). In PI, digestive enzyme activity tended to increase from 2 to 6 h post-feeding and lowest values were observed at 24 h (Fig. 1).

Zymogram analysis of alkaline proteases present in AI and PI sections of fish fed the two diets showed that the number of detectable active proteases was not affected by the dietary lipid source. Five bands with proteolytic activity against a casein substrate were detected in both intestinal sections, with molecular weights ranging between 17 and 52 kDa (band numbers 1–5, Fig. 2). All these proteins are predicted to be serine proteases, since their hydrolytic activity was reduced upon addition of PMSF, an inhibitor of serine protease activity. An additional band with an estimated molecular weight of less than 17 kDa (band number 6, Fig. 2) was visible in PI section. This band was apparently not inhibited by any of the inhibitors used (TLCK, ZPCK or PMSF), indicating that it probably corresponds to a nonserine protease (different catalytic residue). The two bands with estimated molecular weight between 17 and 24 kDa (bands 4 and 5, Fig. 2) were identified as chymotrypsin-like serine enzymes, as observed by a reduction in the band intensity when using the chymotrypsin-specific inhibitor ZPCK. On opposite, the band immediately above (band 3, Fig. 2), with an estimated molecular weight higher than 24 kDa, corresponds most probably to a trypsin-like serine protease, since a reduction in its intensity could be observed with TLCK (trypsin inhibitor). The two uppermost bands (bands 1 and 2, Fig. 2), with highest molecular weights, could putatively be identified as serine protease enzymes (other than trypsin-like or chymotrypsin-like), since their intensity was only reduced with PMSF treatment.

The inclusion of VO in the diets had no effects on the intestinal histomorphology (Table 4; Fig. 3). Fish from both dietary treatments showed similar histological features, with long, well-defined villus without fusion. Lamina propria and submucosa presented normal thickness without increased cellularity. The frequency of eosinophilic granulocytes and intraepithelial leucocytes was similar in fish from both dietary groups.

Both dietary treatments resulted in great hepatocyte vacuolization, and the group fed VO diets exhibited a tendency ($p = 0.071$) towards higher degree of vacuolization than the FO group (Fig. 4). The cytoplasmic vacuoles appeared clear and optically empty, often pushing the nucleus to the cell periphery without altering the shape of the nuclei. The Sudan black staining (Fig. 4c, d) confirmed the lipidic nature of the material contained in the cytoplasmic vacuoles,

Table 3 Postprandial specific digestive enzyme activities (mU mg⁻¹ protein) in the anterior and posterior intestine of European sea bass fed the experimental diets

Enzymes		Amylase			Lipase			Alkaline phosphatase		
Section	LS	2 h	6 h	24 h	2 h	6 h	24 h	2 h	6 h	24 h
AI	FO	42 ± 25	34 ± 12	3.3 ± 1.7	3.9 ± 2.3	3.5 ± 1.4	0.4 ± 0.1	199 ± 100	242 ± 46	46 ± 12
	VO	36 ± 8	35 ± 16	3.9 ± 1.8	3.8 ± 0.7	2.8 ± 1.0	0.4 ± 0.1	311 ± 102	227 ± 104	73 ± 37
PI	FO	26 ± 15	28 ± 19	3.1 ± 1.6	2.4 ± 1.6	5.7 ± 2.8	0.5 ± 0.1	117 ± 30	150 ± 55	57 ± 12
	VO	11 ± 6	28 ± 17	4.1 ± 1.9	1.1 ± 0.7	6.5 ± 4.5	0.5 ± 0.1	121 ± 45	214 ± 65	68 ± 21
<i>p value</i> *										
LS		0.470			0.174			0.020		
TIME		<0.001			<0.001			<0.001		
SECTION		0.007			0.867			<0.001		
LS × TIME		0.164			0.428			0.509		
LS × SECTION		0.477			0.269			0.607		
TIME × SECTION		0.043			<0.001			0.002		
LS × TIME × SECTION		0.368			0.180			0.019		
Enzymes		Total alkaline proteases			Trypsin					
Section	LS	2 h	6 h	24 h	2 h	6 h	24 h			
AI	FO	193 ± 49	188 ± 30	31 ± 4	133 ± 138	162 ± 98	5.8 ± 1.0			
	VO	158 ± 29	178 ± 57	37 ± 14	106 ± 41	140 ± 67	5.8 ± 0.6			
PI	FO	167 ± 47	341 ± 132	80 ± 27	72 ± 94	312 ± 177	3.9 ± 1.0			
	VO	126 ± 20	252 ± 67	79 ± 19	18 ± 22	294 ± 207	4.2 ± 0.5			
<i>p value</i> *										
LS		0.100			0.182					
TIME		<0.001			<0.001					
SECTION		<0.001			0.022					
LS × TIME		0.230			0.384					
LS × SECTION		0.519			0.297					
TIME × SECTION		<0.001			<0.001					
LS × TIME × SECTION		0.991			0.255					

