

1 **The time course of fish oil wash-out follows a simple dilution model in gilthead sea**
2 **breem (*Sparus aurata* L.) fed graded levels of vegetable oils**

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17 **Abstract**

18

19 The aim of the study was to undertake whether changes in the flesh fatty acid (FA)
20 profiles follow a simple test dilution model after changing of dietary oil sources in
21 gilthead sea bream (*Sparus aurata* L.). A 14-month trial was conducted with juvenile
22 fish of 18 g initial body weight fed either a fish oil-based diet (FO diet) or vegetable oils
23 replacing the 33% (33VO) and 66% (66VO) of fish oil. The trial included 3 finishing
24 months with fish oil to follow the restoration of the FA profile with the FO diet. Fish oil
25 replacement with/without a finishing phase of fish oil re-feeding did not have any
26 detrimental effect on growth, and all groups reached 520-531 g body weight. Changes
27 in proximate body composition with weight gain did not modify the FA profile of fish
28 continuously fed FO, 33VO or 66VO diets. Increased amounts of oleic acid (18:1n-9),
29 linoleic acid (18:2n-6) and linolenic acid (18:3n-3), in combination with reduced
30 proportions of n-3 long chain polyunsaturated FAs, were found with the partial
31 replacement of fish oil. Hence, multivariate component analysis highlighted a gradient
32 of fish oil load determined by the total intake of fish oil over the entire production cycle.
33 The simple dilution model was a good descriptor of these flesh FA changes, and
34 excellent correlations between observed and predicted values were found at the end of
35 finishing period in fish grown out with either 33VO or 66VO diets.

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37 **Key words:** fish, growth, flesh, fatty acids, plant proteins

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39 **1. Introduction**

40

41 Fish oil supplies are finite (FAO, 2006) and the continuous increase in the global
42 aquaculture production has necessitated research on alternative lipid sources for fish
43 feeds (Watanabe, 2002). Since fish oils are also highly susceptible to contamination
44 with persistent organic pollutants, the use of vegetable oils can contribute towards the
45 reduction of contaminant loadings in the flesh of farmed fish (Sargent et al., 1995; Bell
46 et al., 2005). However, vegetable oils are devoid of n-3 long chain polyunsaturated fatty
47 acids (LC-PUFA), and can adversely affect the flesh fatty acid (FA) composition if
48 added at high inclusion levels (Sargent and Tacon, 1999; Torstensen et al., 2005). Thus,
49 it may be desirable the use of finishing diets formulated with
50 uncontaminated/decontaminated fish oils to restore the wild flesh FA profile of farmed
51 fish. For instance, southern hemisphere fish oils are cleaner than the northern
52 hemisphere fish oils and can deliver similar levels of n-3 LC-PUFA at lower dietary
53 inclusion levels (Pratoomyot et al., 2008).

54 Lipid tailoring is, however, a fish-specific process and marine fish show
55 extremely low capabilities for the bioconversion of C₁₈ polyunsaturated FAs into C₂₀
56 and C₂₂ PUFA. In this regard, we have found that the essential FA requirements of fast
57 growing juvenile gilthead sea bream are met in practical diets by less than 25% of fish
58 meal and fish oil inclusion (Benedito-Palos et al., 2007). Besides, n-3 LC-PUFA, in
59 particular eicosapentaenoic acid (EPA, 20:5n-3) and docosahexaenoic acid (DHA,
60 20:6n-3), are selectively incorporated into the polar lipid fraction resulting in a high
61 phospholipid robustness in FA composition (Benedito-Palos et al., 2008). The same
62 study points to that the muscle fat depots highly reflect the composition of the diet with
63 a minor influence of the natural seasonal, thermal or photoperiodical changes. Earlier

64 studies in gilthead sea bream also monitored the effect of fish oil re-feeding on the
65 tissue FA profile (Izquierdo et al., 2005). In turbot and brown trout, a simple dilution
66 model was proposed and validated by Robin et al. (2003) to follow the time-course of
67 FA changes after shift levels in dietary lipid sources. The same model was re-evaluated
68 with different success in Atlantic salmon (Jobling, 2004a; Jobling, 2004b), red sea
69 bream (Jobling, 2004a) and Murray cod (Turchini et al., 2006). The rationale for the
70 present study was to investigate whether dietary FAs are incorporated in the flesh of
71 gilthead sea bream following similar patterns. Specifically, we monitored FA dynamics
72 after fish oil re-feeding in fish previously fed plant protein-based diets at two levels of
73 fish oil replacement. Economy of fish oil usage was analysed for the entire 14-month
74 production cycle, including both grow-out and finishing periods.

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77 **2. Materials and methods**

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79 *2.1. Diets*

80

81 Three diets (2/5 mm pellets) with the same basal composition were coated with
82 fish and vegetable oils to contain 53% crude protein and 21% crude lipid on a dry
83 weight basis (Table 1). Fish oil of southern hemisphere was the only lipid source in the
84 reference/finishing diet (FO). The other two diets contained a blend of vegetable oils
85 (rapeseed oil: linseed oil: palm oil), replacing the 33% (33VO) and 66% (66VO) of fish
86 oil. The fatty acid composition of diets is reported in Table 2; reduction in fish oil
87 levels decreased the proportion of n-3 LC-PUFA (predominantly EPA and DHA) from
88 19.4% in the FO diet, to 6.6% in the 66VO diet. All diets were manufactured using a

89 twin-screw extruder (Clextral, BC 45) at the INRA experimental research station of
90 Donzacq (Landes, France), dried under hot air, sealed and kept in air-tight bags until
91 use.

92

93 2.2. *Grow-out trial*

94

95 Juvenile gilthead sea bream (*Sparus aurata* L.) of Atlantic origin (Ferme Marine
96 de Douhet, Ile d'Oléron, France) were acclimated to laboratory conditions at the
97 Institute of Aquaculture Torre de la Sal (IATS) for 20 days before the start of the study.
98 Fish of 18 g initial mean body weight were distributed into 9 fibreglass tanks (3000
99 litres) in groups of 150 fish per tank. Water flow was 20 l/min, and oxygen content of
100 outlet water remained higher than 5 ppm. The growth study was undertaken over 11
101 months (July 11th 2006 – June 6th 2007), and day-length and water temperature (10-
102 26°C) varied over the course of the trial following natural changes (40° 5'N; 0° 10'E).

103 Each diet was randomly allocated to triplicate groups of fish. Feed was offered
104 by hand to apparent visual satiety twice a day (0900 and 1400 hours; 6-7days per week)
105 from July 2006 to September 2006, and once a day (1200 hours, 4-6 days per week)
106 from October 2006 to June 2007. At regular intervals, fish were counted and group-
107 weighed under moderate anaesthesia (3-aminobenzoic acid ethyl ester, MS 222; 100
108 µg/ml) (Sigma-Aldrich, Madrid, Spain). Feed distributed and mortalities (< 5% during
109 the course of the whole 14-month period) were registered daily.

110

111 *2.3. Fish oil finishing trial and sampling protocol*

112

113 To follow the restoration of marine FA profile in fish fed vegetable oils, two of
114 the three replicates of fish fed 33VO and 66VO diets were fed once a day (6 days per
115 week) the FO diet from June 2007 to September 2007 (12 weeks). These duplicate
116 groups became 33VO/FO and 66VO/FO, respectively. The remaining fish were
117 continued to be fed with the FO (3 tanks), 33VO (1 tank) and 66VO (1 tank) diets.

