

1                   **Dietary fatty acids affect mitochondrial phospholipid compositions and**  
2                   **mitochondrial gene expression of rainbow trout liver at different ages**

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24                  **Abbreviations:** ANT, nucleotide translocase; BHT: butylated hydroxytoluene; cDNA,  
25                  complementary DNA; CL, cardiolipin; COX, cytochrome c oxidase complex; DHA,  
26                  docosahexaenoic acid; E, PCR efficiency; EPA, eicosapentaenoic acid; ETC, electron transport  
27                  chain; FA, fatty acid; FAME, fatty acid methyl esters; HP-TLC, high performance thin layer  
28                  chromatography; HUFA, highly unsaturated fatty acids; LA, linoleic acid; LC-PUFA, long chain  
29                  polyunsaturated fatty acid; MPH, membrane pacemaker hypothesis; mtDNA, mitochondrial  
30                  DNA; MUFA, monounsaturated fatty acids; NAC, no-amplification control; ND, NADH-coenzyme  
31                  Q oxidoreductase complex; NTC, no-template control; PC, phosphatidylcholine, PE,  
32                  phosphatidylethanolamine; PI, phosphatidylinositol; PIn, peroxidation index; PL, phospholipid;  
33                  PS, phosphatidylserine; PUFA, polyunsaturated fatty acid; RO, rapeseed oil; ROS, reactive  
34                  oxygen species; SFA, saturated fatty acids; FO, fish oil; SM, sphingomyelin; RT-PCR, real-time  
35                  PCR; TBARS, thiobarbituric acid reactive substances; TBA, thiobarbituric acid; TCA,  
36                  trichloroacetic acid; TLC, thin layer chromatography.

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38                  **Running title:** Effects of diet lipid composition on rainbow trout liver mitochondria

39 **Abstract**

40 Mitochondria are among the first responders to various stressors that challenge the  
41 homeostasis of cells and organisms. Mitochondrial decay is generally associated with  
42 impairment in the organelle bioenergetics function and increased oxidative stress, and it  
43 appears that deterioration of mitochondrial inner membrane phospholipids (PL), particularly  
44 cardiolipin (CL), and accumulation of mitochondrial DNA (mtDNA) mutations are among the  
45 main mechanisms involved in this process. In the present study, liver mitochondrial membrane  
46 PL compositions, lipid peroxidation and mtDNA gene expression were analyzed in rainbow trout  
47 fed three diets with the same base formulation but with lipid supplied either by fish oil (FO),  
48 rapeseed oil (RO) or a high DHA oil (DHA) during six weeks. Specifically, two feeding trials were  
49 performed using fish from the same population of two ages (1 and 3 years), and PL class  
50 compositions of liver mitochondria, fatty acid composition of individual PL classes, TBARS  
51 content and mtDNA expression were determined. Dietary fatty acid composition strongly  
52 affected mitochondrial membrane composition from trout liver but observed changes did not  
53 fully reflect the diet, particularly when it contained high DHA. The changes were PL specific, CL  
54 being particularly resistant to changes in DHA. Some significant differences observed in  
55 expression of mtDNA with diet may suggest long-term dietary effects in mitochondrial gene  
56 expression which could affect electron transport chain function. All the changes were influenced  
57 by fish age, which could be related to the different growth rates observed between 1- and 3-  
58 year-old trout but that could also indicate age-related changes in the ability to maintain  
59 structural homeostasis of mitochondrial membranes.

60

61 **Keywords:** Cardiolipin, diet, fatty acid, mitochondria, rainbow trout, oxidative stress,  
62 phospholipid.

### 63 1. Introduction

64 Membrane lipid composition can be affected by three main factors in fish: temperature,  
65 age and diet. Ectotherm vertebrates are well known to perform cold-induced lipid remodelling in  
66 the process of homeoviscous adaptation and the majority of the cold-active ectotherm species  
67 investigated so far respond to cold by either increasing the ratio of unsaturated to saturated fatty  
68 acids (FA) or increasing polyunsaturated fatty acids (PUFA) in membranes (Hazel and Williams  
69 1990; Moyes and Ballantyne 2011). Membrane FA composition can also change over the  
70 course of an organism's life. In vertebrates and invertebrates with finite growth, PUFA have  
71 either been shown to increase or remain constant with age (Hulbert *et al.* 2006, Hulbert 2007),  
72 Finally, It is also known that changes in the FA composition of the diet can drastically modify  
73 membrane lipid composition in a wide range of taxa including fish (Hulbert *et al.* 2005, Guderley  
74 *et al.* 2008; Martin *et al.* 2013). These changes affect not only plasma membranes but also  
75 those from subcellular organelles, including mitochondria (Rohrbach 2009), altering their  
76 physical properties and influencing the activities of membrane enzymes and, thus, cellular  
77 metabolic pathways (Barzanti *et al.* 1994). This would be a crucial process in the life-cycle of an  
78 organism as it has been observed in a wide number of animal species that membrane  
79 composition and lifespan are linked through the regulation of metabolic rate (Hulbert 2008). The  
80 membrane pacemaker hypothesis of aging (MPH, also referred to as the homeoviscous-  
81 longevity theory of aging) is an extension of the oxidative stress theory of aging that  
82 emphasizes variation in the FA composition of membranes as an important influence on lipid  
83 peroxidation and consequently on the rate of aging and determination of lifespan. Increased

84 polyunsaturation of cell membrane lipids (high number of double bonds) results in more fluid  
85 membranes that can promote higher molecular activity of membrane proteins and, in turn,  
86 increase the metabolic activity of cells, tissues and, consequently, whole animals (Pamplona *et*  
87 *al.* 2002; Sanz *et al.* 2006). At the same time, membranes with increased levels of PUFA are  
88 more prone to attack by oxidizing agents and will then participate in long, free radical reactions  
89 that will propagate oxidative damage throughout the organelle (Hulbert *et al.* 2005). This  
90 hypothesis would explain the huge differences in life span existing among animal species with  
91 different levels of unsaturation in their membranes, but could also explain how situations such  
92 as thermal or dietary changes in an animal life-cycle, could compromise metabolic activity,  
93 oxidative stress and tissue function.

94         The key role of FA composition in membrane processes suggests that dietary effects  
95 upon membrane lipid structure should be, at least in part, regulated. Both factors, the overall  
96 status of the membrane and the specific characteristics of lipid components, must be balanced  
97 as they, very likely, participate in maintaining membrane function (Paradies *et al.*, 1992;  
98 Zabelinskii *et al.* 1999; Guderley *et al.* 2008; Modi *et al.* 2008). It is known that mitochondrial  
99 membranes have different compositions of lipids such as phospholipids (PL), glycolipids and  
100 cholesterol compared with plasma membranes (Wiseman 1996), this being related with the role  
101 of the organelle in oxygen consumption (Hoch 1992). Mitochondrial inner membrane (MIM)  
102 uniquely contains cardiolipin (CL), a key molecule associated with complexes I, III and IV of the  
103 electron transport chain (ETC),  $F_1F_0$ ATPase and nucleotide translocase (ANT) (Yamaoka *et al.*  
104 1988; Paradies *et al.* 2002). CL has a high content of PUFA which makes it, along with its  
105 proximity to the site of ROS production, particularly prone to peroxidation (Paradies *et al.* 2011).  
106 Mitochondrial membranes also contain small amounts of sphingomyelin (SM), another

107 potentially important PL that is present in all cell membranes and has also been suggested as a  
108 mediator of aging and determinant of life-span (Cutler and Mattson 2001). SM not only has  
109 membrane-rigidifying properties, which retard the lateral propagation of free radicals (Subbaiah  
110 *et al.* 1999), but also is a precursor of many signalling molecules, some associated with  
111 apoptosis (Hannum and Obeid, 1994). The particular roles of individual PL classes are  
112 associated with characteristic FA compositions that confer specific properties related to  
113 membrane fluidity and functions (Zabelinskii *et al.* 1999). Therefore, not only changes in PL  
114 class composition, but also alterations in PL FA compositions would modify their molecular  
115 properties and their roles in membrane functions (Shigenaga *et al.* 1994; Chicco and Sparagna  
116 2007; Crimi and Esposti 2011; Monteiro *et al.* 2013).

117         Although lipid peroxidation is quantitatively the main oxidative process inside  
118 mitochondria, other organelle molecules are also attacked and damaged by ROS including  
119 proteins and nucleic acids (Sanz *et al.* 2006). Mitochondrial DNA (mtDNA) is very exposed to  
120 ROS production as it is located very close to the site of ROS production. Moreover, mtDNA is  
121 not extremely condensed and protected by histones, as it is nuclear DNA, and its repair activity  
122 is limited (Paradies *et al.* 2011). It has been suggested that mtDNA is also a primary target of  
123 ROS and, as more mutations in critical coding regions accumulate, complexes of the ETC  
124 become less efficient or inactive, leading to a decline in mitochondrial function (Paradies *et al.*  
125 2002). Since mtDNA encodes either polypeptides of ETC or components required for their  
126 synthesis, mutations in mtDNA will affect the ETC as a whole. ETC-deficient cells are apoptosis  
127 prone and increased cell loss is therefore a likely important consequence of mitochondrial  
128 dysfunction in situations of high oxidative stress (Trifunovic and Larsson 2008).

129 In summary, mitochondria are among the first responders to various stressors that challenge  
130 the homeostasis of cells and organisms (Manoli *et al.* 2007). Mitochondrial decay is generally  
131 associated with impairment in mitochondrial bioenergetics function and increased oxidative  
132 stress (Paradies *et al.* 2011) and it seems clear that deterioration of MIM PLs, particularly of CL,  
133 and accumulation of mtDNA mutations are mechanisms involved in this process. Changes in  
134 the FA composition of the diet modify mitochondrial membrane composition (reviewed in  
135 Hulbert *et al.* 2005) and can alter the organelle function (Clandinin *et al.* 1985; Barzanti *et al.*  
136 1994; Guderley *et al.* 2008; Martin *et al.* 2013) which can lead to an imbalance in organelle  
137 oxidative status. The aim of the present study was to determine the impact of diet lipid  
138 composition on mitochondrial membrane composition and mtDNA damage as possible  
139 regulators of the processes associated with mitochondrial decay under high oxidative stress  
140 situations in fish. Rainbow trout (*Oncorhynchus mykiss*) was used as a vertebrate model  
141 because it is a well-studied species, widely reared in Europe, its age can be easily monitored  
142 (Almroth *et al.* 2010) and, along with other species of salmonids, it has been used previously in  
143 studies on dietary lipid, oxidative stress and mitochondrial function (Otto and Moon 1996;  
144 Zabelinskii *et al.* 1999; Kraffe *et al.* 2007; Østbye *et al.* 2011). For the present study, fish of two  
145 well-differentiated ages (fast growth period and mature period) were used in order to study the  
146 influence of fish age and body mass on these mechanisms.

147

## 148 2. Methods

### 149 2.1. Experimental fish and sampling

150 A population of rainbow trout, *Oncorhynchus mykiss*, was maintained from eggs  
151 (Howietoun Fishery, Stirling, UK) to 3 years under controlled feeding and rearing conditions

152 throughout in the freshwater aquarium facilities at the Institute of Aquaculture, University of  
153 Stirling. Fish were fed twice a day *ad libitum* with a commercial feed (50% protein, 20% fat,  
154 Skretting, Northwich, UK) and kept under ambient water temperature conditions before being  
155 transferred to experimental thermo-regulated tanks. When fish reached one year of age, 240  
156 fish ( $72.4 \pm 12.6$  g average body weight) were distributed into six tanks equipped with a  
157 thermostat to maintain the temperature at  $12 \pm 1$  °C, kept under a 12L:12D photoperiod and fed,  
158 in duplicate, one of three experimental diets consisting of the same base formulation, but with  
159 lipid supplied either by fish oil (FO diet), rapeseed oil (RO) or an oil containing high  
160 docosahexaenoic acid, 22:6n-3 (DHA) (Table 1). Fish were fed twice a day *ad libitum* for 6-  
161 weeks and then euthanized by exposure to the anesthetic benzocaine hydrochloride (400 mg  
162 l<sup>-1</sup>) for 10 min following the cessation of opercular movement. Livers were collected for  
163 analyses. The experiment was repeated identically when the remaining fish (60 fish of  $625.1 \pm$   
164  $33.8$  g of average body weight; 10/tank) reached 3 years of age. For 1-year-old fish, livers of 5  
165 animals were pooled to provide sufficient material for the different analyses and also to reduce  
166 inter-individual variability. For 3-year-old animals, two livers per pool were taken after  
167 considering fish size and numbers. A portion of each pool was immediately taken into  
168 RNA<sup>later</sup>® (Life technologies, Paisley, UK) following the manufacturer instructions and stored at  
169  $-20$  °C for molecular analysis, while the remainder of the tissue sample was processed to  
170 obtain enriched mitochondrial preparations. The purified mitochondrial isolates were analyzed to  
171 determine mitochondrial membrane lipid composition and peroxidation. Fish were treated in  
172 accordance with British national ethical requirements established by the UK Government Home  
173 Office and guidelines determined by the Animals (Scientific Procedures) Act 1986.