Values represent mean ± SD ($n = 6$)

LS dietary lipid source, FO fish oil, VO vegetable oil blend, AI anterior intestine, PI posterior intestine, TIME sampling time, SECTION intestinal section

* Significant differences ($p < 0.05$). Results of one-way ANOVA performed to evaluate differences between sampling time and intestinal section are given in the “Results” section and Fig. 1

showing that fish fed the VO diet had a tendency to higher hepatic lipid accumulation than fish fed FO diet.

Discussion

A few studies have already evaluated the effect of dietary protein source on the profile of digestive enzyme activities over time in fish (Santigosa et al. 2008; Rodiles et al. 2012). But, as far as we are aware,

this is the first study addressing the effect of dietary lipid source on digestive enzyme activities over time carried out in fish. Accordingly, partial or total replacement of fish meal (FM) by plant protein sources in the diets for rainbow trout induced a drastic decrease in postprandial alkaline protease activity and a lack of enzyme activity peak when dietary FM was totally replaced by plant protein (Santigosa et al. 2008). On the contrary, the same authors observed in gilthead sea bream similar postprandial protease activity profiles, independently of diet composition,

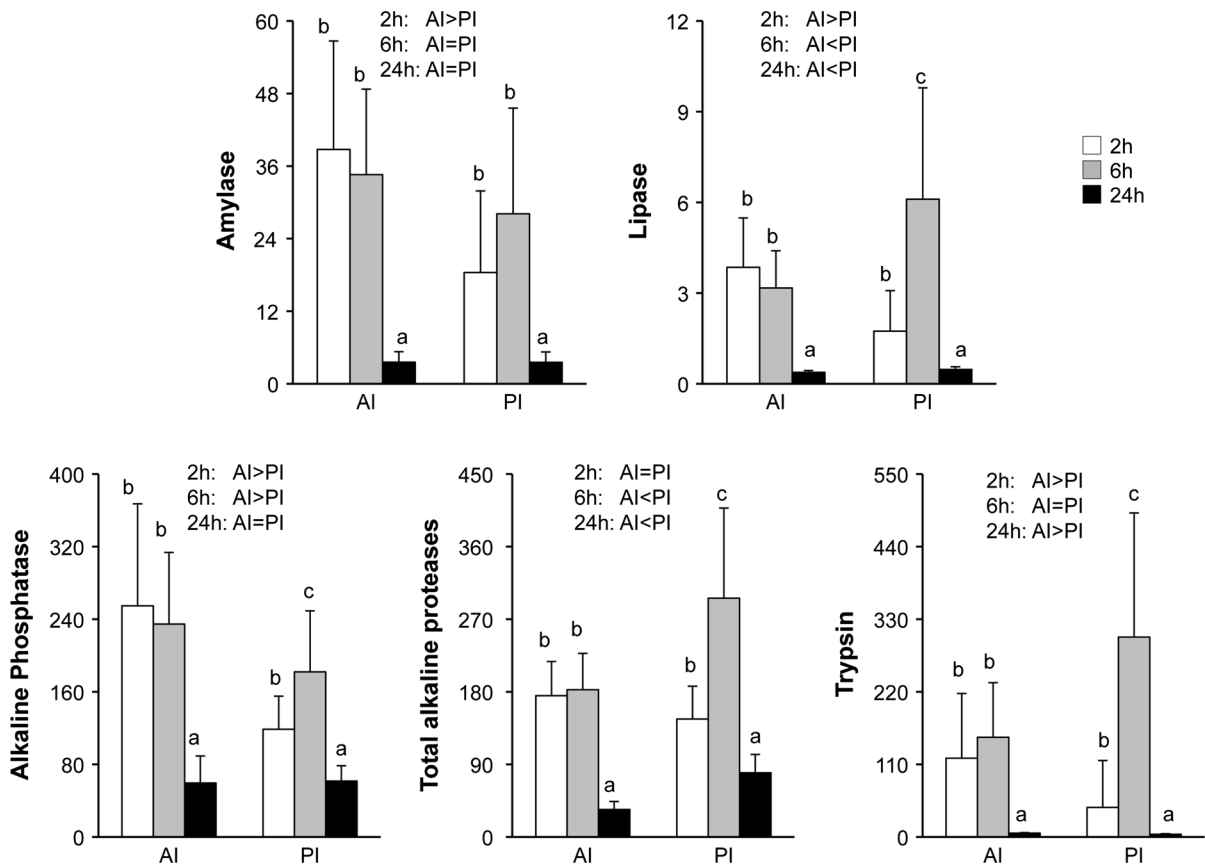


Fig. 1 Summary of the statistical analyses of the specific digestive enzymes (mU mg^{-1} protein) in European sea bass whose activities showed a significant sampling time (TIME) \times intestinal section (SECTION) interaction* (means of the fish fed two experimental diets, $n = 12$). Anterior (AI) and posterior (PI) intestine. *Identified by three-way ANOVA.

