1	VITAMIN C ENHANCES VITAMIN E STATUS AND REDUCES OXIDATIVE
2	STRESS INDICATORS IN SEA BASS LARVAE FED HIGH DHA MICRODIETS
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**Abbreviations**: α-TOH: α-tocopherol; AA: Ascorbic acid; AOE: Antioxidant enzymes; CAT: Catalase; dph: Days post hatching; DHA: Docosahexaenoic acid; EPA: Eicosapentaenoic acid; GPX: Glutathione peroxidase; IGF: insulin-like growth factors; MDA: Malonaldehyde; MUFA: Monoenics fatty acids; MyHC: Myosin heavy chain; ROS: Reactive oxygen species; SAFA: Saturated fatty acids; SOD: Superoxide dismutase; TBARS: Thiobarbituric acid reactive substances; PUFA: Polyunsaturated fatty acids; LC-PUFA: Long chain PUFA

#### 22 Abstract

23 Docosahexaenoic acid (DHA) is an essential fatty acid necessary for many biochemical, 24 cellular and physiological functions in fish. However, high dietary levels of DHA 25 increase free radical injury in sea bass larvae muscle, even when vitamin E (atocopherol,  $\alpha$ -TOH) is increased. Therefore, the inclusion of other nutrients with 26 complementary antioxidant functions, such as vitamin C (ascorbic acid, AA), could 27 28 further contribute to prevent these lesions. The objective of the present study was to 29 determine the effect of AA inclusion (3600 mg/kg) in high DHA (5% DW) and  $\alpha$ -TOH 30 (3000 mg/kg) microdiets (diets 5/3000 and 5/3000+AA) in comparison to a control diet 31 (1% DHA DW and 1500 mg/kg of  $\alpha$ -TOH; diet 1/1500) on sea bass larvae growth, 32 survival, whole body biochemical composition and thiobarbituric acid reactive 33 substances (TBARS) content, muscle morphology, skeletal deformities and antioxidant 34 enzymes, insulin-like growth factors (IGFs) and myosin expression (MyHC). Larvae 35 fed diet 1/150 showed the best performance. IGFs gene expression was elevated in 36 5/3000 larvae, suggesting an increased muscle mitogenesis that was confirmed by the 37 increase in the mRNA copies of MyHC. AA effectively controlled oxidative damages in 38 muscle, increased a-TOH larval contents and reduced TBARS content and the 39 occurrence of skull deformities. The results of the present study showed the antioxidant 40 synergism between vitamins E and C when high contents of DHA are included in sea 41 bass larvae diets.

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45 Key words: DHA, vitamin E, vitamin C, muscle morphology, antioxidant enzymes,
46 IGF, MyHC

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The importance of polyunsaturated fatty acids (PUFA) for marine fish larvae has 49 50 been extensively studied during the last 20 years (1,2,3,4) in order to perfect its culture, 51 especially in relation to eicosapentaenoic acid (EPA; 20:5n-3) and docosahexaenoic 52 acid (DHA; 22:6n-3) (1,5, 6,7). The particular structure of DHA provides this fatty acid 53 with many important functions in fish metabolism (4), and has been demonstrated to be 54 superior to EPA in promoting growth and conferring vitality to larvae (1,8), being preferentially incorporated into biomembranes (6,9). Besides, it has been reported that 55 56 DHA content in marine fish larvae rapidly decreases during the first ten days after 57 hatching, therefore high contents of DHA must be supplied to larvae in order to 58 maintain the adequate levels of DHA in growing larvae (1). Due to their limited 59 capacity to synthesize DHA and EPA from their precursors, fish require diets rich in 60 these fatty acids (10). Nowadays, live prey substitution by compound diets is crucial for 61 lowering costs and increasing production quality, mainly by reducing the incidence of 62 skeletal deformities and increasing welfare in fingerlings, using a formulation having 63 high levels of DHA, similar to those of live prey (10). However, DHA is very 64 susceptible to attack by reactive oxygen species (ROS) due to its high unsaturation 65 degree (11).

ROS are produced during normal cellular function (12), being beneficial or even indispensable at low concentrations in processes such as defense against microorganisms, contributing to phagocitic bactericidal activity. Fish possess enzyme systems and low-molecular-weight molecules with antioxidant functions capable of neutralizing ROS and protecting against their adverse effects (13). However, ROS generation can exceed its removal and oxidative stress can occur (14) with ROS