118 At regular intervals after the start of the finishing phase (zero time, 27, 55 and 88
119 days), 8 randomly selected fish per dietary treatment were killed by a blow on the head.
120 The right-hand side whole fillets (denuded from skin and bone) were excised, vacuum
121 packed and stored at -80 °C until analyses. All procedures were carried out according to
122 national and institutional regulations (Consejo Superior de Investigaciones Científicas,
123 IATS Review Board) and the current European Union legislation on handling
124 experimental animals.

125

126 *2.4. Proximate analyses*

127

128 The proximate composition of diets and fillets were analysed by standard
129 procedures (AOAC, 1990). Moisture content was determined by drying in an oven at
130 105° C for 24 h. Diets and freeze-dried fillets were then blended and used for
131 determinations of lipid, protein and ash contents. Lipid content (0.5 g samples) was
132 determined by the Soxhlet method with extraction in diethyl ether at 120 °C (Soxhlet
133 4001046 Auto extraction apparatus; Selecta, Barcelona, Spain). Crude protein content
134 (N x 6.25) was determined using the automated Kjeldhal method (Kjeldhal Auto

135 4002430 Analyser, Selecta, Barcelona, Spain). Ash contents were determined after
136 heating at 600 °C in a muffle furnace for 2 h.

137

138 2.5. FA analyses

139

140 Total lipids for FA analyses were extracted by the method of Folch et al. (1957),
141 using chloroform:methanol (2:1) containing 0.01% butylated hydroxytoluene (BHT) as
142 antioxidant. After the addition of nonadecanoic FA (19:0) as internal standard, total
143 lipids (TL) were subjected to acid-catalysed transmethylation for 16 hours at 50 °C
144 using 1 ml toluene and 2 ml of 1% (v/v) sulphuric acid in methanol (Christie, 1982). FA
145 methyl esters (FAME) were extracted with hexane:diethyl ether (1:1), and purified by
146 thin layer chromatography (Silica gel G 60, 20 x 20 cm glass plates, Merck, Darmstadt,
147 Germany) using hexane:diethyl-ether:acetic acid (85:15:1.5) as a solvent system. FAME
148 were then analyzed with a Fisons Instruments GC 8000 Series (Rodano, Italy) gas
149 chromatograph, equipped with a fused silica 30 m x 0.25 mm open tubular column
150 (Tracer, TR-WAX; film thickness: 0.25 µm, Teknokroma, Spain) and a cold on-column
151 injection system. Helium was used as a carrier gas, and temperature programming was
152 from 50 to 180 °C at 40 °C/min and then to 220 °C at 3 °C/min. Peaks were recorded in a
153 personal computer using the Azur software package (version 4.0.2.0. Datalys, France).
154 Individual FAME were identified by reference to well characterized fish oil standards,
155 and the relative amount of each FA was expressed as a percentage of the total amount of
156 FA in the analysed sample.

157 BHT and internal standard (19:0) were obtained from Sigma-Aldrich (Madrid,
158 Spain). All solvents in lipid extraction and FA analyses were HPLC grade and were
159 obtained from Merck (Darmstadt, Germany).

160 2.6. Dilution model

161

162 Changes in the flesh FA profile as a result of fish oil re-feeding were described
163 according to Robin et al. (2003) by the following equation:

164
$$P_T = P_{RT} + [(P_0 - P_{RT}) / (Q_T / Q_0)]$$

165 where P_T is the percentage at time T of a given FA, P_0 is the FA percentage at the start
166 of the finishing period, and P_{RT} is the FA percentage at time T in fish continuously fed
167 the reference/finishing diet. Q_0 and Q_T represent the initial and final (at time T) flesh
168 lipid content, respectively.

169 In the present study, P_T is the predicted FA percentage at a given time T in
170 finishing groups (33VO/FO, 66VO/FO), P_0 is the FA percentage of a given FA at the
171 start of the finishing period in 33VO and 66VO groups, P_{RT} represents at time T that of
172 the reference group always fed the finishing FO diet. Q_0 and Q_T are the initial and final
173 flesh lipid content in the respective group. The adequacy of the dilution model was
174 evaluated by direct comparisons of model predictions with the observed values.

175

176 2.8. Statistical analysis

177

178 Growth parameters (fish average values per tank) and the relative amount of FA
179 were checked for normal distribution and homogeneity of variances, and when
180 necessary arcsin transformation was performed. Data were analysed by one-way
181 ANOVA followed by Student-Newman-Keuls (SNK) test at a significance level of 5%.
182 The percentages of each FA were chemometrically analysed by multivariate principal
183 components analysis (MPCA). All analyses were made using the SPSS package version
184 14.0 (SPSS Inc, Chicago, USA).

185 3. Results

186

187 3.1. Growth performance

188

189 Growth rates, feed intake and feed conversion were not affected by the dietary
190 treatment over the course of the study. Hence, at each sampling point, all data on body
191 weight and feed intake were put together and represented in the fitting plot as the mean
192 of the 9 experimental tanks (Fig. 1). Overall, fish grew during the 11-month grow-out
193 period from 18 g to 284 - 294 g with a feed efficiency (wet wt gain/dry feed intake) of
194 0.82 - 0.86 over this whole period. The subsequent trial (3-month period) was
195 conducted over the course of summer, and the cumulative feed intake (g/fish) was of the
196 same order of magnitude than that of the initial period (324 vs 307 g). At the end of the
197 wash-out trial, mean body weight of fish among tanks varied between 520 and 531 g
198 with a feed efficiency for the finishing period of 0.73 – 0.79.

199 In our experimental model, the usage of the added fish oil in the diet (g/fish) in
200 fish always fed FO, 33VO and 66VO diets (growth- plus wash-out periods) was
201 proportional to the percentage of replacement (Fig. 2). At the end of the fish oil wash-
202 out period, fish oil usage in fish fed 33VO became equal to that found in the 66VO/FO
203 group. In the 33VO/FO group, fish oil usage was reduced by a 15% in comparison to
204 fish always fed the FO diet.

205

206 3.2. *Lipid content and flesh FA profile*

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208 Fillet yield and lipid content of skinned fillets were not significantly affected by
209 the dietary treatment (Table 3). However, lipid deposition increased by a 20–30 % over
210 the course of the finishing period regardless of the dietary treatment (Tables 3-5).

211 As shown in Table 3, no consistent changes on the FA profile were found over
212 the course of the finishing period in fish continuously fed with the same diets (FO,
213 33VO and 66VO groups), and only few FAs (16:1 n-7, 17:0, 18:2 n-6, 21:5) showed
214 significant differences (less than 5-30% of variation) in one or two of the three
215 experimental groups. Regarding the effect of dietary treatment, fish fed the FO diet
216 contained 29% saturates (mainly 16:0 and 14:0), almost 32% monoenes (over half of
217 which were 18:1n-9), 1% n-6 LC-PUFA, and 17% n-3 LC-PUFA (predominantly EPA
218 and DHA). Increased amounts of 18:1n-9, 18:2n-6 and 18:3n-3, in combination with
219 reduced proportions of n-3 LC-PUFA and saturated FAs, were found with the
220 progressive replacement of fish oil by vegetable oils.

221 The time course of changes through the finishing period on the flesh FA profile
222 of fish previously fed vegetable oils are shown in Tables 4 and 5. Both in 33VO/FO and
223 66VO/FO groups, the finishing diet caused a progressive increase in the FAs present in
224 higher amounts in fish oil (i.e. 14:0, 16:1n-7, 20:1n-9, 22:1n-11, EPA and DHA), while
225 those characteristic of vegetable oils (i.e. 18:1n-9, 18:2n-6 and 18:3n-3) decreased in
226 proportion to degree of fish oil replacement in the diet.