Values are mean \pm SD. Significant differences ($p < 0.05$) among sampling times are indicated by *different lower-case letters*, while within each sampling times, significant differences ($p < 0.05$) between AI and PI are indicated by symbols ($>$ or $<$), and the lack of differences ($p > 0.05$) between AI and PI is indicated by symbol (=)

though a gradual decrease in alkaline protease activity peak was observed as dietary plant protein increased (Santigosa et al. 2008). The authors also observed a postprandial peak in amylase activity in both species when fed FM-based diets, but no clear peak in amylase activity was observed when fed plant-protein-based diets. In Senegalese sole, different postprandial profiles of total proteases, trypsin and chymotrypsin activities in fish fed FM-based diets or diets with FM partially replaced by plant protein sources were also reported (Rodiles et al. 2012).

It is reasonable to assume that differences in dietary fatty acid (FA) composition may induce changes in digesta residence time along the digestive tract and, consequently, distinctive digestive enzyme activity profiles over time. However, present results indicate a

lack of time-related changes in digestive enzyme activities in response to dietary lipid source. On the other hand, the postprandial digestive enzyme activity profiles evidenced that enzyme secretion was stimulated by the presence of nutrients in the digestive tract. The similar digestive enzyme activities in the AI at 2 and 6 h post-feeding suggest that digesta was still arriving to the AI at 6 h post-feeding, while the higher enzyme activity in the PI at 6 h post-feeding suggests that at 2 h only a small amount of digesta was present in that region, while at 6 h post-feeding PI was full of digesta. The very low activity values of digestive enzymes at 24 h post-feeding in both sections of the intestine indicate that, by then, digestive processes were already accomplished and that pre-feeding enzyme activity values were achieved.