174

175                    *2.2. Experimental Diets*

176                    Practical pelleted diets (2- and 4-mm diameter for 1- and 3-year-old trout, respectively)  
177                    containing 47% crude protein and 24% crude lipid were formulated using the same basal  
178                    ingredients and varying only in the lipid source used: Southern hemisphere fish oil (FO),  
179                    rapeseed oil (RO) and Incromea TG0525 (DHA) (Table 1). The fatty acid compositions of the  
180                    diets are shown in Table 2. The control diet (FO) contained 31.7% saturated fatty acids (SFA),  
181                    mainly 16:0, 27.8% monounsaturated fatty acids (MUFA), 11.7% of which was 18:1n-9, and  
182                    40.6% PUFA, with 6.8% as n-6 PUFA and 30.3% as n-3 PUFA. The FO diet contained a high  
183                    proportion of the n-3 long-chain polyunsaturated fatty acid (LC-PUFA), eicosapentaenoic acid  
184                    (20:5n-3, EPA) and DHA, in approximately equal amounts. Diet RO had lower levels of SFA  
185                    (12.3%), higher percentages of MUFA (56.0% with 48.2% as 18:1n-9) and lower PUFA (31.7%)  
186                    with just 2.1% EPA and 3.6% DHA. Finally, the DHA diet contained similar proportions of SFA,  
187                    MUFA and PUFA to the FO diet but had a lower percentage of EPA (6.6%) and higher  
188                    percentage of DHA (22.3%). The diets were formulated to meet all the known nutritional  
189                    requirements of salmonid fish (National Research Council, 2011).

190

191                    *2.3. Mitochondria isolation*

192                    Approximately 2 g of liver pate was homogenized in 8 mL ice-cold sucrose buffer (0.4 M  
193                    phosphate buffer pH 7.4, 0.25 M sucrose, 0.15 M KCl, 40 mM KF and 1 mM N-acetyl-cysteine)  
194                    using an Ultra-Turrax tissue disrupter (Fisher Scientific, Loughborough, UK). Homogenates  
195                    were centrifuged at 600 g for 6 min, the pellet discarded (cell/nuclei debris) and the supernatant  
196                    re-centrifuged at 600 g. Resulting supernatants were then centrifuged at 6800 g for 10 min and  
197                    the resulting pellet (mitochondrial fraction) used for lipid extraction. To verify that pellets were



198 highly enriched with mitochondria, a portion was fixed in 2.5% glutaraldehyde in 0.1 M  
199 cacodylate buffer overnight at 4 °C, and then processed as specified by Rajapakse *et al.* (2001)  
200 prior to analysis by transmission electron microscopy (Tecnai™ G2 Spirit BioTWIN, FEI Europe,  
201 Eindhoven, The Netherlands).

202

#### 203 *2.4. Lipid extraction and phospholipid class composition*

204 Total lipid contents of livers and diet samples were determined gravimetrically after  
205 extraction by homogenization in chloroform/methanol (2:1, v/v) containing 0.01% butylated  
206 hydroxytoluene (BHT) as antioxidant, basically according to Folch *et al.* (1957). Phospholipid  
207 (PL) classes were separated by high-performance thin layer chromatography (HPTLC) using  
208 10×10 cm silica gel plates (VWR, Lutterworth, England) and methyl acetate/isopropanol/  
209 chloroform/methanol/0.25% (w/v) KCl (25:25:25:10:9, by vol.) as solvent system (Olsen and  
210 Henderson 1989). The lipid classes were visualized by charring at 160 °C for 15 min after  
211 spraying with 3% (w/v) aqueous cupric acetate containing 8% (v/v) phosphoric acid and  
212 quantified by densitometry using a CAMAG-3 TLC scanner (version Firmware 1.14.16)  
213 (Henderson and Tocher 1992). Scanned images were recorded automatically and analyzed by  
214 computer using winCATS (Planar Chromatography Manager, version 1.2.0).

215

#### 216 *2.5. Phospholipid fatty acid composition*

217 Phospholipid classes were separated by preparative-TLC, using silica gel plates (20 x  
218 20 cm) (VWR) and the solvent system as above. Individual phospholipid classes were identified  
219 by comparison with known standards after spraying with 1% (w/v) 2',7'-dichlorofluorescein in

220 97% (v/v) methanol containing 0.05% (w/v) BHT, and visualization under UV light (UVGL-58  
221 Minerallight® Lamp, Ultraviolet Prod. Inc., Calif., USA). Silica corresponding to each  
222 phospholipid class was scraped from the plate into a test tube and subjected directly (on silica)  
223 to acid-catalyzed transmethylation at 50 °C overnight following addition of 2 mL of 1% (v/v)  
224 sulphuric acid in methanol in order to prepare fatty acid methyl esters (FAME) (Christie 2003).  
225 FAME were separated and quantified by gas-liquid chromatography (Carlo Erba Vega 8160,  
226 Milan, Italy) using a 30 m x 0.32 mm i.d. capillary column (CP Wax 52CB, Chrompak, London,  
227 U.K.) and on-column injection at 50 °C. Hydrogen was used as carrier gas and temperature  
228 programming was from 50 °C to 150 °C at 40 °C min<sup>-1</sup> and then to 230 °C at 2.0 °C min<sup>-1</sup>.  
229 Individual methyl esters were identified by comparison with known standards. Data were  
230 collected and processed using Chromcard for Windows (version 1.19).

231

## 232 *2.6. Measurement of thiobarbituric acid reactive substances (TBARS)*

233 Approximately 1 mg of total lipid extract from liver mitochondria was used for the  
234 measurement of TBARS using an adaptation of the protocol of Burk *et al.* (1980). Briefly, 50 µL  
235 of 0.2% (w/v) BHT in ethanol was added to the sample followed by 0.5 mL of 1% (w/v) TBA and  
236 0.5 mL 10% (w/v) TCA, both solutions freshly prepared. The reagents were mixed in a  
237 stoppered test tube and heated at 100 °C for 20 min. After cooling, particulate matter was  
238 removed from the homogenate by centrifugation at 2000 g, and fluorescence in the supernatant  
239 determined in the spectrophotometer at 532 nm against a blank sample. The concentration of  
240 TBARS, expressed as ng g of lipid<sup>-1</sup>, was calculated using the absorption coefficient 0.156 µM<sup>-1</sup>  
241 cm<sup>-1</sup>.

242

243                    *2.7. RNA preparation, cDNA synthesis and quantification*

244                    Total RNA was extracted from individual samples using TRI Reagent (Sigma-Aldrich,  
245 Dorset, UK) according to manufacturer's instructions with high salt precipitation (Chomczynski  
246 and Mackey 1995). RNA quantity, integrity and purity were assessed by agarose gel  
247 electrophoresis and spectrophotometry (NanoDrop ND-1000, Thermo Scientific, Wilmington,  
248 USA). RNA samples were aliquoted and diluted to a final concentration of 200ng/μl. First strand  
249 complementary DNA (cDNA) was synthesized from 2μg of RNA using High Capacity cDNA  
250 Reverse Transcription kits (Applied Biosystem, Paisley, UK) and primed with a mixture of Oligo  
251 dT and random primers (1:3).

252                    Information about target genes was retrieved from the mitochondrion genome sequence  
253 (Genbank accession number DQ288271) and used to design primers for Real-Time PCR (RT-  
254 PCR) with PerlPrimer v1.1.17 (Marshall 2004). Primers were designed to target seven  
255 mitochondrial genes: COX3, ND1, ND3, ND4, ND4L, ND5 and ND6 (Table 3). Three  
256 housekeeping genes were evaluated as internal reference (elongation factor 1 $\alpha$ , glyceraldehyde  
257 3-phosphate dehydrogenase and  $\beta$ -actin) but they were not stable but variable between diets or  
258 age groups and were not used for normalization (data not shown). RT-qPCR reactions were  
259 carried out on a Biometra thermal cycler (Gottingen, Germany) using Luminaris Color HiGreen  
260 qPCR Master mix (Thermo Scientific, Leicestershire, UK) according to the instructions provided  
261 by the manufacturer. To avoid technical variation between runs all treatment and control  
262 samples from both age groups were analyzed on the same plate together with a log<sub>10</sub> serial  
263 dilution of pooled samples used to calculate PCR efficiency (E). Each reaction was conducted in  
264 a total volume of 20 μl in duplicates. All primer pairs were designed to have an optimal  
265 annealing temperature of 60°C. A melting curve analysis was performed after every

266 amplification program to verify specificity of the target and absence of primer dimers, and a no-  
267 template control (NTC) was included with each assay to verify that PCR master mixes were free  
268 of contamination. For each assay E was determined by the equation  $[E = 10^{(-1/\text{slope})}]$  (Table 3).  
269 As the reference genes tested in this study were not usable for normalization, the relative  
270 expression of each gene was calculated using the delta-Ct transformation  $[\text{Ratio}_{(\text{test/calibrator})} = E^{\Delta \text{Ct}(\text{calibrator} - \text{Ct}(\text{test}))}]$  and log transformed before statistical analyses. The average of the 1-  
271 year-old FO treatment was used as the calibrator.  
272

273

## 274 2.8. Indexes and statistical analysis

275 The LC-PUFA index corresponds to the sum of fatty acids with 20 or more carbons and  
276 2 or more double bonds. The peroxidation index (PI<sub>n</sub>) was used as an estimate of PL  
277 susceptibility to oxidation and was calculated using the formula:  $\text{PI}_n = 0.025 \times (\text{percentage of}$   
278  $\text{monoenoics}) + 1 \times (\text{percentage of dienoics}) + 2 \times (\text{percentage of trienoics}) + 4 \times (\text{percentage of}$   
279  $\text{tetraenoics}) + 6 \times (\text{percentage of pentaenoics}) + 8 \times (\text{percentage of hexaenoics})$  (Witting and  
280 Horwitt 1964). Specific growth rate (SGR % day<sup>-1</sup>):  $[(\ln W_t - \ln W_i)/T] \times 100$  where  $W_t$  = mean  
281 final weight,  $W_i$  = mean initial weight and  $T$  = total experimental days (Jaya-Ram *et al.* 2008).  
282 Results are presented as mean ± SD (n = 4 for lipid and TBARS analyses and n = 6 for gene  
283 expression assays). Data were checked for homogeneity of variances by the Levene's test and  
284 percentage data from PL content, PL class composition and PL FA analyses were arc-sin  
285 transformed before further statistical analysis. A two-way analysis of variance (ANOVA) was  
286 used to assess the differences among groups based on diet and age. Post-hoc comparisons  
287 were made using the Bonferroni t-test for multiple comparisons. All statistical analyses were

288 performed using SPSS Statistical Software System version 15.0 (SPSS Inc., Chicago, USA).  
289 Differences were regarded as significant when  $P < 0.05$ .

290

### 291 **3. Results**

292       3.1. *Fish performance* There were no differences in feed intake between the different  
293 dietary groups for either 1- (~1.5% BW /day) or 3-year-old trout (~0.8% BW/day). The 1-year-old  
294 fish showed a considerably higher SGR (around 1.5) than 3-year-old trout (~ 0.45) during the  
295 feeding trial (Fig. 1). No statistically significant differences among the feeding groups were  
296 found for either 1- [ $F(2,88) = 0.786$ ,  $P = 0.459$ ] or 3-year-old trout [ $F(2, 56) = 0.116$ ,  $P = 0.891$ ]. No  
297 significant interaction between fish age and diet was observed for SGR ( $p = 0.549$ ).

298

### 299       3.2. *Effects of diet on mitochondrial phospholipid content of trout liver*

300       The phospholipid (PL) content and PL class composition of liver mitochondria from 1-  
301 and 3-year-old rainbow trout fed the experimental diets are shown in Fig. 2. Phosphatidylcholine  
302 (PC) and phosphatidylethanolamine (PE) were the most abundant PL species in all the  
303 experimental groups, constituting more than 71% of total PL in 1-year-old trout and more than  
304 68% in 3-year-old animals. Among the remaining PL classes and focusing only in the control  
305 group (FO) in both age groups, the rank order was cardiolipin (CL) (9.5 and 11.4% for 1- and 3-  
306 year-old trout, respectively), followed by phosphatidylinositol (PI) (4.4 and 6.6%) with lower  
307 amounts of phosphatidylserine (PS) and sphingomyelin (SM). Mitochondria from liver of 1-year-  
308 old fish fed the FO diet had a different PL composition to that of 3-year-old fish, with significantly

309 more PE ( $p= 0.000$ ), SM ( $p = 0.000$ ) and CL ( $p = 0.020$ ) while the content of PS ( $p = 0.027$ ) and  
310 PI ( $p = 0.000$ ) was lower.