227 The MPCA analysis of fillet FA profiles before, during and at the end of the
228 finishing period revealed that the two first components accounted for 62% of the total
229 variation, with 52.5% of the variation being explained by component 1 itself (Fig. 3A).
230 Some of the most characteristic variables of marine versus vegetable oils had the

231 highest loadings on function 1 and were located at the extremes. The results of the score
232 plot are represented only for the first component since it accounted for the majority of
233 the variation (Fig. 3B). The plot revealed the three invariable groups (FO, 33VO and
234 66VO) well separated from each other, with 66VO and FO at the extremes. The
235 finishing 33VO/FO and 66VO/FO groups were also clearly separated from each other
236 on a time- FO intake-manner (Fig. 3B). Thus, a gradient of fish oil load caused either by
237 the amount of this ingredient in the diet, or by the total intake per unit of body weight,
238 could be easily distinguishable. At the end of the finishing period, the resulting FA
239 profile of the 66VO/FO became equal to that of fish always fed the 33VO diet, and
240 intermediate values between 33VO and FO groups were found for the 33VO/FO group.

241 Regardless of nutritional background (33VO and 66VO diets), the concordance
242 between the observed FA values (x-axis) and those predicted by the dilution model (y-
243 axis) was extremely high at the end of the finishing period (Fig. 4). This gave a
244 regression line with a slope very close to 1 (0.96-0.95) when 32 FAs were considered in
245 the models derived from both 33VO/FO and 66VO/FO fish. Similar slopes (1.04-1.05)
246 were obtained when calculations were repeated for 14 selected FAs having a high
247 weight in the MPCA.

248

249 **4. Discussion**

250

251 Data reported here, along with those of Benedito-Palos et al. (2008) over an 8-
252 month trial, convincingly demonstrate that dietary fish oils of northern and southern
253 origin can be replaced up to 66% without negative effects on the growth performance of
254 gilthead sea bream. These original data are pioneer at in a stenohaline marine teleost
255 maximizing the simultaneous replacement of fish meal and fish oil by alternative plant

256 ingredients without any histological sign of damage in the liver and intestine epithelium
257 (Benedito-Palos et al., 2008). Other metabolic effects of feeding diets with alternative
258 oils are complex, interconnected and, to date, not fully understood in gilthead sea bream
259 and other fish species. However, it must be noted that growth-compensatory
260 mechanisms are orchestrated at the local tissue level (skeletal muscle) by the
261 somatotrophic axis when fast growing juvenile fish of gilthead sea bream are fed with
262 increased levels of vegetable oils (Benedito-Palos et al., 2007). Besides, n-3 LC-PUFA
263 are selectively incorporated into polar lipids, and the stability of muscle phospholipid
264 FA composition is a useful criterion to assess the suitability of the replacement strategy
265 in fish feeds with low levels of marine derived ingredients (Benedito-Palos et al., 2008).

266 In fish and higher vertebrate species, neutral lipids are less conservative than
267 phospholipids (Tocher, 2003; Skalli and Robin, 2004; Schulz et al., 2005). This is
268 because they are the fat storage form and its FA profile highly reflects that of the diet. In
269 the present study, the muscle lipid content was greater than 10% on wet matter basis,
270 and the FA profile of total lipids and thereby that of triglycerides (TAG) remained
271 mostly unchanged (finishing phase) in fish always fed either FO, 33VO or 66VO diets
272 (Table 3). The result of these temporal series agrees with data on a previous seasonal
273 study (Benedito-Palos et al., 2008), and reinforces the idea that accelerated growth
274 overrides major changes in the FA profile in spite of concurrent changes in body
275 composition through weight gain (Grigorakis, 2007). However, it should be born in
276 mind that the tissue-specific FA profile varies in salmonids with the size and age of fish
277 (Bell et al., 2002; Bell et al., 2003a). Probably, this is also true for gilthead sea bream
278 and FA databases considering the effect of different size class, gender, season,
279 maturation state, and nutritional condition are now under construction. Anyway, fish oil
280 replacement by alternative lipid sources has a pronounced effect on the flesh FA profile

281 of most fish. In the present study, we found in fish always fed 33VO and 66VO diets a
282 22-36 % increase of 18:1 n-9 and 18:2 n-6 with a concurrent 20-65 % reduction in EPA
283 and DHA. Similar results have been reported in gilthead sea bream (Izquierdo et al.,
284 2005) and a wide variety of fish species, including Atlantic salmon (Bell et al., 2002;
285 Bransden et al., 2003; Bell et al., 2003b; Torstensen et al., 2004; Bell, 2004; Nanton et
286 al., 2007), rainbow trout (Drew et al., 2007), turbot (Regost et al., 2003), European sea
287 bass (Montero et al., 2005; Mourente and Bell, 2006), Murray cod (Francis et al.,
288 2007a; 2007b), red sea bream (Piedecausa et al., 2007; Huang et al., 2007) and black sea
289 bream (Peng et al., 2008). Since this feature can compromise the beneficial effects of
290 sea food (Din et al., 2004; Psota et al., 2006) as main source of EPA and DHA in the
291 human diet, there is now increased interest in finfish aquaculture for modelling the
292 time-course of FA changes during fish oil re-feeding.

293 Gilthead sea bream shows in our latitude a pronounced growth seasonality
294 (Mingarro et al., 2002), and the window for fish oil wash-out should take place in the
295 broadly active feeding period of May-October. Thus, after the growth stop of cold
296 season, one month (May) was spent before to start the finishing trial that then continued
297 through the summer growth spurt (June-September). Several variables, including among
298 others the growth and lipid deposition rates, need to be considered to analyse the
299 effectiveness of fish oil wash-out. Therefore, one must be cautious before drawing a
300 definitive conclusion, but the literature is prolific on studies in which a complete
301 restoration of the FA profile is not fully achieved after fish oil re-feeding: 32 days red
302 sea bream (Glencross et al., 2003), 8 weeks turbot (Regost et al., 2003); 8 weeks brown
303 trout (Robin et al., 2003), 12-25 weeks Atlantic salmon (Bell et al., 2003a; Bell et al.,
304 2003b; Torstensen et al., 2004; Bell, 2004); 3 months gilthead sea bream (Izquierdo et
305 al., 2005); 4 months Murray cod (Turchini et al., 2006); 5 months European sea bass

306 (Montero et al., 2005). Most evidences point towards a dilution model, which was
307 proved in turbot, brown trout and Murray cod. Using original data and those derived
308 from red sea bream (Glencross et al., 2003) and Atlantic salmon (Bell et al., 2003a)
309 studies, Jobling (2004a; 2004b) concluded that a dilution process also plays a key role
310 in governing the muscle FA profile of these fish species. Our work in gilthead sea
311 bream points clearly towards the same direction, and gradual changes in the FA profiles
312 of 33VO/FO and 66VO/FO groups were found during the finishing period, making
313 them increasingly similar to the FO group. This is particularly highlighted by the results
314 of the MPCA (Fig. 3) that shows the gradient of fish oil load along the ordinate axis.