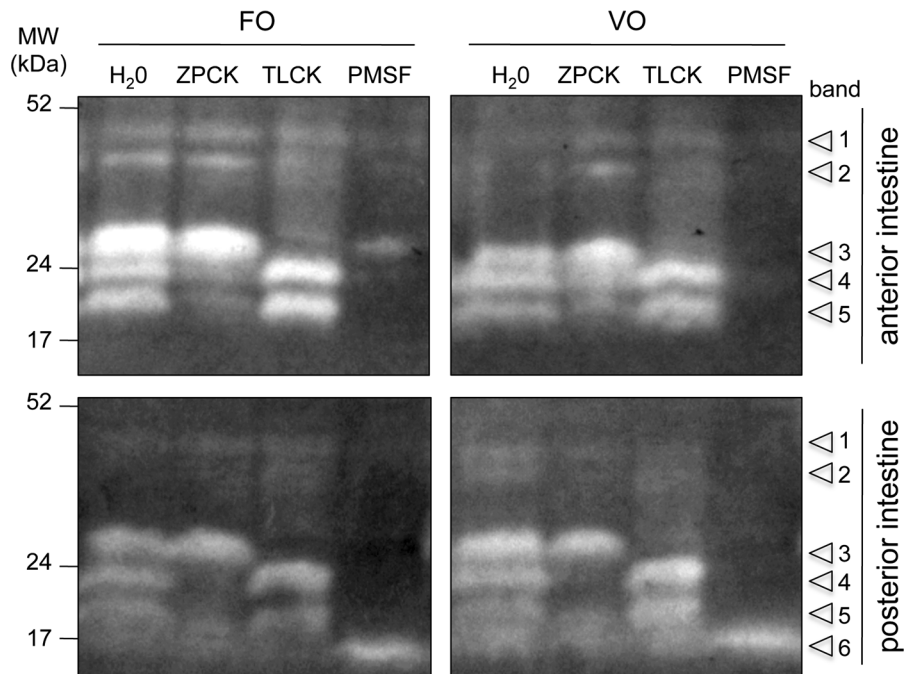


Fig. 2 Zymograms of alkaline proteases in anterior (AI) and posterior (PI) intestine homogenates of European sea bass fed the fish oil (FO) and the vegetable oil blend (VO) diets at 6 h post-feeding and treated with different specific inhibitors:

chymotrypsin inhibitor (ZPCK), trypsin inhibitor (TLCK), general serine protease inhibitor (PMSF) or water (H₂O) used as control. *MW* molecular weight in kDa

Table 4 Mean values of histomorphological semi-quantitative evaluation of the pyloric caeca (PC), anterior intestine (AI), posterior intestine (PI), distal intestine (DI) and liver (Li) of European sea bass fed the experimental diets

	PC	AI	PI	DI	Li
FO	1.7 ± 0.2	1.5 ± 0.2	1.9 ± 0.2	1.6 ± 0.4	3.0 ± 0.9
VO	1.5 ± 0.2	1.5 ± 0.3	1.7 ± 0.2	1.6 ± 0.3	3.8 ± 0.4
<i>p value</i> *					
LS	0.163	0.329	0.411	0.936	0.071

Values represent mean ± SD (*n* = 6)

LS dietary lipid source, FO fish oil, VO vegetable oil blend

* Significant differences (*p* < 0.05)

Digestive enzymes that are synthesized in fish exocrine pancreas are secreted into the anterior section of the digestive tract (Kuz'mina 2008; Bakke et al. 2010). Accordingly, this section seems to contribute more for nutrient digestion and absorption than more posterior sections, as demonstrated by a number of histomorphological, enzymatic and nutrient digestibility studies along the digestive tract of fish (Olsen et al. 1999; Nordrum et al. 2000; Chikwati et al. 2013; Hartviksen et al. 2014). In the present study, activities

of digestive enzymes in both intestinal sections were in general high and differences among intestinal sections were mostly time-related (time × section interaction), reinforcing both the stimulation of pancreatic secretions by the feed ingestion and the evolution of digesta along the digestive tract. Activity of the selected digestive enzymes of pancreatic (total alkaline proteases, trypsin, amylase and lipase) and enterocyte brush border membrane (BBM) (alkaline phosphatase) origin in the present study has also been