72 attacking diverse cellular components. One of the consequences of oxidative stress is 73 the oxidative peroxidation of PUFA, known as lipid peroxidation. The high unsaturation 74 content of these fatty acids renders them very susceptible to lipid oxidation. Thus, the 75 high requirements of marine fish larvae for long chain PUFA (LC-PUFA), mainly DHA 76 and EPA, makes them more prone to suffering peroxidative attack than are adults (15). 77 Therefore, the importance of nutrition in the pro-oxidant-anti-oxidant balance process 78 may be highly critical for fish larvae, as their high LC-PUFA contents disposes larval 79 tissues vulnerable to oxidative stress and an increase in the content of antioxidant 80 nutrients is essential. Among the antioxidant nutrients, vitamin E (tocopherols and 81 tocotrienols) is the major membrane-bound lipid-soluble antioxidant (13), whereas 82 vitamin C (ascorbic acid, AA) is an important water-soluble antioxidant which protects 83 low density lipoproteins from oxidation and is required for the correct formation of 84 cartilage (16). AA is easily oxidized to the unstable dehydroascorbic acid (DHAA), 85 which is not normally detectable in plasma but may develop transiently during oxidative 86 stress (17). The presence of sparing mechanisms between both vitamins was first 87 hypothesized by Tappel (18). This hypothesis proposes that the oxidized  $\alpha$ -tocopherol 88  $(\alpha$ -TOH) is reduced by ascorbate, thereby regenerating  $\alpha$ -TOH. In some fish species the 89 presence of a vitamin C/E sparing mechanism has been suggested (19,20,21,22,23) 90 reporting an influence on growth, tissue composition or immune responses. For 91 instance, supplementation with 100 mg/kg of ascorbyl-2-polyphosphate to an  $\alpha$ -TOH 92 deficient diet in juvenile channel catfish (Ictalurus punctatus) decreased vertebral 93 deformities and improved weight gain, feed intake and feed efficiency rate (23). High 94 supplementation of ascorbate might also spare  $\alpha$ -TOH in diets for hybrid tilapia 95 (Oreochromis niloticus x O. aureus) as shown by the increased weight gain, feed 96 efficiency and  $\alpha$ -TOH concentrations. However, little is known about the effect of both 97 vitamins in preventing oxidative stress in fish larvae, when high levels of LC-PUFA are98 administered.

99 The potentially deleterious effects of ROS are counteracted by a suite of 100 antioxidant enzymes (AOE), including radical-scavenging enzymes such as catalase 101 (CAT; EC 1.15.1.1) and superoxide dismutase (SOD; EC 1.11.1.19) or peroxidases such 102 as glutathione peroxidase (GPX; EC 1.11.1.6). In Manchurian trout larvae 103 (Brachymystax lenok) high lipid content microdiets stimulated the activity of AOE, 104 generally accompanied by an increase in malonaldehyde (MDA) contents (24). In 105 contrast, Mourente et al. (25) did not find a direct relationship between the activity of 106 AOE and the level of dietary n-3 LC-PUFA in Dentex dentex larvae, but decreased a-TOH and increased MDA contents were found in larvae fed high n-3 LC-PUFA 107 108 enriched Artemia. Furthermore, it seems that the level of antioxidant enzymes rises with 109 larval development, whereas the level of antioxidant molecules falls (26). In mammals, 110 ROS can induce changes in gene expression during normal development (27) causing 111 defective embryo development and retardation of embryo growth (28). Similarly, 112 oxidative stress causes embryonic mortality and developmental arrest in sea urchins 113 (Paracentrotus lividus and Spherechinus granularis) larvae (29). Oxygen stress during 114 early larval stages of fish may also alter development. Somatic growth in fish is 115 regulated by insulin-like growth factors I and II (IGF-I and IGF-II), two single chain 116 polypeptides that also have a function as myogenic regulatory factors which increase 117 satellite cell proliferation and differentiation (30,31). In addition, IGFs stimulate 118 cartilage growth by the incorporation of sulphate into cartilage and also affect cell 119 differentiation, growth and proliferation (32,33). Differences in myogenesis regulation, 120 such as myosin isoform expression, have also been observed in the earliest stages and 121 during temperature acclimation (34,35,36). However, no information is available about the effect of different dietary components on the regulation of the various componentsof the IGF signaling pathways, as well as their role on muscle growth in fish (37).

124 In previous studies we have shown the alteration of sea bass (Dicentrarchus 125 *labrax*) larvae oxidative status when they were fed high levels of DHA (5%), with  $\alpha$ -126 TOH having a limited effect in preventing alterations such as muscular dystrophy or 127 hepatic ceroidosis (38,39). Therefore, the aim of the present study was to evaluate the 128 combined effect of  $\alpha$ -TOH and AA in preventing oxidative stress in sea bass larvae fed 129 high levels of DHA in relation to larval performance or antioxidant status as well as the 130 incidence of morphological alterations, including the expression of selected related 131 genes.

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### 133 Material and methods

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135 **Fish** 

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137 The experiment was carried out at the Grupo de Investigación en Acuicultura 138 facilities (Telde, Canary Islands, Spain). Sea bass, Dicentrarchus labrax, larvae were 139 obtained from a natural spawning from the Instituto de Acuicultura de Torre de la Sal 140 (CSIC, Castellón, Spain). Prior to starting the feeding experiment, larvae were fed enriched yeast-fed rotifers (DHA Protein Selco<sup>®</sup>, INVE, Belgium) until they reached 14 141 142 days post hatching (dph). Then, larvae (total length  $8.58 \pm 0.64$  mm, dry body weight 143  $0.36 \pm 0.00$  mg) were randomly distributed in experimental tanks (n=9) at a density of 144 1000 larvae/tank and were fed one of the experimental diets for 21 days, at a water 145 temperature of 19.5 to 21.0°C. All tanks (170 L light grey color cylindrical fibreglass 146 tanks) were supplied with filtered sea water (34 ‰ salinity) at an increasing rate of 1.0 -

147 1.5 L/min along the feeding trial. Sea water entered the tank from bottom to top; water
148 quality was tested daily and no deterioration was observed. Water was continuously
149 aerated (125 mL/min), attaining 5-8 g/L dissolved O<sub>2</sub> and 60-80% saturation.