311 Dietary FA composition affected PL content and composition in rainbow trout. In 1-year-  
312 old trout several significant changes were found among experimental groups. The RO group  
313 showed a higher content of PC [ $F(2,9 = 25.051, p = 0.000$ ], PE [ $F(2,9 = 5.050, p = 0.020$ ] and PI  
314 [ $F(2,9 = 6.173, p = 0.010$ ] and lower proportions of total PL [ $F(2,9 = 30.389, p = 0.000$ ], CL  
315 [ $F(2,9 = 55.566, p = 0.000$ ] and SM [ $F(2,9 = 12.062, p = 0.001$ ] compared with the FO and DHA  
316 groups. The FO and DHA groups had a similar PL composition with only CL showing a  
317 significant difference ( $p = 0.009$ ), being higher in fish fed FO compared to fish fed DHA (Fig. 2).  
318 However, the DHA group had the lowest content of total PL. A different situation was found in 3-  
319 year-old fish, with fewer differences among the feeding groups. The effect of diet fatty acid  
320 composition on mitochondrial PL composition was age-dependent for all classes except PS,  
321 (Age\*Diet, Fig. 2).

322

### 323 *3.3. Effects of diet on phospholipid fatty acid compositions of liver mitochondria*

324 Fatty acid compositions of the three main phospholipid classes (PC, PE and CL) from  
325 liver of 1- and 3-year-old rainbow trout are presented in Tables 4-6. The fatty acid compositions  
326 of the minor PL classes, PI, PS and SM, are included as Supplementary Tables 1-3. The FA  
327 profiles of mitochondrial PC, PE and CL from trout liver were basically similar in both age  
328 groups (Tables 4-6). Further differences were found in PI (S. Table 1) and, particularly, in PS  
329 which showed higher DHA ( $p = 0.000$ ) and PIn ( $p = 0.000$ ) in 1-year-old trout fed the FO diet  
330 when compared to 3-year-old fish (S. Table 2). Diet lipid composition significantly affected the  
331 FA profiles of individual PL species from 1-year-old fish mitochondria and these effects were

332 different for each PL class. Mitochondria from 1-year-old fish fed the RO diet showed lower SFA  
333 in the main PL classes (around 8% in PC,  $p = 0.000$ ; 3% in PE,  $p = 0.000$  and CL,  $p = 0.030$ )  
334 and in PS ( $p = 0.000$ ), increased MUFA in PC (around 4%,  $p = 0.0003$ ), PE (2%,  $p = 0.038$ ), PS  
335 (4%,  $p = 0.001$ ) and PI (5%,  $p = 0.002$ ) and decreased MUFA in CL (6%,  $p = 0.000$ ) when  
336 compared with the FO group (Tables 4-6, S. Tables 1 and 2). There were also significant  
337 increased percentages of n-6 PUFA in the three main PL classes, particularly CL ( $p = 0.000$ ),  
338 mainly due to increased linoleic acid (18:2n-6, LA,  $p = 0.000$ ) compared to the FO group. There  
339 was also higher n-6 PUFA content in mitochondrial PS ( $p = 0.002$ ) and particularly, SM (10%,  $p$   
340  $= 0.000$ ) from fish fed the RO diet. Regarding n-3 PUFA, mitochondria from 1-year-old fish fed  
341 the RO diet had lower percentages of n-3 PUFA in the main PL classes compared to that of the  
342 FO group, although this varied quantitatively between the groups. The lowest decrease in n-3  
343 PUFA was observed in PC (around 4%,  $p = 0.012$ ) due to lower EPA ( $p = 0.000$ ) with no  
344 change in DHA ( $p = 0.945$ ), while a higher decrease was obtained in CL (8%,  $p = 0.000$ ), mainly  
345 due to lower DHA content ( $p = 0.000$ ) and very little change in EPA ( $p = 0.001$ ). These changes  
346 were reflected in the peroxidation index (PI<sub>n</sub>) of PC and CL, which was significantly lower in 1-  
347 year-old rainbow trout fed the RO diet ( $p = 0.003$  and  $p = 0.001$ , respectively). Mitochondrial PS  
348 from fish fed the RO diet had around 26 % less n-3 LC-PUFA than that from fish fed the FO diet  
349 ( $p = 0.000$ ) which was reflected in PS PI<sub>n</sub> ( $p = 0.000$ ). Mitochondria from 1-year-old rainbow  
350 trout fed the DHA diet showed fewer significant changes in the main PL classes, particularly CL,  
351 where almost no statistical differences were found when compared with the FO group (Table 6).  
352 Regarding PC and PE from 1-year-old fish fed the DHA diet, lower SFA in PC (~ 3%,  $p =$   
353  $0.000$ ), lower MUFA in PE (2%,  $p = 0.011$ ) and higher n-6 PUFA in both (2 and 3% in PC,  $p =$   
354  $0.001$  and PE,  $p = 0.005$ , respectively) were found (Tables 4 and 5). Total n-3 PUFA was not  
355 significantly altered in any of the three PL classes in mitochondria from 1-year-old fish fed the

356 DHA diet ( $p = 0.294$ ,  $1.000$  and  $0.560$  for PC, PE and CL, respectively). In PC and PE, this was  
357 mainly due to the fact that EPA content decreased (4%,  $p = 0.000$ , and 3%,  $p = 0.048$ ,  
358 respectively) while DHA increased (7%,  $p = 0.000$ , and 3%,  $p = 0.035$ , respectively). PIn  
359 significantly increased in mitochondrial PC ( $p = 0.003$ ) and PE ( $p = 0.032$ ) but not in CL ( $p =$   
360  $1.000$ ) from 1-year-old trout fed the DHA diet compared with those from fish fed the FO diet.  
361 Regarding the minor PL classes, SM showed virtually no differences between DHA and FO  
362 groups (S. Table 3) while PI and PS from fish fed the DHA diet showed lower n-3 LC-PUFA,  
363 particularly PS (around 20% less,  $p = 0.001$ ) and PIn ( $p = 0.000$ ) than those from trout fed the  
364 FO diet (S. Tables 1 and 2).

365 The effects of diet on mitochondrial PL FA compositions varied between the two age  
366 groups as indicated by the significant Age\*Diet interactions (Tables 4-6, S. Tables 1-3). In  
367 mitochondrial PC and PE from 3-year-old trout fed the DHA diet, lower DHA levels compared to  
368 that of 1-year-old fish were observed (Tables 4 and 5). This was reflected in the different impact  
369 of diet composition on PIn of mitochondrial PC and PE between the two age groups ( $p = 0.015$   
370 and  $0.003$ , respectively). Diet DHA had a similar impact on mitochondrial CL FA composition in  
371 both age groups. A similar trend was found in the three minor PL classes. Mitochondrial PI, PS  
372 and SM from 3-year-old fish fed the DHA diet showed fewer changes than those from 1-year-old  
373 trout (S. Tables 1-3). Regarding the RO diet, CL from 3-year-old trout showed less n-6 PUFA  
374 than 1-year-old fish (28.2 vs. 33.2%,  $p = 0.001$ ). While mitochondrial CL from 1-year-old fish fed  
375 the RO diet had around 20% more n-6 PUFA than that from fish fed the FO diet, mitochondrial  
376 CL from 3-year-old animals fed the RO diet had just around 14% more n-6 PUFA than that from  
377 3-year-old fish fed the FO diet. There were no significant differences in n-6 PUFA levels in the  
378 remaining PL classes between the two age groups when fed the RO diet.



379

### 380 3.4. Lipid peroxidation

381 Lipid peroxidation in rainbow trout mitochondria, estimated by measuring the TBARS  
382 content, did not show significant differences among feeding groups in 1-year-old fish [F(2, 9) =  
383 2.626,  $p = 0.097$ ] (Fig. 3). Some changes, however, were found in 3-year-old trout [F(2, 9) =  
384 4.852,  $p = 0.019$ ], with higher lipid peroxidation in the FO group compared to that of the RO  
385 group (1164.4 vs. 590.1 ng/g lipid, respectively). Although 3-year-old fish fed the DHA diet  
386 showed lower lipid peroxidation (652.3 ng/g lipid) than that of the FO group, it was not  
387 statistically significant ( $p = 0.060$ ). Regarding age differences, lipid peroxidation was statistically  
388 higher in 1-year-old trout fed the DHA diet than in 3-year-old fish fed the same diet ( $p = 0.020$ )  
389 (Fig. 3). No significant differences were found between age groups in fish fed the FO or RO  
390 diets. The effect of diet lipid composition on mitochondria lipid peroxidation was not influenced  
391 by age as determined by two-way ANOVA ( $p = 0.430$ ).

392

### 393 3.5. Mitochondrial gene expression

394 One-year old rainbow trout fed the different diets showed some consistent trends in  
395 mitochondrial gene expression but these were only statistically significant for ND4 [F(2, 15) =  
396 4.191,  $p = 0.025$ ] (Fig. 4). Fish fed the RO diet showed higher expression for almost all the  
397 studied genes compared with the FO group while trout fed the DHA diet had in general lower  
398 expression than the FO group. Values for 3-year-old trout were in general more stable between  
399 diet groups. When the two age groups were compared, 3-year-old fish showed generally lower  
400 gene expression than 1-year-old animals (except for fish fed the DHA diet) (Fig. 4), but statistical

401 significant differences were only found in fish fed the RO diet for COX3 ( $p= 0.021$ ), ND1 ( $p=$   
402  $0.013$ ), ND4 ( $p= 0.018$ ) and ND5 ( $p= 0.022$ ). The interaction between age and diet was only  
403 significant for ND4 ( $p= 0.037$ ).

404

## 405 **4. Discussion**

### 406 *4.1. Diet fatty acid composition and mitochondrial lipids*

407 Dietary lipid composition markedly influenced PL contents and FA compositions of  
408 individual PL classes of mitochondrial membranes from rainbow trout liver, but had generally  
409 low impact upon mitochondrial DNA (mtDNA) gene expression. The effects differed in relation to  
410 fish age. One-year-old trout fed RO and, mainly, DHA diet showed a significantly lower total PL  
411 content (percentage of total lipid) in their liver mitochondrial membranes compared with the FO  
412 group. Moreover, the RO group showed significantly higher proportions of PC, PE and PI and  
413 lower percentages of CL and SM, while trout fed the DHA diet only showed a significantly lower  
414 CL content compared with the FO group. These changes in the proportions of individual PL  
415 classes may alter mitochondrial membrane function by affecting charge distribution across the  
416 membrane, membrane permeability properties, catalytic activities or specific enzymes and ETC  
417 function (Daum 1985). Especially interesting was the observed loss of CL considering the  
418 critical role that CL plays in the MIM as a regulator of processes related to oxidative  
419 phosphorylation and mitochondrial integrity (Paradies *et al.* 2002). Loss of CL has been shown  
420 in a variety of tissues in mammals and fish (Chicco and Sparagna 2007; Almaida-Pagán *et al.*  
421 2012) and suggested as one of the first signs of damage caused by high oxidative stress in  
422 mitochondrial membranes. Decreased SM has also been related with oxidative stress since SM  
423 can retard the lateral propagation of free radicals through the membrane and it is an important  
424 mediator of mitochondrial pathways including apoptosis (Hannum and Obeid 1997; Subbaiah *et*  
425 *al.* 1999; Cutler and Mattson 2001).

426 The FA compositions of individual PL classes from liver mitochondria of 1-year-old trout  
427 were strongly affected by the diet lipid composition but the observed changes did not entirely

428 reflect differences among the diets. These data agree with those showing marked modifications  
429 of FA of total mitochondrial PL by diet in rats (Lemieux *et al.* 2008; Abbott *et al.* 2010) and fish  
430 (Ushio *et al.* 1997; Robin *et al.* 2003; Guderley *et al.* 2008; Østbye *et al.* 2011; Martin *et al.*  
431 2013), and indicates the existence of mechanisms regulating FA within mitochondrial PL  
432 classes. Moreover, these data point to the importance of FA composition of specific PL classes  
433 in mitochondrial membranes. Individual mitochondrial PL classes were affected by the diet in  
434 different ways. CL in liver mitochondria from one-year-old fish fed the DHA diet had almost the  
435 same FA composition than that from the FO group, while CL from the RO group had around 20  
436 % more n-6 PUFA than that from the FO group, considerably larger than the difference in n-6  
437 PUFA content between the two diets (12%). The FA compositions of PC and PE reflected more  
438 clearly the diet composition although changes in these two PL classes were attenuated  
439 compared with the diets. The influence of diet on liver mitochondrial DHA level was generally  
440 smaller than expected, considering the large differences between diets, especially between RO  
441 and DHA (3.6 vs. 22.3%). This can be in part explained by preferential retention of DHA in fish  
442 tissues when this FA is very low in the diet, as found in Atlantic salmon and other fish species  
443 (Bell *et al.* 2004, NRC 2011) and/or increased endogenous synthesis of DHA from 18:3n-3 in  
444 rainbow trout when dietary levels of n-3 LC-PUFA are reduced (Tocher *et al.* 2002). Selective  
445 incorporation of FA from the diet and biosynthesis of FA are regulatory mechanisms that have  
446 been shown in trout (Guderley *et al.* 2008). Another explanation may be an increase in the  
447 degree of oxidative stress and, particularly, lipid peroxidation that would decrease membrane  
448 levels of PUFAs (Hulbert *et al.* 2007). The observed decrease in mitochondrial total PL content  
449 in RO and especially, DHA groups could indicate the existence of high oxidative stress as was  
450 found in liver and muscle mitochondria of Atlantic salmon fed diets with a high level of n-3 LC-  
451 PUFA (Kjaer *et al.* 2008; Østbye *et al.* 2011). These studies also showed a decrease in CL  
452 content in mitochondrial membranes that agrees with data in the present study. PL classes  
453 incorporating LC-PUFA would become more susceptible to oxidative attack and would  
454 contribute to the observed reduction in total PL. Among the minor PL classes, a notable  
455 decrease in DHA content in mitochondrial PS from fish fed the RO and DHA diets was found  
456 which could be also indicating high oxidative stress since PS is the PL with the highest PI<sub>n</sub>.