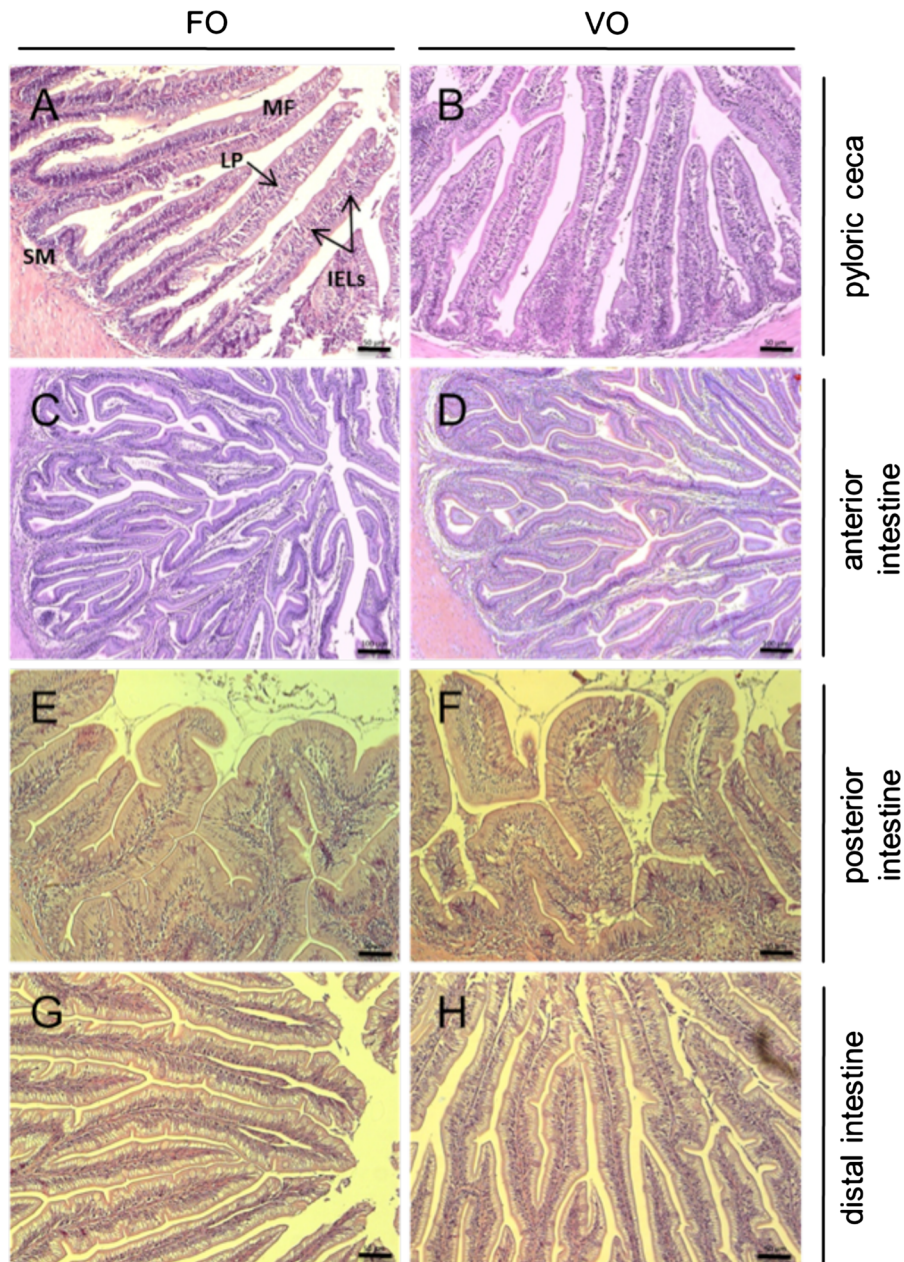


Fig. 3 Histomorphological features of the pyloric ceca (a, b), anterior intestine (c, d), posterior intestine (e, f) and distal intestine (g, h) of European sea bass fed the fish oil—FO (a, c, e,