315 Therefore, changes in the FA profile arise because the existing stores become
316 diluted as fish grow and deposit increasing amounts of dietary-derived FAs. In other
317 words, nutritional background in fish with no apparent signs of FA deficiencies has a
318 marginal role on FA turnover and flesh FA profiles, although age- and nutritional
319 condition affect the expression pattern of cytokines and key limiting enzymes on tissue
320 FA uptake and mobilization (Saera-Vila et al., 2005; 2007). Thus, using the simple
321 dilution model, a reliable FA prediction was found herein at the end of the fish oil
322 finishing period regardless of the level of fish oil replacement (Fig. 4). The model is in
323 fact a good general descriptor of FAs, and regression curves (predicted *vs* observed
324 values) give slopes nearby to the line of equality when either selected or almost all FAs
325 were considered in the model. In Atlantic salmon, Jobling (2004b) tested three FAs
326 (18:1 isomers, 18:2 n-6 and 18:3 n-3) and confirmed closely the predictions made with
327 the dilution model. Jobling (2004a) again evaluated the dilution model with data from
328 red sea bream studies (Glencross et al., 2003), and a high degree of concordance was
329 found between the predicted and observed values. Turchini et al. (2006) also indicated
330 that deviations for some individual FAs in Murray cod does not invalidate totally the

331 FA dilution model, although FA oxidation and biosynthetic capacities vary from one
332 fish species to another. In spite of these limitations, the dilution model is a useful tool
333 for predicting the FA composition of aquaculture growing fish. This is especially true
334 for gilthead sea bream, and MPCA highlights that fish grow-out with the 33VO diet
335 need more than 12 weeks to revert back the FA composition towards the normal
336 variability of fish fed fish oil-based diets. Besides, low and intermediate levels of fish
337 oil replacement can produce equally acceptable fillets when the latter is accompanied by
338 a fish oil finishing phase. This is because temporal changes on fish oil intake gave a
339 minor effect on the muscle FA profile if the absolute amount becomes equal at the end
340 of the trial. In our experimental model, this was the case of 33VO and 66VO/FO groups
341 (Fig. 2), and the flesh FA profile at the end of the 3-month finishing period was very
342 close in both groups (Fig. 3B).

343 Savings on fish oil resources are therefore limited to a simple dilution, and new
344 approaches are required to improve any mobilisation or turnover of pre-existing FAs.
345 Intake of conjugate linoleic acid complex (CLA) has a lipid-lowering effect in gilthead
346 sea bream juveniles, promotes the diversion of dietary-derived TAG from muscle and
347 adipose tissue to liver, and increase hepatic peroxisomal β oxidation (Diez et al., 2007).
348 At the same time, however, the LC-PUFA biosynthesis is reduced and changes in the
349 muscle FA profile indicate that the inclusion of CLA in aquaculture diets would be of
350 little benefit in gilthead sea bream. Nevertheless, the use of more specific
351 agonists/antagonists of peroxisome proliferators-activated receptors (PPARs) cannot be
352 excluded to improve the retention of n-3 LC-PUFA. Attention also needs to be focused
353 on the transfer from fish oil of PCBs, dioxins and other harmful lipophilic organic
354 chemicals that are now ubiquitous contaminants of marine ecosystems (Sargent et al.,
355 1995; Jacobs et al., 2002; Bell et al., 2005; Domingo, 2007). The effects of feeding

356 strategies on toxic-kinetics will be reported separately to have a more complete
357 framework of nutritional fish tailoring, and to gain public acceptance for the now
358 becoming fish fed with alternative and sustainable diets.
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362

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517 **Legends**

518

519 Figure 1. (A) Seasonal changes on temperature (solid line) and day length (dashed line).

520 White and black boxes at the top of figure refer to summer and winter period

521 respectively. (B) Body weight representation as mean \pm SEM of all experimental groups

522 in the experimental design. Arrows indicate the start of the study and finishing periods.

523 Cumulative feed intake is indicated at the top of figure for each period.

524

525 Figure 2. Effects of diet composition and feeding protocol on fish oil (added fish oil in

526 the diet) intake through grow-out (330 days, black bars) and finishing (88 days, grey

527 bars) periods.

528

529 Figure 3. Component plot (A) and factor score plot (B) of the MPCA for the fillet FA

530 profile through the 3-month finishing period (June-September). Temporal series (July,

531 August and September) derived from 33VO/FO and 66VO/FO fish are represented in

532 factor score 1 as mean \pm SEM (n = 8). No temporal changes were found for FO, 33VO

533 and 66VO groups and data from June and September (initial and final steps of finishing

534 period) are represented as one point for each group.

535

536 Figure 4. Plot prediction (dilution model) in 33VO/FO (A) and 66VO/FO (B) groups of

537 the flesh FA profile at the end of the finishing period. Observed values are the mean \pm

538 SEM of 8 fish per treatment. The solid line is the plotted regression. The equations were

539 calculated considering both the selected and all (square brackets) the identified FAs.

540

541 **Table 1.** Ingredients and chemical composition of experimental diets.

Ingredient (%)	FO	33VO	66VO
Fish meal (CP 70%) ¹	15	15	15
CPSP 90 ²	5	5	5
Corn gluten	40	40	40
Soybean meal	14.3	14.3	14.3
Extruded wheat	4	4	4
Fish oil ³	15.15	10.15	5.15
Rapeseed oil	0	0.85	1.7
Linseed oil	0	2.9	5.8
Palm oil	0	1.25	2.5
Soya lecithin	1	1	1
Binder	1	1	1
Mineral premix ⁴	1	1	1
Vitamin premix ⁵	1	1	1
CaHPO ₄ ·2H ₂ O (18%P)	2	2	2
L-Lys	0.55	0.55	0.55
<i>Proximate composition</i>			
Dry matter (DM, %)	93.13	92.9	92.77
Protein (% DM)	53.2	52.81	52.62
Fat (% DM)	21.09	21	20.99
Ash (% DM)	6.52	6.69	6.57

542

543 ¹Fish meal (Scandinavian LT)544 ²Fish soluble protein concentrate (Sopropêche, France)545 ³Fish oil (Sopropêche, France)

546 ⁴Supplied the following (mg / kg diet, except as noted): calcium
547 carbonate (40% Ca) 2.15 g, magnesium hydroxide (60% Mg)
548 1.24 g, potassium chloride 0.9 g, ferric citrate 0.2 g, potassium
549 iodine 4 mg, sodium chloride 0.4 g, calcium hydrogen phosphate
550 50 g, copper sulphate 0.3, zinc sulphate 40, cobalt sulphate 2,
551 manganese sulphate 30, sodium selenite 0.3.

552 ⁵Supplied the following (mg / kg diet): retinyl acetate 2.58, DL-
553 cholecalciferol 0.037, DL- α tocopheryl acetate 30, menadione
554 sodium bisulphite 2.5, thiamin 7.5, riboflavin 15, pyridoxine 7.5,
555 nicotinic acid 87.5, folic acid 2.5, calcium pantothenate 2.5,
556 vitamin B₁₂ 0.025, ascorbic acid 250, inositol 500, biotin 1.25
557 and choline chloride 500.