457 However, these data were not consistent with lipid peroxidation results from the present study  
458 that showed no significant differences among feeding groups.

#### 459 *4.2. Diet lipid composition and mitochondrial genes*

460 The expression of mtDNA genes showed significant differences for only ND4 among  
461 feeding groups although there was a consistent trend among all the studied genes. One-year-  
462 old fish fed the RO diet tended to display higher expression of almost all the analysed genes  
463 compared to the other two groups, while trout fed the DHA diet tended to show lower values.  
464 Although only significant for ND4, the consistency of the trend among all the genes could  
465 suggest a minor effect of diet FA composition on mtDNA expression which could, over a longer  
466 time, influence the functioning of the ETC and mitochondrial respiratory rate as found in rainbow  
467 trout red muscle (Guderley *et al.* 2008). The studied genes belong to a specific region of the  
468 mitochondrial genome that has been reported to be especially prone to oxidative attack in  
469 several tissues of humans and mice (Cortopassi and Wong 1999; Vu *et al.* 2000). Genes in this  
470 area encode predominantly for subunits of complexes I [NADH-coenzyme Q (CoQ)  
471 oxidoreductase] and IV (cytochrome c oxidase) which appear to be particularly prone to age-  
472 related decline in activity in several tissues (Shigenaga *et al.* 1994; Richter 1995; Trifunovic and  
473 Larsson 2008). This is not surprising taking into consideration that 7 out of the 13 mtDNA  
474 encoded polypeptides in the ETC are found in complex I while 3 are found in complex IV.  
475 Moreover, there is a strong connection between complexes I and IV, and CL. It has been  
476 reported that CL is specifically required for electron transfer in complex I of the mitochondrial  
477 ETC (Paradies *et al.* 2002), complex I being considered the main site of ROS production in  
478 mitochondria. It is reasonable to suggest that the defect of mitochondrial complex I activity in  
479 addition to that of complex IV (very tightly bonded to CL), due to oxidation/ depletion of CL  
480 molecules and/or mutations affecting the corresponding area in mitochondrial genome, may  
481 increase electron leak from the ETC, generating more oxidative stress, mitochondrial damage  
482 and, ultimately, mitochondrial dysfunction and bioenergetics decay associated with disease and  
483 aging.

#### 484 *4.3. Fish age*

485           Some differences were observed in the effects of dietary FA composition on  
486 mitochondrial membranes and mtDNA expression between age groups. Compared with 1-year-  
487 old fish, more mature animals showed less susceptibility to diet composition, especially to DHA  
488 diet. This can be due mainly to the fact that younger fish had a higher growth rate resulting in  
489 greater incorporation of dietary lipids into their tissues, but it could also indicate age-related  
490 changes in the animal's capacity for maintaining the structural homoeostasis of mitochondrial  
491 membranes. Mitochondrial membrane PL composition was significantly different in 1- and 3-  
492 year-old fish and many of the observed changes in one age group were in the opposite direction  
493 in the other, or simply remained unchanged. Moreover, mtDNA gene expression was generally  
494 lower in 3-year-old trout fed the FO and RO diets and, although only significant for ND4, there  
495 were different trends in most of the studied genes. These observations could be associated with  
496 the well-known feature of the ageing process involving a reduction in the rate of lipid  
497 metabolism and turnover of FA (Hansford & Castro 1982). Martin *et al.* (2013) also found that  
498 several functional properties of mitochondria were affected by trout body mass which could also  
499 influence the effect of diet composition on mitochondrial processes.

#### 500           4.4. Membrane pacemaker hypothesis (MPH)

501           In conclusion, the present study showed marked changes in mitochondrial PL content  
502 and composition from trout liver when diet lipid composition was modified. These changes were  
503 PL specific and showed some regulatory mechanisms operated on mitochondrial lipids from  
504 trout liver as they did not clearly reflect differences among the diets. The main PL classes from  
505 trout mitochondria showed the PIn was higher in fish fed the DHA diet than in the RO group.  
506 Following the MPH, this would render the former more prone to oxidative attack and would  
507 explain the decrease in total PL and CL content observed in 1-year-old animals. Membranes  
508 with higher n-3 LC-PUFA content are also related with higher metabolism and ROS production  
509 (Pamplona *et al.* 2002; Sanz *et al.* 2006) which, eventually, could affect mtDNA. TBARS values,  
510 however, did not show any differences in lipid peroxidation among feeding groups. Changes  
511 observed in liver mitochondrial lipids from trout fed the RO diet could not be explained following  
512 the MPH. The three main PL classes in liver mitochondria had lower PIn and, therefore, were  
513 less susceptible to oxidative attack. In this case, fish could be suffering stress due to the

514 influence of dietary lipid composition, which has been shown to induce changes in metabolism  
515 (Tocher *et al.* 2001). Therefore, changes observed in liver mitochondrial PL content and  
516 composition from trout fed the RO diet could be reflecting the existence of compensatory  
517 mechanisms in mitochondrial membranes as a response to dietary FA composition. In both  
518 cases, mitochondrial function could be compromised by diet lipid composition which would  
519 affect animal well-being and longevity. Mitochondrial lipid composition and mtDNA expression  
520 from trout liver were affected by diet in a different way when the two age groups were  
521 compared. The observed changes could in part be explained by the different SGR of the two  
522 age groups, this being related to different fatty acid incorporation into fish tissues, but may also  
523 indicate age-related changes in the animal's capacity for maintaining the structural  
524 homoeostasis of mitochondrial membranes.

525

## 526 **Acknowledgments**

527 The authors gratefully acknowledge our colleagues Professor Gordon Bell for formulation and  
528 manufacture of the experimental feeds and Niall Auchinachie for fish husbandry. This research  
529 and P.F.A.-P. were funded by a Marie Curie Intra-European Fellowship within the 7<sup>th</sup>  
530 Community Framework Programme (PIEF-GA-2011-297964, OLDMITO). The authors report no  
531 conflicts of interest.

532

533

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673

#### 674 **Figure legends**

675 Figure 1. Specific growth rate (SGR % day<sup>-1</sup>) of 1- and 3-year-old trout fed the three  
676 experimental diets. Results are means ± SEM (n = 91 for 1- and n = 59 for 3-year-old fish). No  
677 statistical differences among feeding groups for either age group were found.

678 Figure 2. Phospholipid content (percentage of total lipid weight) and phospholipid class  
679 composition (percentage of total phospholipids) of mitochondria isolated from liver of 1- and 3-  
680 year-old rainbow trout fed three different diets. Results are means ± SEM. (n=4). Different

681 superscript letters represent differences between feeding groups for each phospholipid class as  
682 determined by two-way ANOVA ( $P<0.05$ ). Table represents  $P$  values for interaction Diet and  
683 Age for each phospholipid class ( $P<0.05$ ). Asterisks denote significant differences between age  
684 groups for each phospholipid class when compared using a Bonferroni test ( $P<0.05$ ). CL,  
685 cardiolipin; PC, phosphatidylcholine; PE, phosphatidylethanolamine; PI, phosphatidylinositol;  
686 PL, phospholipid;  $\Sigma$ PL, total phospholipids; PS, phosphatidylserine; SM, sphingomyelin. SM  
687 content found in mitochondria of 3-year-old trout fed the RO and DHA diet was lower than 0.4%.

688 Figure 3. TBARS contents (ng/g lipid) of liver mitochondria of 1- and 3-year-old rainbow trout  
689 fed the three experimental diets. Data expressed as mean  $\pm$  SEM (n=4). Letters represent the  
690 existence of statistical differences among feeding treatments for each age group as determined  
691 by a two-way ANOVA ( $P<0.05$ ). Asterisks denote significant differences between 1- and 3-year-  
692 old trout mitochondria for each feeding group when compared using a Bonferroni test ( $P<0.05$ ).  
693  $P$  value for interaction between Age and Diet was 0.430.

694 Figure 4. Relative gene expression of COX3, ND1, ND3, ND4, ND4L, ND5 and ND6 from liver  
695 mitochondria of 1- and 3-year-old rainbow trout fed the three experimental diets. Data are  
696 expressed as mean  $\pm$  SEM (n=6). Asterisks indicate significant differences between age groups  
697 for a given diet group as determined by a Bonferroni test ( $P<0.05$ ). Table presents signification  
698 values for the interaction between Age and Diet for each gene.

699

Table 1. Ingredients and feed composition by proximate analysis (n= 4)

	SHFO	RO	DHA
<i>Ingredients (g per 100 g dry weight)</i>			
Fish meal <sup>a</sup>	40.0	40.0	40.0
Soya meal <sup>a</sup>	15.0	15.0	15.0
Wheat <sup>b</sup>	10.0	10.0	10.0
Corn gluten <sup>c</sup>	10.0	10.0	10.0
Lysine	0.2	0.2	0.2
Carboxy-methyl-cellulose (CMC)	3.06	3.06	3.06
Southern hemisphere fish oil <sup>a</sup>	15.6	0.0	0.0
Rapeseed oil <sup>d</sup>	0.0	15.6	0.0
DHA oil <sup>e</sup>	0.0	0.0	15.6
Choline chloride (40% w/v)	0.4	0.4	0.4
Premixes <sup>f</sup>	5.74	5.74	5.74
<i>Proximate analysis (% dry matter)</i>			
Dry matter	85.3	82.5	88.3
Crude protein	47.1	46.7	46.9
Crude fat	24.5	24.5	23.4
Nitrogen-free extract (NFE)	17.1	17.8	18.1
Ash	11.2	10.9	11.6

<sup>a</sup> BioMar Ltd., Grangemouth, UK.

<sup>b</sup> Aquatic Feeds, Denmark.

<sup>c</sup> MP Biomedicals, LLC.

<sup>d</sup> Tesco Ltd., UK.

<sup>e</sup> Incromega TG0525. Croda International Plc, East Yorkshire, UK

<sup>f</sup> Vitamins, minerals and antioxidants, University of Stirling, UK

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

	SHFO	RO	DHA
14:0	7.5	1.3	3.3
16:0	19.9	8.8	18.4
18:0	3.7	2.0	3.9
Σsaturated	31.7	12.3	26.4
16:1n-7	8.8	1.5	5.2
18:1n-9	11.7	48.2	18.5
18:1n-7	3.3	2.6	2.7
20:1n-9	1.6	1.8	2.4
22:1n-11	1.5	1.4	2.0
ΣMonounsaturated	27.8	56.0	31.8
18:2n-6	4.7	18.5	5.6
20:3n-6	0.2	0.0	1.6
20:4n-6	1.2	0.0	0.0
22:5n-6	0.3	0.0	1.1
Σn-6 PUFA	6.8	18.5	8.3
18:3n-3	2.2	6.7	1.3
20:5n-3	13.8	2.1	6.6
22:5n-3	1.7	0.2	1.1
22:6n-3	12.1	3.6	22.3
Σn-3 PUFA	30.3	12.9	31.9
ΣPUFA	40.6	31.7	41.8
n-3/n-6	4.5	0.7	3.8
PI <sub>n</sub>	218.0	77.7	248.5

Fatty acids representing less than 1.0% of total fatty acids are not shown.

PI<sub>n</sub>, peroxidation index; PUFA, polyunsaturated fatty acids.



Table 3. Forward and reverse primers used in gene expression studies.

<i>Gene Name</i>	<i>Forward</i>	<i>Reverse</i>	<i>Amplicon Size</i>	<i>Efficiency</i>
COX 3	GTAACATGAGCCCACCACAG	CGACAAAGAAAGTAGAGCCGT	168	92.3%
ND1	CAACGTAGCCCAAGAAAGCA	ACTAATTCTGACTCTCCTTCTGTG	128	100.5%
ND3	CTATTACCATCACATTATCCGCAG	GAAAGAAGCGTAAGGAGAAGGG	146	96.5%
ND4	TGAACTACATCAATCGCCCA	GTTGCTAAATAGAGGTTGGAGG	106	97.2%
ND4L	CTCTCAGCCCTTCTATGCCT	AACGCTAGGAGAAGTATCGGG	190	100.5%
ND5	CCTATTGCCCTGTATGTAACC	ATGATATAATTCCGACTCCCTCTC	183	92.3%
ND6	ACTCCTTAAACTCGTCCACTG	GGGATGCTTGTGGTATTTGCT	183	92.3%

Table 4. Fatty acid composition (percentage of total fatty acids) of phosphatidylcholine of mitochondria isolated from liver of 1- and 3-year old rainbow trout fed with one of three experimental diets.