g) or the vegetable oil—VO (b, d, f, h) diets. *MF* mucosal fold, *SM* submucosa, *LP* lamina propria, *IELs* intraepithelial leucocytes. Scale a, b, e, f, g, h 50 μ m; c, d 100 μ m; H–E staining

detected throughout the digestive tract in several fish species (Krogdahl et al. 2005; Pérez-Jiménez et al. 2009).

The digestive enzyme activity profiles in response to a single meal fit well with the postprandial profiles of serum metabolites (reported in Castro et al. 2015a). However, present results do not allow linking the

distinctive postprandial profile of serum lipids observed in fish fed the two experimental diets to different specificities of lipases in relation to FA profile of the tested diets. Lipase specificity is known to change in function of the unsaturation degree and of the chain length of dietary FA (Koven et al. 1994; Olsen and Ringø 1997; Tocher 2003). However,

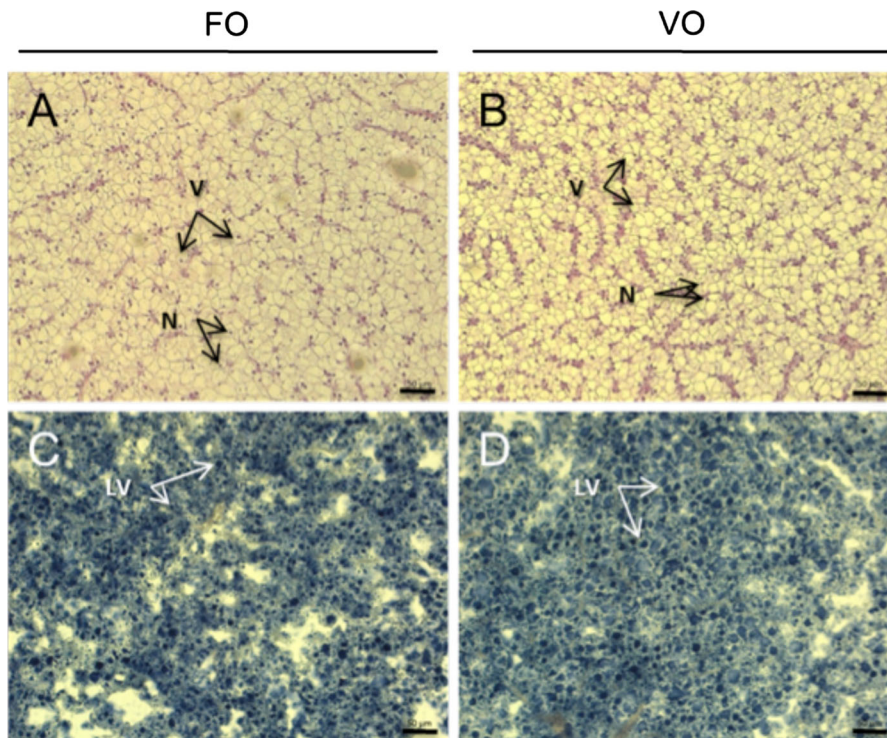


Fig. 4 Histomorphological features of the liver of European sea bass fed fish oil—FO (a, c) or vegetable oil—VO (b, d). *V* cytoplasmic vacuoles, *N* nuclei, *LV* lipid-filled vacuoles. Scale bar 50 μ m; H–E (a, b) and Sudan Black (c, d) stainings