558

559 **Table 2.** FA composition of experimental diets (% of total FAME).
 560

FA %	FO	33VO	66VO
14:0	7.18	5	2.7
15:0	0.12	0.09	tr
16:0	22.26	20.30	18.48
16:1n-7	7.06	4.85	2.62
16:2	0.47	0.31	0.15
16:3	1.66	1.09	0.46
16:3n-3	0.11	0.07	0.03
16:4	1.8	1.1	0.47
17:0	0.96	0.64	0.32
18:0	4.27	3.92	3.55
18:1n-9	12.49	20.39	24.59
18:1n-7	2.97	0.23	tr
18:2n-6	10.35	14.03	17.48
18:3 n-6	0.34	0.21	0.09
18:3n-3	0.81	9.16	17.33
18:4n-3	1.8	1.17	0.62
20:0	0.07	0.07	0.06
20:1n-9	0.92	0.97	1.03
20:2n-6	0.6	0.16	0.19
20:3 n-6	0.07	0.11	tr
20:4n-6	0.69	0.43	0.18
20:4n-3	0.3	0.21	0.15
20:5n-3	13.57	8.84	4.38
21:5	0.4	0.22	0.06
22:1n-11	0.97	0.82	0.76
22:5n-3	0.81	0.56	0.23
22:6n-3	4.78	3.3	1.88
Total	97.83	98.25	97.81
Saturates	34.86	30.02	25.11
Monoenes	24.41	27.26	29
n-3 LC-PUFA ¹	19.46	12.91	6.64
n-6 LC-PUFA ²	1.36	0.7	0.37

561 tr = trace values

562 ¹Calculated excluding 16 C and 18 C.

563 ²Calculated excluding 18 C.

564 Table 3. Fillet weight, wet lipid content and FA profile (% of total FAME) in fish always fed
 565 FO, 33VO and 66VO diets. Fish were sampled at the beginning (June 2007) and at the end of
 566 the finishing period (September 2007). Mean values and standard deviations are presented (n =
 567 8). Mean values within dietary groups with unlike superscript letters are significantly different
 568 (P<0.05).

	FO				33VO				66VO			
	June		September		June		September		June		September	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Fillet (g)	61.50 ^a	12.5	114.7 ^b	14.4	59.52 ^a	5.73	113.1 ^b	20.5	58.23 ^a	13.2	124.2 ^b	14.9
Lipids (%)	9.19 ^a	2.01	11.36 ^b	1.21	10.97 ^a	3.20	12.84 ^b	3.24	9.33 ^a	2.02	11.40 ^b	0.88
Σ FAs (mg/g)	53.99	6.61	61.67	10.8	62.46	18.8	66.78	16.5	59.76	14.2	60.54	7.86
FA (%)												
14:0	5.34	0.47	5.72	0.51	4.24	0.52	4.21	0.43	2.78	0.32	2.92	0.42
15:0	0.20	0.09	0.17	0.05	0.27	0.14	0.19	0.08	0.27	0.22	0.21	0.18
16:0	17.78	1.37	19.34	0.51	17.62	0.70	18.65	0.99	16.58	0.36	18.00	0.68
16:1n-7	7.96	0.64	8.36	0.25	5.96	0.47	5.94	0.18	3.85 ^a	0.14	4.08 ^b	0.17
16:2	1.03	0.20	0.90	0.29	0.60	0.10	0.69	0.15	0.27	0.09	0.35	0.08
16:3	1.21	0.21	1.35	0.06	0.82	0.08	0.91	0.17	0.43	0.23	0.55	0.14
16:4	1.09	0.30	0.95	0.08	0.52	0.06	0.53	0.07	0.24	0.02	0.26	0.02
17:0	0.66 ^a	0.31	0.45 ^b	0.37	0.24	0.02	0.25	0.01	0.25	0.09	0.25	0.14
18:0	3.70	0.38	3.71	0.30	3.76	0.17	3.90	0.19	3.85	0.12	3.84	0.12
18:1 n-9	18.13	2.24	18.11	0.77	23.21	1.05	23.27	0.59	27.06	0.71	26.97	0.59
18:1 n-7	3.23	0.25	3.30	0.14	2.56	0.12	2.52	0.10	2.00	0.12	1.98	0.19
18:2 n-6	9.39 ^a	0.83	8.95 ^b	0.31	12.32 ^a	0.75	11.96 ^b	0.55	15.02	0.31	14.34	0.31
18:3 n-6	0.38	0.06	0.39	0.02	0.25	0.02	0.25	0.01	0.14	0.01	0.15	0.03
18:3 n-3	1.16	1.20	0.71	0.04	7.39	0.93	7.11	0.26	12.94	0.33	10.82	4.33
18:4 n-3	1.16	0.14	1.23	0.07	0.82	0.02	0.82	0.05	0.60	0.06	2.03	4.07
20:0	0.20	0.02	0.20	0.01	0.19	0.01	0.22	0.06	0.18	0.02	0.19	0.01
20:1 n-7	0.21	0.02	0.22	0.01	0.15	0.01	0.16	0.01	0.11	0.01	0.11	0.01
20:1 n-9	1.14	0.08	1.21	0.02	1.02	0.05	1.08	0.03	0.96	0.03	1.02	0.03
20:1 n-11	0.24	0.02	0.24	0.02	0.20	0.02	0.20	0.02	0.17	0.02	0.17	0.01
20:2 n-6	0.25	0.08	0.26	0.04	0.28	0.06	0.29	0.05	0.32	0.07	0.32	0.07
20:3 n-6	0.26	0.07	0.28	0.03	0.23	0.04	0.25	0.07	0.24	0.11	0.21	0.06
20:3 n-3	0.16	0.20	0.23	0.25	0.23	0.09	0.26	0.09	0.34	0.04	0.36	0.05
20:4 n-6	0.57	0.18	0.53	0.19	0.40	0.12	0.36	0.09	0.23	0.03	0.24	0.03
20:4 n-3	0.61	0.06	0.62	0.03	0.51	0.03	0.51	0.03	0.42	0.02	0.40	0.03
20:5 n-3	8.49	0.89	8.84	0.36	5.57	0.39	5.25	0.38	2.88	0.18	2.98	0.31
21:5	0.41 ^a	0.05	0.50 ^b	0.06	0.31	0.03	0.30	0.04	0.17	0.02	0.13	0.06
22:0	0.17	0.06	0.16	0.06	0.16	0.06	0.15	0.05	0.14	0.03	0.15	0.03
22:1 n-9	0.32	0.02	0.32	0.03	0.31	0.02	0.30	0.02	0.29	0.01	0.28	0.02
22:1 n-11	0.89	0.06	0.95	0.03	0.72	0.05	0.73	0.05	0.58	0.01	0.59	0.02
22:5 n-3	2.66	0.33	2.63	0.15	1.92	0.12	1.94	0.19	1.19	0.04	1.20	0.17
22:6 n-3	5.48	0.90	5.16	0.20	3.92	0.21	3.58	0.27	2.57	0.21	2.29	0.18
24:1 n-9	0.35	0.06	0.28	0.11	0.31	0.03	0.26	0.11	0.31	0.04	0.30	0.03
Saturates	28.04	1.92	29.75	0.74	26.47	1.64	27.26	1.29	24.04	0.67	25.56	0.90
Monoenes	32.53	2.58	33.22	0.84	34.45	1.81	34.51	0.64	35.18	0.84	35.32	0.73
n-3 LC-PUFA ¹	17.40	2.06	17.47	0.60	12.15	0.84	11.54	0.76	7.40	0.22	7.23	0.62
n-6 LC-PUFA ²	1.09	0.21	1.07	0.18	0.92	0.22	0.90	0.09	0.79	0.06	0.78	0.06

¹Calculated excluding 18 C. ²Calculated excluding 18 C.

569 Table 4. Effects of finishing diet on fillet weight, wet lipid content and FA profile (% of total
 570 FAME) in fish fed 33VO diet and then FO diet (33VO/FO group). Fish were sequentially
 571 sampled through the finishing period (June, +0; July, +27; August, +55; September, +88).
 572 Mean values and standard deviations are presented (n = 8). Raw values with unlike superscript
 573 letters are significantly different over sampling time (P<0.05).