Fatty acid	1 year			3 years			Age*Diet
	SHFO	RO	DHA	SHFO	RO	DHA	<i>P</i>
14:0	2.5±0.1 <sup>b*</sup>	1.0±0.1 <sup>a</sup>	1.2±0.1 <sup>a*</sup>	2.1±0.4 <sup>c</sup>	1.1±0.1 <sup>a</sup>	1.6±0.1 <sup>b</sup>	0.003
16:0	23.0±0.4 <sup>c*</sup>	18.3±0.7 <sup>a</sup>	20.8±1.3 <sup>b</sup>	19.4±0.3 <sup>a</sup>	18.2±0.6 <sup>a</sup>	21.8±0.8 <sup>b</sup>	<0.001
18:0	4.4±0.3 <sup>b</sup>	3.3±0.3 <sup>a</sup>	3.7±0.2 <sup>a*</sup>	4.9±0.5	3.5±1.1	5.1±0.8	0.166
∑Saturated	30.6±0.4 <sup>c*</sup>	22.9±0.8 <sup>a</sup>	26.4±1.4 <sup>b*</sup>	26.8±0.4 <sup>b</sup>	23.2±1.6 <sup>a</sup>	29.2±0.2 <sup>c</sup>	<0.001
16:1n-7	2.6±0.1 <sup>c*</sup>	0.9±0.1 <sup>a</sup>	1.9±0.2 <sup>b</sup>	2.0±0.3 <sup>b</sup>	1.0±0.1 <sup>a</sup>	2.0±0.1 <sup>b</sup>	<0.001
18:1n-9	6.6±0.3 <sup>a</sup>	11.5±0.9 <sup>b*</sup>	7.2±0.6 <sup>a</sup>	7.4±1.3 <sup>a</sup>	12.9±0.3 <sup>b</sup>	7.6±0.4 <sup>a</sup>	0.359
18:1n-7	1.8±0.1	1.1±0.1	1.2±0.1	3.2±1.2	1.9±0.7	2.3±0.7	0.599
20:1n-9	0.9±0.1 <sup>a*</sup>	2.3±0.3 <sup>b</sup>	1.0±0.1 <sup>a*</sup>	2.0±0.6	2.8±0.7	1.9±0.3	0.277
∑Monounsaturated	12.8±0.5 <sup>a</sup>	16.9±1.4 <sup>b</sup>	12.2±1.1 <sup>a*</sup>	15.5±2.8 <sup>ab</sup>	19.5±1.3 <sup>b</sup>	14.7±1.0 <sup>a</sup>	0.994
18:2n-6	1.1±0.0 <sup>a</sup>	5.4±0.4 <sup>b</sup>	1.2±0.1 <sup>a</sup>	1.7±0.5 <sup>a</sup>	6.0±1.0 <sup>b</sup>	1.4±0.2 <sup>a</sup>	0.523
20:2n-6	0.5±0.0 <sup>a</sup>	2.1±0.1 <sup>b*</sup>	0.6±0.1 <sup>a</sup>	0.6±0.1 <sup>a</sup>	1.6±0.2 <sup>b</sup>	0.6±0.1 <sup>a</sup>	0.001
20:3n-6	0.2±0.0 <sup>a</sup>	1.7±0.2 <sup>b</sup>	0.1±0.0 <sup>a</sup>	0.3±0.0 <sup>a</sup>	1.7±0.3 <sup>b</sup>	0.2±0.0 <sup>a</sup>	0.957
20:4n-6	1.4±0.0 <sup>a</sup>	1.7±0.2 <sup>a</sup>	2.7±0.1 <sup>b</sup>	1.3±0.1 <sup>a</sup>	1.7±0.5 <sup>a</sup>	2.7±0.1 <sup>b</sup>	0.965
22:5n-6	0.4±0.0 <sup>a</sup>	0.5±0.1 <sup>a</sup>	1.4±0.1 <sup>b</sup>	0.3±0.1 <sup>a</sup>	0.4±0.1 <sup>a</sup>	1.4±0.4 <sup>b</sup>	0.558
∑n-6 PUFA	3.9±0.0 <sup>a</sup>	11.6±0.7 <sup>c</sup>	6.2±0.5 <sup>b</sup>	4.3±0.4 <sup>a</sup>	11.7±1.4 <sup>c</sup>	6.6±0.2 <sup>b</sup>	0.927
20:5n-3	10.1±0.3 <sup>c</sup>	4.8±0.1 <sup>a*</sup>	5.7±0.3 <sup>b</sup>	11.6±1.2 <sup>b</sup>	7.0±0.7 <sup>a</sup>	6.8±0.9 <sup>a</sup>	<0.001
22:5n-3	1.8±0.1 <sup>b*</sup>	0.9±0.1 <sup>a</sup>	1.0±0.0 <sup>a</sup>	2.5±0.2 <sup>b</sup>	1.2±0.3 <sup>a</sup>	1.4±0.3 <sup>a</sup>	0.176
22:6n-3	39.7±0.5 <sup>a</sup>	41.0±1.2 <sup>a*</sup>	47.0±1.9 <sup>b*</sup>	38.0±1.7 <sup>ab</sup>	35.1±2.2 <sup>a</sup>	40.0±2.1 <sup>b</sup>	0.024
∑n-3 PUFA	52.3±0.6 <sup>b</sup>	48.3±1.3 <sup>a*</sup>	54.4±2.1 <sup>b*</sup>	53.3±2.9 <sup>c</sup>	45.4±1.9 <sup>a</sup>	49.0±0.8 <sup>b</sup>	0.008
∑PUFA	56.5±0.5 <sup>a</sup>	60.2±0.8 <sup>b</sup>	61.4±2.0 <sup>b*</sup>	57.8±2.6	57.2±2.7	56.1±1.1	0.008
∑n-3 LC-PUFA	51.8±0.5 <sup>b</sup>	46.8±1.2 <sup>a</sup>	53.4±2.0 <sup>b*</sup>	52.8±2.0 <sup>c</sup>	44.0±2.0 <sup>a</sup>	49.1±0.5 <sup>b</sup>	0.009
n-3/n-6	13.3±0.2 <sup>c</sup>	4.1±0.3 <sup>a</sup>	8.3±0.1 <sup>b*</sup>	12.7±2.1 <sup>c</sup>	4.1±0.4 <sup>a</sup>	7.5±0.2 <sup>b</sup>	0.482
PI <sub>n</sub>	400.3±4.6 <sup>b</sup>	380.9±0.1 <sup>a*</sup>	442.4±21.0 <sup>c*</sup>	405.9±30.4 <sup>b</sup>	347.8±17.3 <sup>a</sup>	402.0±10.3 <sup>b</sup>	0.015

Data expressed as mean ± SD (n=4). Different superscript letters within a row and for each age group represent significant differences between diet treatments as determined by a two-way ANOVA (*p*<0.05). Asterisks denote statistical differences between 1- and 3-year-old trout for each diet group (SHFO, RO and DHA) when

compared using a Bonferroni test ( $p < 0.05$ ). Right column represent signification values for the interaction between Diet and Age ( $p < 0.05$ ). SHFO, fish oil diet group; RO, rapeseed oil group; DHA, Incromega TG0525 oil group; LC-PUFA, long-chain polyunsaturated fatty acids; PIn, peroxidation index; PUFA, polyunsaturated fatty acids. Fatty acids representing less than 1% of total fatty acids are not shown.

Table 5. Fatty acid composition (percentage of total fatty acids) of phosphatidylethanolamine of mitochondria isolated from liver of 1- and 3-year old rainbow trout fed with one of three experimental diets.

Fatty acid	1 year			3 years			Age*Diet
	SHFO	RO	DHA	SHFO	RO	DHA	<i>P</i>
16:0	10.1±0.6	9.8±0.5 <sup>*</sup>	9.4±0.6	10.3±1.3 <sup>b</sup>	8.1±0.5 <sup>a</sup>	10.4±0.5 <sup>b</sup>	0.01
18:0	9.3±0.2 <sup>b*</sup>	7.0±0.4 <sup>a*</sup>	8.8±0.8 <sup>b</sup>	7.9±1.1 <sup>b</sup>	5.8±0.3 <sup>a</sup>	9.2±1.0 <sup>b</sup>	0.046
∑Saturated	20.0±0.6 <sup>b</sup>	17.3±0.9 <sup>a*</sup>	18.7±0.7 <sup>b*</sup>	18.8±0.7 <sup>b</sup>	14.4±0.6 <sup>a</sup>	20.2±0.4 <sup>b</sup>	<0.001
16:1n-7	1.1±0.0 <sup>c</sup>	0.3±0.1 <sup>a</sup>	0.6±0.0 <sup>b</sup>	1.0±0.4 <sup>b</sup>	0.3±0.0 <sup>a</sup>	0.7±0.1 <sup>b</sup>	0.416
18:1n-9	8.0±0.5 <sup>a</sup>	12.9±0.4 <sup>b</sup>	8.3±1.1 <sup>a</sup>	8.0±0.6 <sup>a</sup>	13.1±0.5 <sup>b</sup>	7.9±1.0 <sup>a</sup>	0.746
18:1n-7	5.1±0.3 <sup>c</sup>	2.3±0.0 <sup>a</sup>	3.0±0.3 <sup>b*</sup>	5.8±1.3 <sup>b</sup>	2.9±0.3 <sup>a</sup>	5.5±1.0 <sup>b</sup>	0.035
20:1n-9	2.3±0.2 <sup>a*</sup>	3.9±0.2 <sup>b</sup>	2.4±0.4 <sup>a*</sup>	3.6±0.4	3.9±0.3	3.7±0.4	0.004
∑Monounsaturated	17.5±0.8 <sup>b</sup>	19.8±0.4 <sup>c</sup>	14.7±1.8 <sup>a*</sup>	19.1±1.4	20.8±0.3	18.3±1.0	0.128
18:2n-6	2.4±0.1 <sup>a</sup>	5.1±0.6 <sup>b*</sup>	2.6±0.2 <sup>a</sup>	3.0±1.0 <sup>a</sup>	6.5±0.1 <sup>b</sup>	2.8±0.3 <sup>a</sup>	0.122
20:2n-6	1.2±0.2 <sup>a</sup>	2.7±0.1 <sup>b*</sup>	1.3±0.2 <sup>a</sup>	0.9±0.2 <sup>a</sup>	2.1±0.3 <sup>c</sup>	1.3±0.1 <sup>b</sup>	0.016
20:4n-6	2.2±0.1 <sup>a</sup>	2.6±0.3 <sup>a</sup>	3.8±0.2 <sup>b*</sup>	2.9±1.4 <sup>a</sup>	2.6±0.9 <sup>a</sup>	6.4±1.9 <sup>b</sup>	0.046
∑n-6 PUFA	7.3±0.3 <sup>a</sup>	12.5±0.6 <sup>c</sup>	10.0±0.4 <sup>b*</sup>	7.7±1.5 <sup>a</sup>	13.5±1.5 <sup>b</sup>	12.8±1.5 <sup>b</sup>	0.129
20:5n-3	4.9±0.1 <sup>c*</sup>	3.1±0.3 <sup>b</sup>	2.1±0.1 <sup>a*</sup>	9.6±4.1 <sup>b</sup>	4.7±1.0 <sup>a</sup>	7.5±3.7 <sup>ab</sup>	0.229
22:5n-3	1.2±0.2 <sup>b</sup>	0.5±0.1 <sup>a</sup>	0.9±0.2 <sup>b</sup>	1.4±0.3 <sup>b</sup>	0.7±0.1 <sup>a</sup>	1.0±0.1 <sup>ab</sup>	0.653
22:6n-3	47.4±0.4 <sup>b*</sup>	44.9±1.1 <sup>a</sup>	51.8±2.7 <sup>c*</sup>	41.7±5.8	43.4±0.6	38.8±6.0	0.019
∑n-3 PUFA	54.6±0.7 <sup>b</sup>	49.8±1.0 <sup>a</sup>	55.6±2.7 <sup>b*</sup>	54.0±1.5 <sup>b</sup>	50.7±0.8 <sup>ab</sup>	48.3±2.4 <sup>a</sup>	0.001
∑PUFA	62.5±0.5 <sup>a</sup>	63.0±0.8 <sup>a</sup>	66.6±2.3 <sup>b*</sup>	62.1±1.0 <sup>ab</sup>	64.8±0.8 <sup>b</sup>	61.5±1.2 <sup>a</sup>	0.001
∑n-3 LC-PUFA	54.3±0.8 <sup>b</sup>	49.4±1.3 <sup>a</sup>	56.2±2.6 <sup>b*</sup>	53.5±1.5 <sup>b</sup>	49.8±0.7 <sup>ab</sup>	47.9±2.3 <sup>a</sup>	0.001
n-3/n-6	7.6±0.3 <sup>c</sup>	4.0±0.3 <sup>a</sup>	5.7±0.5 <sup>b*</sup>	7.2±1.5 <sup>b</sup>	3.8±0.5 <sup>a</sup>	3.8±0.6 <sup>a</sup>	0.106
PI <sub>n</sub>	438.3±7.7 <sup>a</sup>	416.1±10.6 <sup>a</sup>	472.5±27.9 <sup>b*</sup>	417.9±17.5	412.3±4.1	406.5±21.4	0.003

Data expressed as mean ± SD (n=4). Different superscript letters within a row and for each age group represent significant differences between diet treatments as determined by a two-way ANOVA ( $p < 0.05$ ). Asterisks denote statistical differences between 1- and 3-year-old trout for each diet group (SHFO, RO and DHA) when compared using a Bonferroni test ( $p < 0.05$ ). Right column represent signification values for the interaction between Diet and Age ( $p < 0.05$ ). SHFO, fish oil diet group; RO, rapeseed oil group; DHA, Incromega TG0525 oil group; LC-PUFA, long-chain polyunsaturated fatty acids; PI<sub>n</sub>, peroxidation index; PUFA, polyunsaturated fatty acids. Fatty acids representing less than 1% of total fatty acids are not shown.