present results did not evidence a lipase activity modulation in relation to dietary lipids. This was not unexpected, as FA chain length and unsaturation degree of the dietary lipids were not much different. These results are also in accordance with the small differences recorded in apparent lipid digestibility of the tested diets (see Castro et al. 2015a). Similarly, no difference on lipase activity was observed in gilthead sea bream juveniles fed diets including FO or a blend of VO (Santigosa et al. 2011). On the other hand, in yellowtail kingfish, lower lipase activity was observed in fish fed diets with canola oil than with FO (Bowyer et al. 2012). On the contrary, in European sea bass larvae, an increment in lipase activity was observed with the ingestion of coconut oil (Morais et al. 2004).

In the present study, no differences in the number of alkaline proteases present in AI or PI sections of intestine were observed between dietary treatments. Also, and consistent with zymogram results, no effect on trypsin activity was detected with the dietary replacement of FO by VO. On the contrary, a decrease in trypsin activity was observed in yellowtail kingfish juveniles fed a diet with canola oil (Bowyer et al.

2012) and in Senegalese sole larvae fed a diet with soybean oil (Morais et al. 2006). The presence of anti-nutritional factors (ANFs), in particular trypsin inhibitors, which may be fat soluble in some types of oils, could explain the differences between the two previous studies and the present results. However, in gilthead sea bream, Santigosa et al. (2011) reported higher total alkaline protease activity in the pyloric caeca of fish fed VO than FO diets, while the opposite was observed in the proximal intestine. Inhibition of luminal proteases due to the presence of ANFs in VO diets was at first hypothesized as the possible cause for the observed differences. However, the authors concluded that the increment in proteolytic activity in pyloric caeca and the higher intensity of trypsin- and chymotrypsin-like proteolytic bands in zymograms promoted by the VO were mainly linked to a slower release of proteases into the intestinal lumen in the VO group due to a decrease in digesta transit rate.

Due to its close nature with the hydrophobic core of intestinal microvillus membrane, the activity of alkaline phosphatase is often used as indicator of intestinal membrane integrity. Available data on the effect of

dietary plant protein (Krogdahl et al. 2003) or lipid level (Cahu et al. 2000) on alkaline phosphatase indicate a decline in activity as indicator of malnutrition. According to Cahu et al. (2000), the decline in alkaline phosphatase activity in fish fed high-lipid diets was due to modifications of FA composition of brush border membrane and, consequently, modification in membrane fluidity. In rainbow trout, Ducasse-Cabanot et al. (2007) found that alkaline phosphatase activity in the AI decreased with the removal of FO from the diet. The authors pointed that the lower digestive capacity in the enterocyte membrane of fish fed low-lipid diets was related to the lower levels of dietary lipids rather than to a modification in the enterocyte brush border membrane FA composition, which remained unchanged. Contrarily to the present data, a recent study from Ribeiro et al. (2015) reported that alkaline phosphatase activity in intestine of meagre was not affected when dietary FO was replaced by a VO blend (rapeseed, linseed and palm oils). Although increased alkaline phosphatase activity in our study may suggest some modification at the enterocyte level in response to the presence of VO in the diet, no histopathological alterations were observed (see below). Even so, and considering that alkaline phosphatase is involved in several other cellular functions, further studies are required to understand potential mechanisms of action of dietary VO.