	Jun (+0)		Jul (+27)		Aug (+55)		Sep (+88)	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Fillet (g)	59.52 ^a	5.73	74.58 ^b	14.8	98.66 ^c	8.59	126.1 ^d	12.5
Lipids (%)	10.97	3.20	11.44	1.82	13.19	1.73	12.36	1.62
Σ FAs (mg/g)	62.46	18.8	68.26	23.5	69.64	12.6	63.84	19.9
FA (%)								
14:0	4.24 ^a	0.52	4.39 ^{ab}	0.36	4.55 ^{ab}	0.35	4.98 ^b	0.32
15:0	0.27	0.14	0.17	0.08	0.17	0.07	0.15	0.08
16:0	17.62 ^a	0.70	18.29 ^b	0.40	18.77 ^b	0.52	19.30 ^b	0.36
16:1n-7	5.96 ^a	0.47	6.48 ^{ab}	0.20	6.97 ^b	0.32	7.15 ^b	0.12
16:2	0.60 ^a	0.10	0.78 ^b	0.14	0.74 ^b	0.25	0.72 ^b	0.22
16:3	0.82 ^a	0.08	0.94 ^{ab}	0.07	1.01 ^b	0.06	1.08 ^b	0.10
16:4	0.52 ^a	0.06	0.55 ^{ab}	0.24	0.62 ^{ab}	0.21	0.71 ^b	0.12
17:0	0.24	0.02	0.26	0.03	0.40	0.28	0.30	0.05
18:0	3.76	0.17	3.89	0.08	3.70	0.18	3.93	0.10
18:1 n-9	23.21 ^a	1.05	22.61 ^{ab}	0.98	21.84 ^{ab}	0.88	21.58 ^b	0.78
18:1 n-7	2.56 ^a	0.12	2.75 ^b	0.12	2.89 ^{bc}	0.14	2.94 ^c	0.12
18:2 n-6	12.32 ^a	0.75	11.61 ^b	0.60	10.77 ^c	0.29	10.16 ^d	0.17
18:3 n-6	0.25 ^a	0.02	0.29 ^b	0.01	0.31 ^c	0.01	0.33 ^d	0.01
18:3 n-3	7.39 ^a	0.93	5.78 ^{ab}	0.45	4.74 ^{bc}	0.12	3.61 ^c	0.13
18:4 n-3	0.82	0.02	0.76	0.30	0.97	0.07	0.90	0.34
20:0	0.19	0.01	0.19	0.01	0.18	0.01	0.20	0.02
20:1 n-7	0.15 ^a	0.01	0.17 ^{ab}	0.01	0.18 ^b	0.01	0.19 ^b	0.01
20:1 n-9	1.02 ^a	0.05	1.11 ^b	0.09	1.14 ^b	0.05	1.11 ^b	0.04
20:1 n-11	0.20	0.02	0.21	0.01	0.22	0.03	0.22	0.02
20:2 n-6	0.28	0.06	0.32	0.03	0.29	0.04	0.25	0.07
20:3 n-6	0.23	0.04	0.24	0.03	0.24	0.03	0.24	0.03
20:3 n-3	0.23 ^a	0.09	0.22 ^a	0.13	0.15 ^b	0.01	0.17 ^b	0.13
20:4 n-6	0.40	0.12	0.43	0.11	0.47	0.03	0.46	0.11
20:4 n-3	0.51	0.03	0.55	0.03	0.58	0.03	0.55	0.02
20:5 n-3	5.57 ^a	0.39	6.06 ^a	0.52	6.83 ^b	0.31	6.91 ^b	0.22
21:5	0.31	0.03	0.32	0.04	0.36	0.05	0.36	0.04
22:0	0.16	0.06	0.14	0.03	0.13	0.05	0.17	0.07
22:1 n-9	0.31	0.02	0.30	0.01	0.31	0.03	0.30	0.01
22:1 n-11	0.72 ^a	0.05	0.77 ^{ab}	0.03	0.80 ^b	0.05	0.80 ^b	0.02
22:5 n-3	1.92	0.12	2.10	0.09	2.20	0.20	2.16	0.11
22:6 n-3	3.92	0.21	4.08	0.41	4.08	0.28	4.19	0.21
24:1 n-9	0.31	0.03	0.33	0.03	0.32	0.04	0.30	0.02
Saturates	26.47 ^a	1.64	27.34 ^a	0.19	27.89 ^{ab}	0.88	29.03 ^b	0.46
Monoenes	34.45	1.81	34.72	1.11	34.71	0.79	34.67	0.86
n-3 LC-PUFA ¹	12.15 ^a	0.84	13.00 ^{ab}	0.83	13.85 ^{ab}	0.65	13.97 ^b	0.46
n-6 LC-PUFA ²	0.92	0.22	0.99	0.10	1.01	0.05	0.95	0.11

¹Calculated excluding 18 C. ²Calculated excluding 18 C.

574 Table 5. Effects of finishing diet on fillet weight, wet lipid content and FA profile (% of total
 575 FAME) in fish fed 66VO diet and then FO diet (66VO/FO group). Fish were sequentially
 576 sampled through the finishing period (June, +0; July, +27; August, +55; September, +88).
 577 Mean values and standard deviations are presented (n = 8). Raw values with unlike superscript
 578 letters are significantly different over sampling time (P<0.05).
 579