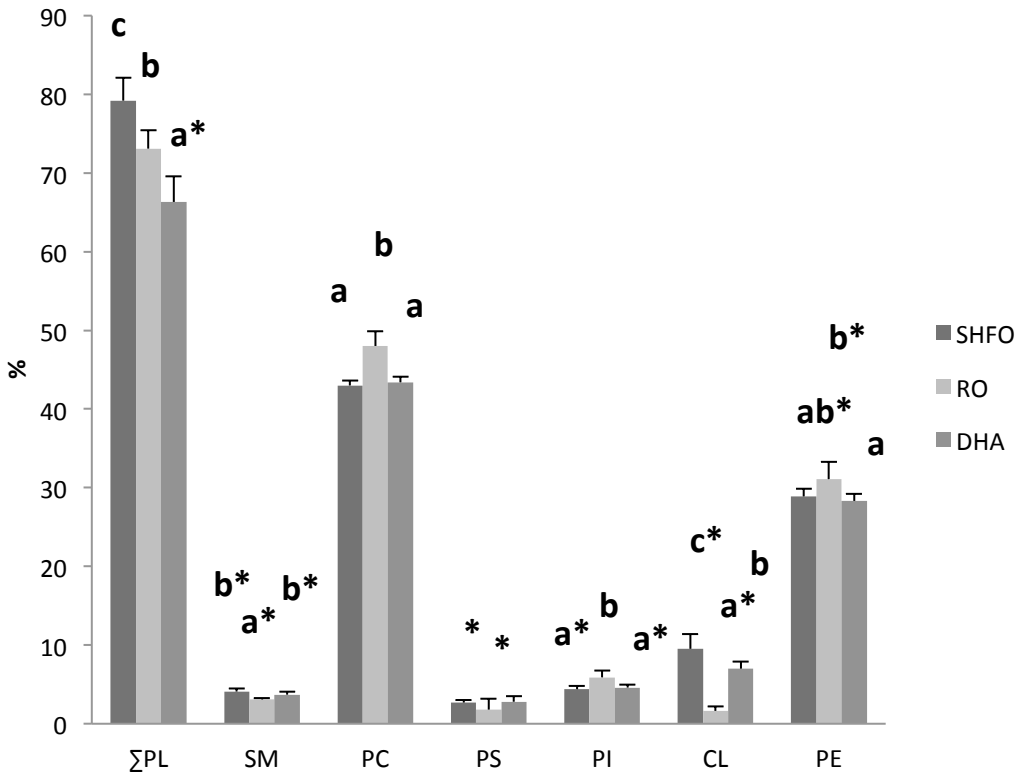
Table 6. Fatty acid composition (percentage of total fatty acids) of cardiolipin of mitochondria isolated from liver of 1- and 3-year old rainbow trout fed with one of three experimental diets.

Fatty acid	1 year			3 years			Age*Diet
	SHFO	RO	DHA	SHFO	RO	DHA	<i>P</i>
14:0	2.3±0.1 <sup>c*</sup>	0.9±0.2 <sup>a</sup>	1.2±0.1 <sup>b</sup>	1.8±0.1 <sup>c</sup>	0.9±0.2 <sup>a</sup>	1.3±0.1 <sup>b</sup>	0.001
16:0	8.7±1.1 <sup>b</sup>	5.7±1.2 <sup>a*</sup>	9.9±1.4 <sup>b</sup>	8.3±0.6	8.8±1.9	10.2±1.4	0.053
18:0	1.4±0.3	1.2±0.4	2.5±1.4	1.5±0.2	2.6±1.5	2.4±1.2	0.347
∑saturated	12.9±1.3 <sup>b</sup>	8.1±1.9 <sup>a*</sup>	14.2±2.7 <sup>b</sup>	12.0±0.8	12.7±3.4	14.5±2.5	0.81
16:1n-7	7.4±0.5 <sup>c*</sup>	1.4±0.2 <sup>a*</sup>	4.4±0.6 <sup>b</sup>	5.7±0.8 <sup>b</sup>	0.0±0.0 <sup>a</sup>	4.7±0.6 <sup>b</sup>	0.016
18:1n-9	6.3±0.1	7.8±0.6	7.0±0.2	6.6±1.0 <sup>a</sup>	9.7±1.5 <sup>b</sup>	7.3±0.4 <sup>a</sup>	0.109
18:1n-7	4.0±0.3 <sup>b</sup>	1.6±0.3 <sup>a*</sup>	3.5±0.2 <sup>b</sup>	4.0±0.6 <sup>b</sup>	2.4±0.3 <sup>a</sup>	4.3±0.9 <sup>b</sup>	0.239
20:1n-9	1.0±0.1 <sup>*</sup>	1.4±0.1 <sup>*</sup>	1.3±0.1	1.7±0.3	2.2±0.2	1.7±0.4	0.36
∑monounsaturated	19.5±0.2 <sup>b</sup>	13.5±1.2 <sup>a*</sup>	17.3±1.6 <sup>b</sup>	18.5±0.9	16.5±2.7	18.9±0.7	0.056
18:2n-6	6.5±0.4 <sup>a*</sup>	21.1±0.9 <sup>b*</sup>	7.4±1.3 <sup>a</sup>	8.6±0.3 <sup>a</sup>	18.0±1.7 <sup>b</sup>	8.4±1.7 <sup>a</sup>	0.002
20:2n-6	3.9±0.3 <sup>a*</sup>	6.3±0.5 <sup>b*</sup>	5.6±0.8 <sup>b</sup>	2.8±0.2 <sup>a</sup>	4.8±0.9 <sup>b</sup>	4.4±0.8 <sup>b</sup>	0.899
20:3n-6	0.9±0.1 <sup>a</sup>	4.8±0.6 <sup>b</sup>	0.7±0.1 <sup>a</sup>	1.2±0.4 <sup>a</sup>	3.9±0.5 <sup>b</sup>	0.9±0.1 <sup>a</sup>	0.01
20:4n-6	0.8±0.2 <sup>a</sup>	0.7±0.3 <sup>a</sup>	1.7±0.5 <sup>b</sup>	1.0±0.1 <sup>a</sup>	1.2±0.1 <sup>a</sup>	1.8±0.4 <sup>b</sup>	0.616
∑n-6 PUFA	12.9±0.5 <sup>a</sup>	33.2±1.0 <sup>b*</sup>	16.2±1.5 <sup>a</sup>	14.1±0.8 <sup>a</sup>	28.2±2.8 <sup>b</sup>	16.4±1.9 <sup>a</sup>	0.04
18:3n-3	1.0±0.1 <sup>a*</sup>	3.1±0.2 <sup>b</sup>	1.1±0.2 <sup>a</sup>	2.1±0.1 <sup>ab</sup>	2.8±0.4 <sup>b</sup>	1.5±0.6 <sup>a</sup>	0.007
20:4n-3	1.2±0.1 <sup>b*</sup>	1.2±0.2 <sup>b</sup>	0.6±0.1 <sup>a</sup>	2.2±0.2 <sup>b</sup>	1.3±0.3 <sup>a</sup>	1.0±0.1 <sup>a</sup>	0.005
20:5n-3	1.6±0.2 <sup>b*</sup>	0.6±0.0 <sup>a*</sup>	1.0±0.4 <sup>ab</sup>	3.0±0.4 <sup>b</sup>	1.2±0.4 <sup>a</sup>	1.2±0.3 <sup>a</sup>	0.007
22:5n-3	3.5±0.5 <sup>b</sup>	1.6±0.2 <sup>a</sup>	1.3±0.3 <sup>a*</sup>	3.2±0.4 <sup>b</sup>	1.8±0.5 <sup>a</sup>	2.2±0.3 <sup>a</sup>	0.032
22:6n-3	45.6±0.7 <sup>b</sup>	37.6±1.8 <sup>a</sup>	46.4±0.8 <sup>b*</sup>	43.2±1.2 <sup>b</sup>	34.1±3.0 <sup>a</sup>	42.4±2.7 <sup>b</sup>	0.732
∑n-3 PUFA	53.8±1.1 <sup>b</sup>	45.0±2.1 <sup>a</sup>	51.4±0.4 <sup>b</sup>	54.7±1.0 <sup>c</sup>	42.2±4.1 <sup>a</sup>	49.3±2.2 <sup>b</sup>	0.305
Total PUFA	67.7±1.4 <sup>a</sup>	78.4±3.1 <sup>b</sup>	68.5±1.4 <sup>a</sup>	69.5±0.3	70.7±5.5	66.5±1.9	0.021
∑n-3 LC-PUFA	52.7±1.1 <sup>b</sup>	42.6±1.0 <sup>a</sup>	50.2±0.7 <sup>b</sup>	52.1±0.9 <sup>b</sup>	40.5±3.1 <sup>a</sup>	48.6±1.7 <sup>b</sup>	0.58
n-3/n-6	4.2±0.2 <sup>c</sup>	1.4±0.0 <sup>a</sup>	3.2±0.3 <sup>b</sup>	3.9±0.3 <sup>b</sup>	1.6±0.1 <sup>a</sup>	3.2±0.3 <sup>b</sup>	0.279
PIIn	421.8±7.0 <sup>b</sup>	376.7±10.6 <sup>a</sup>	423.1±2.3 <sup>b*</sup>	416.1±8.2 <sup>b</sup>	365.5±17.7 <sup>a</sup>	399.4±5.5 <sup>b</sup>	0.464

Data expressed as mean ± SD (n=4). Different superscript letters within a row and for each age group represent significant differences between diet treatments as determined by a two-way ANOVA ( $p < 0.05$ ). Asterisks denote statistical differences between 1- and 3-year-old trout for each diet group (SHFO, RO and DHA) when

compared using a Bonferroni test ( $p < 0.05$ ). Right column represent signification values for the interaction between Diet and Age ( $p < 0.05$ ). SHFO, fish oil diet group; RO, rapeseed oil group; DHA, Incromega TG0525 oil group; LC-PUFA, long-chain polyunsaturated fatty acids; PIn, peroxidation index; PUFA, polyunsaturated fatty acids. Fatty acids representing less than 1% of total fatty acids are not shown.