Although accumulation of lipid droplets in the enterocytes due to dietary VO has been reported in previous studies in other fish species (Caballero et al. 2003; Kowalska et al. 2010), in the present study, dietary VO did not affect intestinal histomorphology of European sea bass. A previous study in this species (Mourente et al. 2007) using identical VO blend and dietary inclusion levels also found no major histomorphological alterations on mid- or distal intestine, just increased absorptive vacuolization in the proximal intestine. Apart from species differences, the degree of lipid droplet accumulation in the enterocytes was also shown to vary with the level and type of dietary FA composition (Caballero et al. 2002, 2003; Santigosa et al. 2011). Accordingly, Caballero et al. (2003) found high accumulation of lipid droplets in gilthead sea bream fed diets containing linseed or soybean oils replacing 60 % FO, and even higher accumulation when fish were fed the same VO replacing 80 % FO, or with rapeseed oil at 60 %. Also in gilthead sea

bream, Santigosa et al. (2011) found increased lipid droplet content in proximal intestine as the percentage of dietary VO blend increased, replacing from 33 to 100 % of dietary FO. These histomorphological alterations are thought to occur due to modifications in lipid transport mechanisms, more precisely with interferences of dietary lipid source on the intestinal lipoprotein assembly, namely in the reacylation mechanisms of the absorbed lipids (Caballero et al. 2003, 2006). The lack of alterations on tissue histomorphology due to dietary treatments in the present study is in accordance with the lack of transcriptional modifications of proteins related to lipid transport and lipoprotein metabolism (FABP, DGAT, ApoB, ApoA4, ApoA1, MTP) (see Castro et al. 2015a).

The effects of dietary VO on fish liver histomorphology are very diverse. In European sea bass fed diets with dietary soybean oil replacing 50 % of FO (Figueiredo-Silva et al. 2005) or a blend of VO replacing 60 % of FO (Mourente et al. 2007), no histomorphological effects were observed. Similarly, Ribeiro et al. (2015) observed in meagre that histological structure of liver was little affected by the 60 % replacement of FO by a VO blend. Contrarily, occurrence of variable size vacuoles and large amounts of lipid droplets in the liver of European sea bass fed rapeseed and linseed oils replacing 60 % of FO was described by Mourente et al. (2005). In the liver of rainbow trout (Caballero et al. 2002), gilthead sea bream (Caballero et al. 2004; Wassef et al. 2007; Fountoulaki et al. 2009) and Atlantic salmon (Kjaer et al. 2008), increased lipid accumulation was detected when fish were fed VO-rich diets. In the present study, the tendency towards higher hepatocellular vacuolization caused by an accumulation of lipid droplets in the VO group may be related to an induction of lipogenesis. In several fish species, VO administration promoted an increase (Jordal et al. 2007; Morais et al. 2011; Peng et al. 2014) in the transcript or enzymatic activity of lipogenic-related enzymes. In addition, and even though no changes were reported on the hepatic gene expression level in our fish (Castro et al. 2015a), it is also possible that factors involved in the assembly and secretion of the lipoprotein particles may also be involved. However, these hypotheses need further investigation, as available literature on the effects of FO replacement by VO on lipogenesis (Regost et al. 2003; Menoyo et al. 2004; Torstensen et al. 2004; Richard et al. 2006a, b; Castro et al. 2015b)

and on the mechanisms of lipoprotein assembly and secretion (Richard et al. 2006a, b; Jordal et al. 2007; Kjaer et al. 2008; Leaver et al. 2008) is not clear and contradictory. Nevertheless, this kind of tissue histomorphological alteration at the hepatic level seems to be nonpathological and reversible when fish are re-fed with a balanced diet (Caballero et al. 2004).

Overall, FO replacement by circa 70 % of a balanced VO blend in European sea bass juveniles did not substantially modulate digestive enzyme activities, suggesting that luminal digestive processes were not affected by dietary lipids tested. The kinetics of digestive enzyme activity response to a single meal reflected feed ingestion and was not affected by dietary oil source. VO blend promoted no hepatic or intestinal histomorphological alterations. Nevertheless, a trend towards higher lipid accumulation in the hepatocytes was observed in the fish fed the VO diet, which may result in effects at long term, and requires to be further investigated.

In the context of the development of more sustainable and economic aquafeeds, the present data seem promising as it suggests that FO can be replaced up to 70 % with VO in diets for European sea bass without major alterations in the digestive function and intestinal histomorphology.

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