	Jun (+0)		Jul (+27)		Aug (+55)		Sep (+88)	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Fillet (g)	58.23 ^a	13.2	71.61 ^b	11.1	97.65 ^c	9.33	116.2 ^d	12.9
Lipids (%)	9.33	2.02	10.30	3.27	11.81	2.51	11.15	1.11
Σ FAs (mg/g)	59.76	14.2	56.27	20.6	67.46	17.6	55.14	7.23
FA (%)								
14:0	2.78 ^a	0.32	3.27 ^a	0.34	3.68 ^{ab}	0.27	4.38 ^b	0.39
15:0	0.27	0.22	0.10	0.06	0.18	0.09	0.16	0.09
16:0	16.58 ^a	0.36	17.63 ^b	0.62	18.14 ^b	0.54	18.81 ^c	0.61
16:1n-7	3.85 ^a	0.14	4.81 ^{ab}	0.31	5.60 ^{bc}	0.19	6.20 ^c	0.13
16:2	0.27 ^a	0.09	0.39 ^{ab}	0.11	0.55 ^b	0.20	0.59 ^b	0.19
16:3	0.43 ^a	0.23	0.63 ^a	0.09	0.70 ^{ab}	0.21	0.91 ^b	0.07
16:4	0.24 ^a	0.02	0.31 ^{ab}	0.15	0.50 ^{bc}	0.05	0.60 ^c	0.08
17:0	0.25	0.09	0.28	0.17	0.28	0.12	0.34	0.20
18:0	3.85	0.12	3.86	0.18	3.81	0.13	3.85	0.21
18:1 n-9	27.06 ^a	0.71	25.89 ^b	1.43	24.62 ^c	0.58	22.80 ^d	0.90
18:1 n-7	2.00 ^a	0.12	2.31 ^b	0.11	2.46 ^c	0.16	2.66 ^d	0.09
18:2 n-6	15.02 ^a	0.31	13.81 ^b	0.48	12.67 ^c	0.31	11.69 ^d	0.34
18:3 n-6	0.14 ^a	0.01	0.20 ^{ab}	0.02	0.23 ^{bc}	0.01	0.27 ^c	0.02
18:3 n-3	12.94 ^a	0.33	10.35 ^b	0.47	8.36 ^c	0.45	6.31 ^d	0.50
18:4 n-3	0.60 ^a	0.06	0.61 ^{ab}	0.23	0.83 ^b	0.03	0.92 ^c	0.07
20:0	0.18	0.02	0.19	0.02	0.18	0.00	0.20	0.02
20:1 n-7	0.11 ^a	0.01	0.13 ^{bc}	0.01	0.14 ^c	0.01	0.17 ^d	0.01
20:1 n-9	0.96 ^a	0.03	1.00 ^{ab}	0.06	1.05 ^{bc}	0.03	1.08 ^c	0.06
20:1 n-11	0.17	0.02	0.20	0.02	0.18	0.02	0.20	0.02
20:2 n-6	0.32	0.07	0.30	0.13	0.24	0.09	0.30	0.04
20:3 n-6	0.24	0.11	0.25	0.06	0.22	0.04	0.24	0.05
20:3 n-3	0.34 ^a	0.04	0.30 ^{ab}	0.03	0.26 ^{ab}	0.05	0.24 ^b	0.09
20:4 n-6	0.23 ^a	0.03	0.32 ^{ab}	0.10	0.33 ^{ab}	0.06	0.39 ^b	0.08
20:4 n-3	0.42 ^a	0.02	0.47 ^b	0.06	0.49 ^{bc}	0.02	0.52 ^c	0.02
20:5 n-3	2.88 ^a	0.18	3.90 ^b	0.61	5.01 ^c	0.22	5.86 ^d	0.31
21:5	0.17 ^a	0.02	0.21 ^{bc}	0.08	0.26 ^{bc}	0.03	0.29 ^c	0.07
22:0	0.14	0.03	0.17	0.06	0.13	0.04	0.18	0.04
22:1 n-9	0.29	0.01	0.30	0.02	0.29	0.01	0.30	0.02
22:1 n-11	0.58 ^a	0.01	0.65 ^b	0.03	0.69 ^c	0.03	0.76 ^d	0.03
22:5 n-3	1.19 ^a	0.04	1.40 ^b	0.09	1.61 ^c	0.06	1.87 ^d	0.11
22:6 n-3	2.57 ^a	0.21	3.14 ^b	0.38	3.28 ^b	0.26	3.65 ^c	0.42
24:1 n-9	0.31	0.04	0.33	0.03	0.32	0.04	0.32	0.03
Saturates	24.04 ^a	0.67	25.50 ^b	1.10	26.39 ^b	0.74	27.92 ^b	0.92
Monoenes	35.18	0.84	35.50	1.66	35.29	0.46	34.47	0.86
n-3 LC-PUFA ¹	7.40 ^a	0.22	9.21 ^b	0.65	10.65 ^c	0.47	12.13 ^d	0.75
n-6 LC-PUFA ²	0.79	0.06	0.86	0.18	0.80	0.09	0.93	0.09

¹Calculated excluding 18 C. ²Calculated excluding 18 C.