# 1 YEAR



	Age*Diet <i>P</i>
ΣPL	<0.001
SM	<0.001
PC	<0.001
PS	0.131
PI	0.033
CL	<0.001
PE	0.002

# 3 YEARS

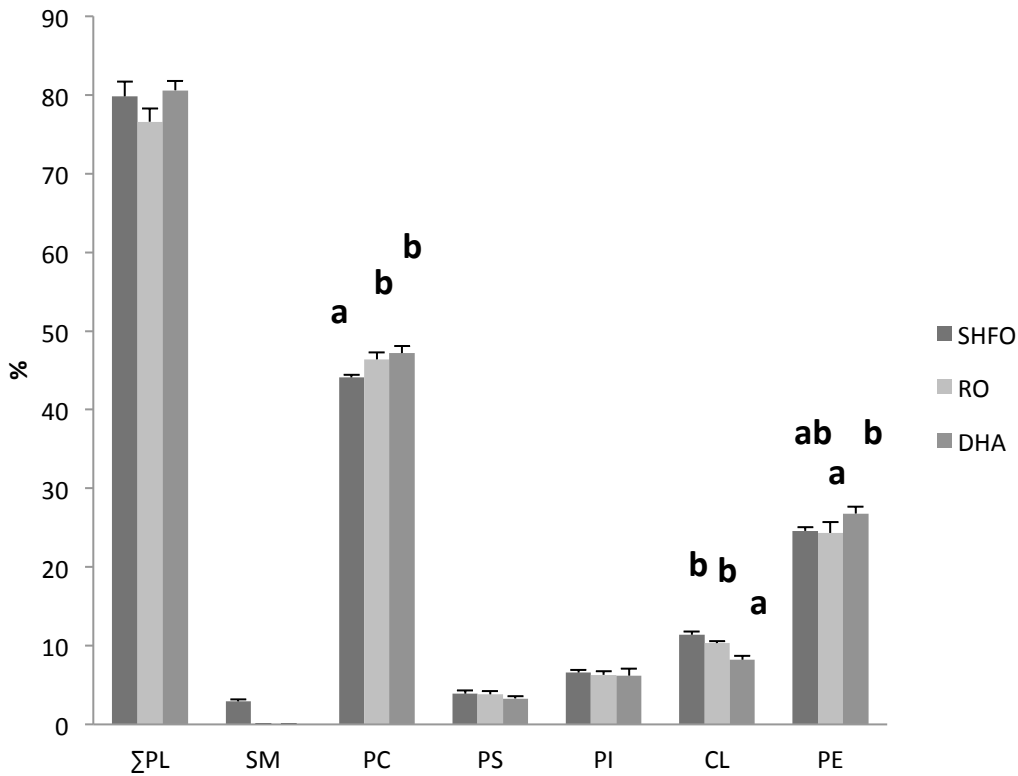


Figure 1. Almáida-Pagán *et al.*

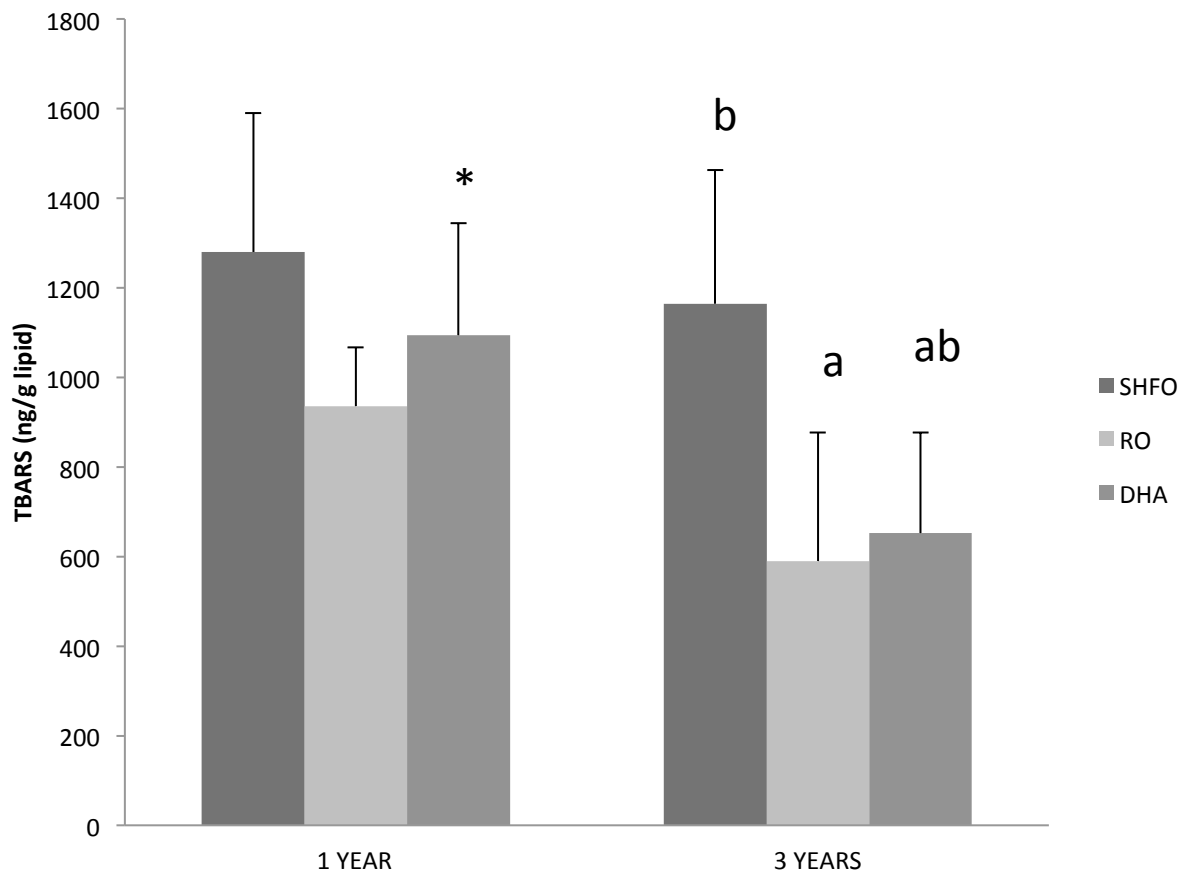


Figure 2. Almaida-Pagán *et al.*



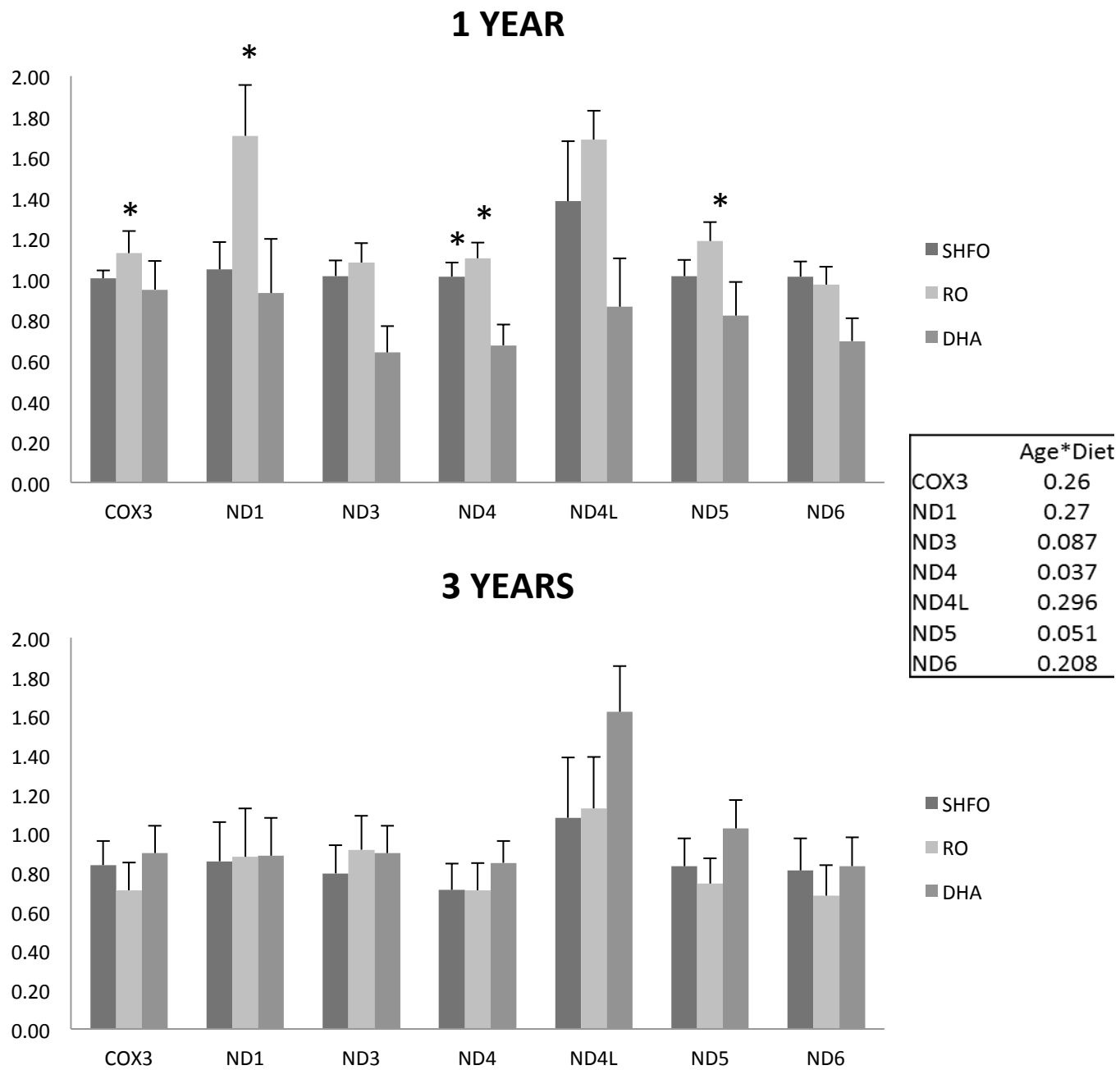


Figure 3. Almaida-Pagán *et al.*

Supplementary Table 1. Fatty acid composition (percentage of total fatty acids) of phosphatidylinositol of mitochondria isolated from liver of 1- and 3-year old rainbow trout fed with one of three experimental diets.

Fatty acid	Liver			3 years			Age*Diet
	SHFO	RO	DHA	SHFO	RO	DHA	<i>P</i>
16:0	4.3±0.3*	5.1±1.1	3.8±0.2*	9.2±1.2 <sup>b</sup>	4.7±0.7 <sup>a</sup>	5.6±1.2 <sup>a</sup>	<0.001
18:0	35.0±0.9*	30.4±1.5*	33.1±1.5*	23.2±4.4	26.8±1.9	28.4±3.2	0.008
∑Saturated	40.3±0.8*	36.5±2.7	37.7±1.5	33.7±3.2	32.2±1.5	34.7±2.1	0.239
18:1n-9	4.2±0.1 <sup>a*</sup>	8.6±1.2 <sup>b</sup>	5.4±0.8 <sup>a</sup>	9.4±1.9 <sup>b</sup>	10.5±0. <sup>b</sup>	6.7±1.6 <sup>a</sup>	0.009
18:1n-7	1.0±0.1*	0.7±0.1	0.9±0.1*	2.0±0.5 <sup>b</sup>	1.0±0.2 <sup>a</sup>	1.4±0.3 <sup>ab</sup>	0.025
20:1n-9	1.0±0.1*	1.5±0.2*	1.2±0.2*	2.4±0.5	2.2±0.3	1.9±0.2	0.029
∑Monounsaturated	7.1±0.2 <sup>a*</sup>	12.6±2.2 <sup>b</sup>	8.6±1.5 <sup>a</sup>	16.4±3.0 <sup>b</sup>	14.2±1.0 <sup>b</sup>	10.8±2.1 <sup>a</sup>	0.001
18:2n-6	0.4±0.0 <sup>a*</sup>	1.4±0.3 <sup>b</sup>	0.6±0.2 <sup>a</sup>	1.3±0.4 <sup>ab</sup>	1.5±0.2 <sup>b</sup>	0.8±0.2 <sup>a</sup>	0.011
20:4n-6	34.5±2.3 <sup>a</sup>	32.9±3.8 <sup>a</sup>	41.1±1.2 <sup>b</sup>	31.4±4.2 <sup>a</sup>	34.9±2.2 <sup>a</sup>	40.0±1.2 <sup>b</sup>	0.197
∑n-6 PUFA	36.0±2.3 <sup>a</sup>	39.0±4.1 <sup>ab</sup>	43.2±0.9 <sup>b</sup>	35.0±4.2 <sup>a</sup>	41.3±0.6 <sup>b</sup>	42.7±1.2 <sup>b</sup>	0.438
20:5n-3	1.1±0.1 <sup>b*</sup>	0.8±0.1 <sup>b*</sup>	0.4±0.1 <sup>a*</sup>	2.2±0.4 <sup>b</sup>	1.5±0.5 <sup>ab</sup>	0.8±0.3 <sup>a</sup>	0.036
22:5n-3	0.8±0.0 <sup>b*</sup>	0.3±0.1 <sup>a*</sup>	0.3±0.0 <sup>a*</sup>	1.4±0.3 <sup>b</sup>	0.5±0.2 <sup>a</sup>	0.5±0.1 <sup>a</sup>	0.374
22:6n-3	13.8±3.0 <sup>b</sup>	9.4±3.7 <sup>ab</sup>	9.1±0.5 <sup>a</sup>	10.3±1.2	9.6±0.2	10.1±0.4	0.081
∑n-3 PUFA	15.8±3.0 <sup>b</sup>	10.8±3.7 <sup>ab</sup>	10.0±0.4 <sup>a</sup>	14.4±1.4 <sup>b</sup>	12.0±0.5 <sup>a</sup>	11.7±0.3 <sup>a</sup>	0.302
∑PUFA	52.6±0.8	50.9±0.6*	53.8±0.2	50.0±3.8	53.5±0.5	54.6±0.9	0.015
∑n-3 LC-PUFA	16.3±3.4	9.0±2.5*	9.7±0.5*	14.1±1.6 <sup>b</sup>	12.0±0.1 <sup>a</sup>	11.5±0.3 <sup>a</sup>	0.003
n-3/n-6	0.5±0.1 <sup>b</sup>	0.2±0.1 <sup>a</sup>	0.2±0.0 <sup>a</sup>	0.4±0.1	0.3±0.0	0.3±0.0	0.744
PI <sub>n</sub>	274.0±19.2 <sup>c*</sup>	213.8±8.1 <sup>a*</sup>	248.8±3.1 <sup>b*</sup>	243.6±10.8 <sup>a</sup>	240.3±7.4 <sup>a</sup>	258.5±0.8 <sup>b</sup>	0.016

Data expressed as mean ± SD (n=4). Different superscript letters within a row and for each age group represent significant differences between diet treatments as determined by a two-way ANOVA ( $p < 0.05$ ). Asterisks denote statistical differences between 1- and 3-year-old trout for each diet group (SHFO, RO and DHA) when compared using a Bonferroni test ( $p < 0.05$ ). Right column represent signification values for the interaction between Diet and Age ( $p < 0.05$ ). SHFO, fish oil diet group; RO, rapeseed oil group; DHA, Incromega TG0525 oil group; LC-PUFA, long-chain polyunsaturated fatty acids; PI<sub>n</sub>, peroxidation index; PUFA, polyunsaturated fatty acids. Fatty acids representing less than 1% of total fatty acids are not shown.