Figure 1

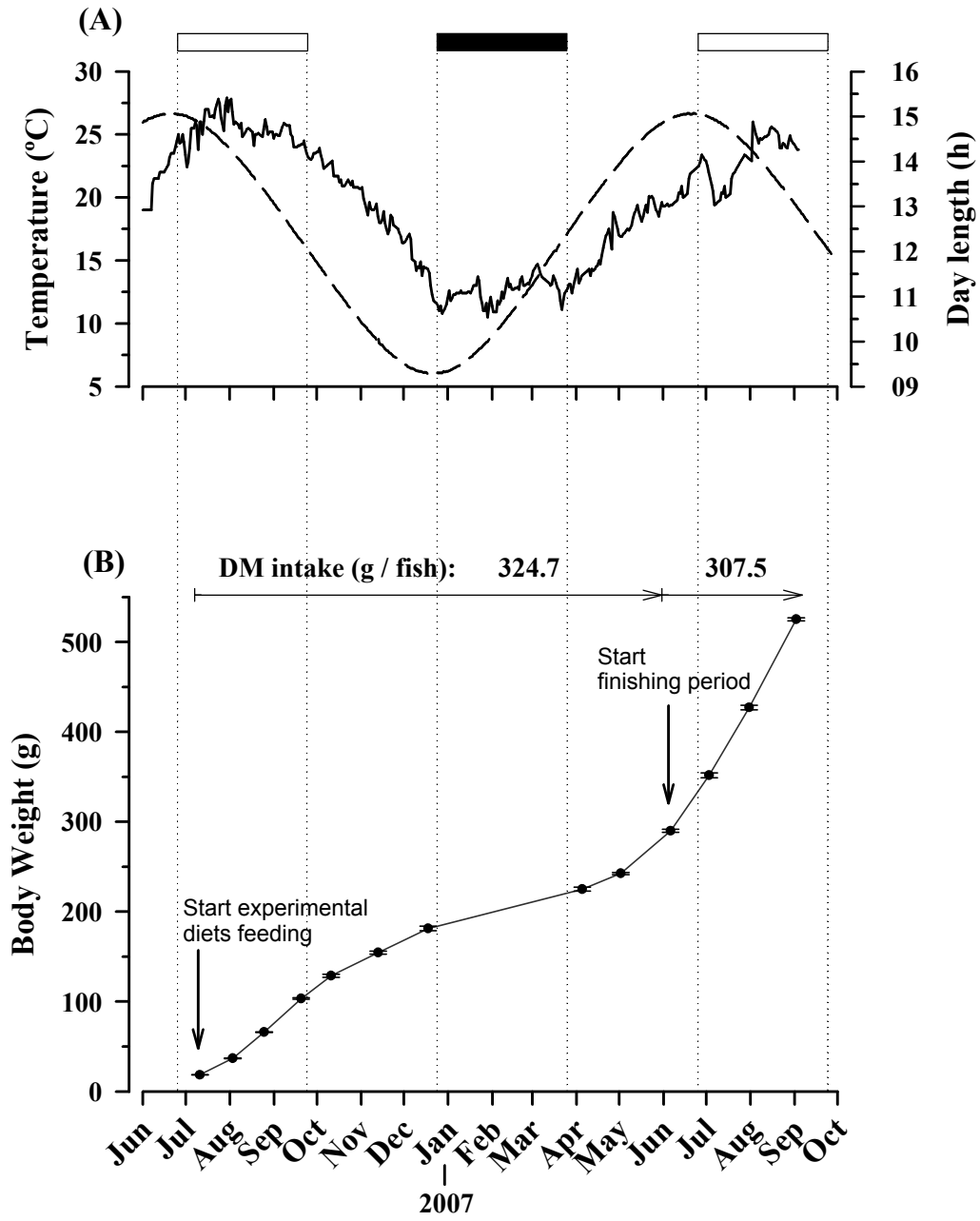


Figure 2

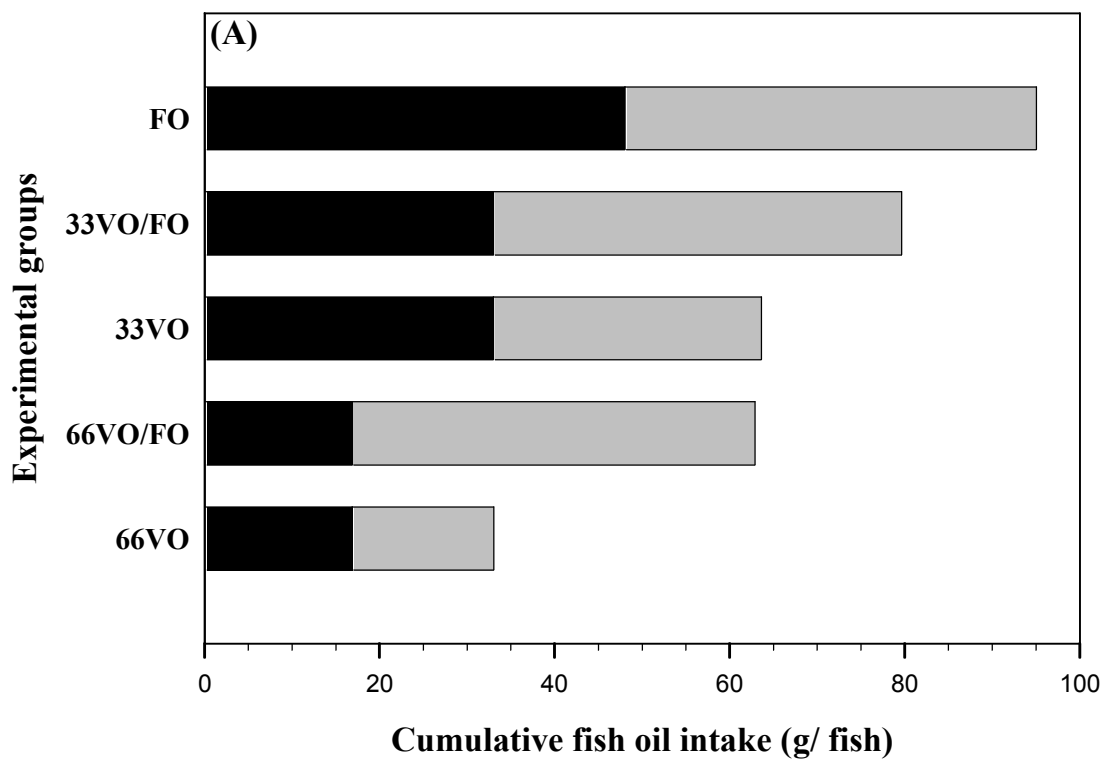


Figure 3

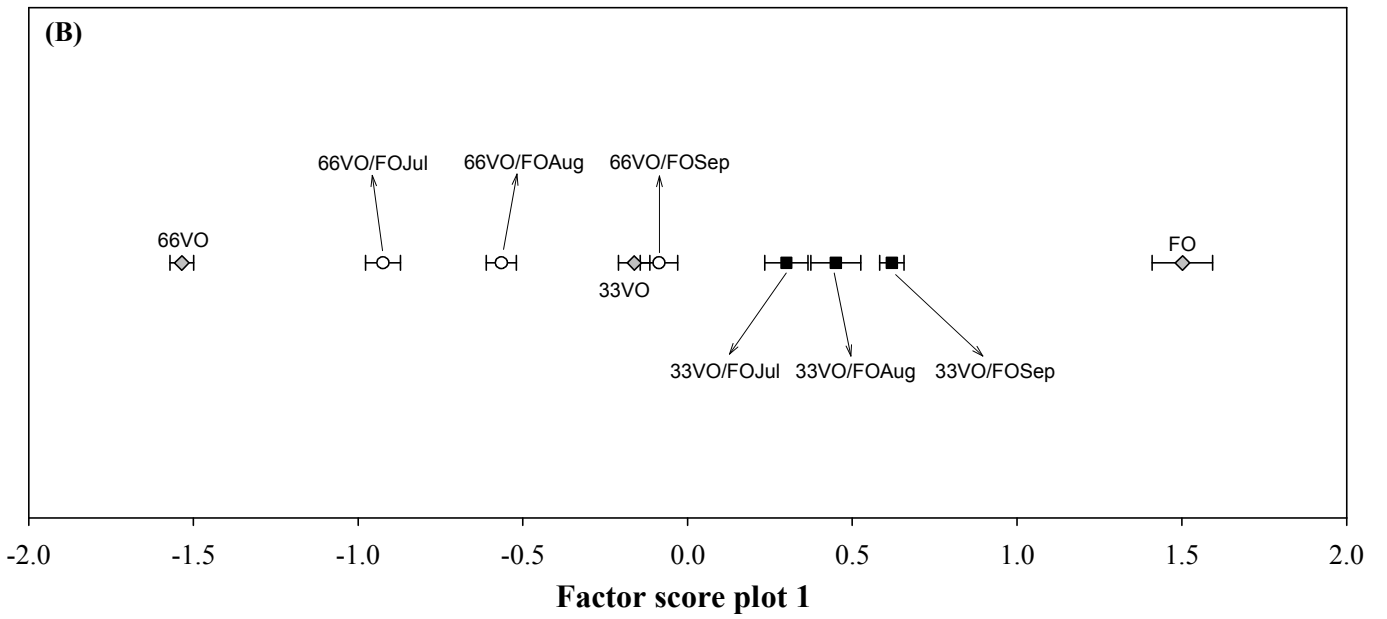
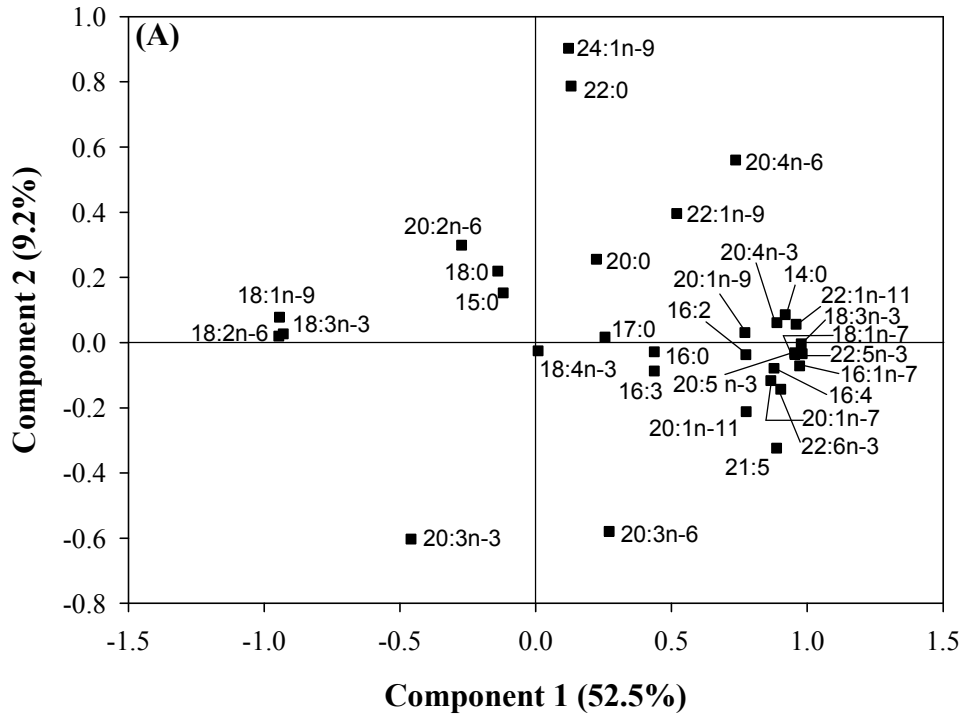


Figure 4