Supplementary Table 2. Fatty acid composition (percentage of total fatty acids) of phosphatidylserine of mitochondria isolated from liver of 1- and 3-year old rainbow trout fed with one of three experimental diets.

Fatty acid	Liver			3 years			Age*Diet
	SHFO	RO	DHA	SHFO	RO	DHA	<i>P</i>
16:0	8.5±1.4 <sup>a*</sup>	13.5±1.7 <sup>b*</sup>	10.6±0.9 <sup>a</sup>	12.3±2.4	10.4±1.5	11.8±0.9	0.002
18:0	9.0±1.5 <sup>a*</sup>	21.5±1.1 <sup>b</sup>	25.2±0.9 <sup>b*</sup>	15.9±1.4 <sup>a</sup>	18.9±4.2 <sup>ab</sup>	21.0±1.0 <sup>b</sup>	<0.001
∑Saturated	18.6±2.4 <sup>a*</sup>	36.8±2.4 <sup>b</sup>	37.2±0.9 <sup>b</sup>	30.3±3.2	30.9±5.9	34.2±0.8	<0.001
18:1n-9	2.9±0.5 <sup>a*</sup>	5.2±0.0 <sup>b*</sup>	2.8±0.2 <sup>a</sup>	4.2±0.6 <sup>b</sup>	4.5±0.5 <sup>b</sup>	3.2±0.3 <sup>a</sup>	0.001
18:1n-7	1.3±0.3 <sup>*</sup>	1.2±0.1	1.3±0.1	2.7±0.9 <sup>b</sup>	1.4±0.1 <sup>a</sup>	1.9±0.6 <sup>ab</sup>	0.068
20:1n-9	0.7±0.2 <sup>a*</sup>	2.0±0.5 <sup>b</sup>	1.2±0.2 <sup>a</sup>	1.9±0.6	1.9±0.2	1.6±0.4	0.017
∑Monounsaturated	6.7±1.2 <sup>a*</sup>	10.9±1.1 <sup>b*</sup>	6.9±0.6 <sup>a</sup>	10.7±1.8	8.9±0.6	7.9±1.3	0.001
18:2n-6	0.6±0.2 <sup>a</sup>	1.5±0.1 <sup>b</sup>	0.4±0.1 <sup>a</sup>	0.6±0.3 <sup>a</sup>	1.3±0.2 <sup>b</sup>	0.5±0.1 <sup>a</sup>	0.515
20:4n-6	1.3±0.2	0.7±0.2	0.9±0.1 <sup>*</sup>	1.2±0.4	0.9±0.4	1.6±0.4	0.128
22:5n-6	0.8±0.0 <sup>a</sup>	0.7±0.0 <sup>a</sup>	2.0±0.1 <sup>b</sup>	0.5±0.1 <sup>a</sup>	0.8±0.2 <sup>a</sup>	2.4±0.3 <sup>b</sup>	0.009
∑n-6 PUFA	3.4±0.4 <sup>a</sup>	6.1±2.0 <sup>b</sup>	4.4±0.2 <sup>b*</sup>	3.4±0.5 <sup>a</sup>	5.1±0.6 <sup>b</sup>	6.2±0.4 <sup>b</sup>	0.015
20:5n-3	8.2±0.7 <sup>b*</sup>	1.0±0.2 <sup>a</sup>	0.7±0.1 <sup>a</sup>	5.0±0.4 <sup>b</sup>	2.4±1.1 <sup>a</sup>	1.9±0.6 <sup>a</sup>	0.001
22:5n-3	3.1±0.3 <sup>b</sup>	0.6±0.0 <sup>a</sup>	0.4±0.0 <sup>a</sup>	3.0±0.6 <sup>b</sup>	1.3±0.9 <sup>a</sup>	0.9±0.3 <sup>a</sup>	0.306
22:6n-3	58.9±3.4 <sup>c*</sup>	42.1±1.3 <sup>a*</sup>	48.5±0.4 <sup>b</sup>	46.3±4.9	50.0±3.8	48.2±2.7	<0.001
∑n-3 PUFA	70.8±3.7 <sup>c*</sup>	44.2±0.9 <sup>a*</sup>	50.0±0.4 <sup>b</sup>	54.6±5.0	54.5±6.6	51.4±2.0	<0.001
∑PUFA	74.7±3.4 <sup>b*</sup>	52.2±1.3 <sup>a*</sup>	55.8±0.5 <sup>a</sup>	59.0±3.8	60.2±4.5	57.9±1.7	<0.001
∑n-3 LC-PUFA	70.6±4.6 <sup>c*</sup>	43.9±1.1 <sup>a*</sup>	49.9±0.4 <sup>b</sup>	54.4±4.8	55.8±4.2	52.0±0.9	<0.001
n-3/n-6	21.2±4.0 <sup>c</sup>	7.7±2.1 <sup>b</sup>	11.4±0.7 <sup>a*</sup>	16.4±4.0 <sup>b</sup>	11.4±2.2 <sup>ab</sup>	8.7±0.5 <sup>a</sup>	0.018
PI <sub>n</sub>	547.7±46.5 <sup>c*</sup>	361.3±4.6 <sup>a*</sup>	417.7±1.4 <sup>b</sup>	427.9±49.0	448.6±73.8	434.6±13.4	<0.001

Data expressed as mean ± SD (n=4). Different superscript letters within a row and for each age group represent significant differences between diet treatments as determined by a two-way ANOVA ( $p < 0.05$ ). Asterisks denote statistical differences between 1- and 3-year-old trout for each diet group (SHFO, RO and DHA) when compared using a Bonferroni test ( $p < 0.05$ ). Right column represent signification values for the interaction between Diet and Age ( $p < 0.05$ ). SHFO, fish oil diet group; RO, rapeseed oil group; DHA, Incromege TG0525 oil group; LC-PUFA, long-chain polyunsaturated fatty acids; PI<sub>n</sub>, peroxidation index; PUFA, polyunsaturated fatty acids.

Fatty acids representing less than 1% of total fatty acids are not shown.

Supplementary Table 3. Fatty acid composition (percentage of total fatty acids) of sphingomyelin of mitochondria isolated from liver of 1- and 3-year old rainbow trout fed with one of three experimental diets.

Fatty acid	Liver			3 years			Age*Diet
	SHFO	RO	DHA	SHFO	RO	DHA	<i>P</i>
14:0	4.3±0.4 <sup>b*</sup>	1.8±0.2 <sup>a*</sup>	2.1±0.2 <sup>a*</sup>	6.0±0.9 <sup>b</sup>	3.8±1.2 <sup>a</sup>	3.7±0.5 <sup>a</sup>	0.901
16:0	19.3±2.5 <sup>a</sup>	24.1±1.7 <sup>b*</sup>	25.2±1.5 <sup>b*</sup>	16.7±3.9 <sup>b</sup>	8.5±1.1 <sup>a</sup>	12.8±0.1 <sup>b</sup>	<0.001
18:0	7.8±0.6 <sup>*</sup>	7.6±0.4 <sup>*</sup>	7.3±0.4	5.9±0.3	5.7±0.4	6.5±0.9	0.252
∑Saturated	33.2±2.3	35.1±1.2 <sup>*</sup>	35.9±1.5 <sup>*</sup>	30.3±4.4 <sup>b</sup>	19.8±1.0 <sup>a</sup>	24.1±1.3 <sup>a</sup>	0.002
16:1n-9	1.2±0.5	0.8±0.4 <sup>*</sup>	0.6±0.2	1.7±0.5 <sup>ab</sup>	2.2±0.2 <sup>b</sup>	1.3±0.3 <sup>a</sup>	0.002
16:1n-7	1.6±0.5 <sup>ab</sup>	0.9±0.1 <sup>a*</sup>	2.0±0.1 <sup>b</sup>	2.9±1.7 <sup>b</sup>	0.2±0.1 <sup>a</sup>	0.0±0.0 <sup>a</sup>	<0.001
18:1n-9	12.1±4.8 <sup>a</sup>	25.3±2.1 <sup>b*</sup>	23.3±2.7 <sup>b*</sup>	12.3±2.9 <sup>b</sup>	6.2±1.3 <sup>a</sup>	9.8±1.1 <sup>b</sup>	<0.001
18:1n-7	1.4±0.2	1.4±0.1 <sup>*</sup>	1.4±0.1	2.1±0.6 <sup>c</sup>	0.7±0.1 <sup>a</sup>	1.0±0.1 <sup>b</sup>	<0.001
24:1n-9	35.8±5.2 <sup>b</sup>	14.3±4.9 <sup>a*</sup>	16.7±2.8 <sup>a*</sup>	30.6±6.3 <sup>a</sup>	56.9±1.2 <sup>b</sup>	50.3±4.7 <sup>b</sup>	<0.001
∑Monounsaturated	53.6±2.2	45.3±3.2 <sup>*</sup>	45.4±2.4 <sup>*</sup>	53.2±2.5	67.1±1.0	63.8±2.9	0.062
18:2n-6	0.8±0.0 <sup>a</sup>	8.0±1.1 <sup>b*</sup>	1.3±0.1 <sup>a</sup>	1.1±0.3	1.7±0.2	0.8±0.3	<0.001
∑n-6 PUFA	3.1±0.3 <sup>a*</sup>	13.7±2.0 <sup>b*</sup>	4.7±0.7 <sup>a*</sup>	5.3±1.9	3.6±0.2	2.9±0.7	<0.001
20:5n-3	0.7±0.0	0.3±0.1	1.1±0.2	1.7±1.3	1.0±0.8	1.0±0.4	0.319
22:6n-3	7.4±1.5 <sup>b</sup>	4.3±0.9 <sup>a</sup>	11.3±2.9 <sup>b*</sup>	7.3±1.5	6.7±1.1	7.0±0.2	0.01
∑n-3 PUFA	8.7±1.6 <sup>b</sup>	5.4±1.1 <sup>a*</sup>	13.1±3.1 <sup>b*</sup>	10.1±3.0	8.6±1.7	8.5±0.7	0.017
Total PUFA	13.2±1.8 <sup>a</sup>	19.6±2.1 <sup>b*</sup>	18.7±3.0 <sup>b*</sup>	16.5±4.2	13.1±1.9	12.0±1.6	0.009
∑n-3 LC-PUFA	8.2±1.4 <sup>ab</sup>	5.0±1.1 <sup>a</sup>	12.8±3.1 <sup>b*</sup>	9.6±2.9	8.0±1.9	8.3±0.6	0.018
n-3/n-6	2.9±0.8 <sup>b</sup>	0.4±0.1 <sup>a*</sup>	2.8±0.8 <sup>b</sup>	2.0±0.7	2.4±0.4	3.0±0.4	0.005
PI <sub>n</sub>	74.8±4.2 <sup>a</sup>	61.6±11.8 <sup>a</sup>	110.3±29.1 <sup>b*</sup>	97.1±1.4 <sup>b</sup>	81.5±0.8 <sup>a</sup>	78.2±8.1 <sup>a</sup>	0.021

Data expressed as mean ± SD (n=4). Different superscript letters within a row and for each age group represent significant differences between diet treatments as determined by a two-way ANOVA ( $p < 0.05$ ). Asterisks denote statistical differences between 1- and 3-year-old trout for each diet group (SHFO, RO and DHA) when compared using a Bonferroni test ( $p < 0.05$ ). Right column represent signification values for the interaction between Diet and Age ( $p < 0.05$ ). SHFO, fish oil diet group; RO, rapeseed oil group; DHA, Incromege TG0525 oil group; LC-PUFA, long-chain polyunsaturated fatty acids; PI<sub>n</sub>, peroxidation index; PUFA, polyunsaturated fatty acids.

Fatty acids representing less than 1% of total fatty acids are not shown.