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151 **Diets** 

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153 Three isonitrogenous and isolipidic experimental microdiets (pellet size  $< 250 \mu m$ ) 154 were prepared containing two levels of DHA, AA and  $\alpha$ -TOH (Table 1). A low 155 oxidation risk diet (1/1500) contained low DHA (1% DW) and α-TOH (1500 mg/kg); a 156 high oxidation risk diet contained the highest DHA (5% DW) and  $\alpha$ -TOH (3000 mg/kg) 157 levels and a third diet contained high DHA (5% DW) and  $\alpha$ -TOH (3000 mg/kg) 158 together with increased AA (3600 mg/kg). The protein source used was squid meal 159 defatted 3 consecutive times with a chloroform:squid meal ratio of 3:1 to allow a better 160 control of the fatty acid profile of the microdiet. EPA, DHA, a-TOH and AA sources 161 used in the experimental diets were EPA50 and DHA50 (CRODA, East Yorkshire, 162 England, UK), DL- a- Tocopheryl Acetate (Sigma-Aldrich, Madrid, Spain) and 163 Rovimix Stay-C 35 (Roche, Paris, France) respectively. Oleic acid (Merck, Darmstadt, 164 Germany) was added to equalize the lipid content in all diets. Microdiets were prepared 165 according to Liu et al. (40) by first mixing the squid powder and water soluble 166 components, followed by lipid- and fat- soluble vitamins and, finally, warm water 167 dissolved gelatin. The paste was pelleted and oven dried at 38°C for 24 h. Pellets were 168 ground and sieved to obtain particle size below 250 µm. Diets were analyzed for 169 proximate and fatty acid composition of dry matter and kept in plastic bags under 170 nitrogen atmosphere at -20°C until the beginning of the experimental trial. Diets were 171 manually supplied 14 times per day every 45 min from 9:00-19:00. Daily feed supplied was 2, 2.5 and 3 g/tank during the first, second and third week of feeding respectively.
Each diet was tested in triplicate.

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#### 175 **Growth and survival**

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Final survival was calculated by individually counting live larvae at the beginning and end of the experiment. Growth was determined by measuring dry body weight (105°C for 24 hours) and total length (Profile Projector V-12A Tokyo, Nikon) of 30 larvae/tank at the beginning, middle and end of the trial.

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#### 182 **Biochemical analysis**

All remaining larvae in each tank were washed with distilled water, sampled and kept at -80°C for biochemical composition and TBARS analyses after 12 hours of starvation at the end of the trial. Moisture, protein (41) and lipid (42) contents of the larvae and diets were analyzed.

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## 188 Total lipid fatty acid analysis

Fatty acid methyl esters (FAMEs) were obtained by transmethylation of total lipids as described by Christie (43). FAMEs were separated by GLC, quantified by FID (GC -14A, Shimadzu, Tokyo, Japan) under the conditions described in Izquierdo *et al.* (44) and identified by comparison with previously characterized standards and GLC-MS.

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## 195 *Determination of* α-TOH *content*

196 α-TOH concentrations were determined in diets and larvae samples using high-197 pressure liquid chromatography (HPLC) with UV detection. Samples were weighed, homogenized in pyrogallol and saponified as described by McMurray et al. (45) for 198 199 diets or according to Cowey et al. (46) for larvae. HPLC analysis was performed using a 200 150 x 4.60 mm, reverse-phase Luna 5µm C18 column (Phenomenox, California, USA). 201 The mobile phase was 98% methanol pumped at 1.0 mL/min. The effluent from the 202 column was monitored at a wavelength of 293 nm and quantification achieved by 203 comparison with  $(+)-\alpha$ -tocopherol (Sigma-Aldrich, Madrid, Spain) as external standard.

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### Determination of AA content

206 Ascorbil-2-monophosphate concentrations were determined in diets using a 207 HPLC procedure with UV detection. The HPLC system comprised of a 150 x 4.6 mm, 5 208 µm particle size, Gemini C18 column fitted with a Gemini pre-column of the same 209 material. The mobile phase consisting of phosphate buffer was delivered at a flow rate 210 of 0.8 mL/min. Samples were dissolved with 0.4 M phosphate buffer (pH 3.0) and 211 centrifuged at 1610 x g for 5 minutes at room temperature. The supernatants were kept 212 at 4°C until assayed. AA concentrations were determined at a wavelength of 293 nm 213 and quantification achieved by comparison with tris(cyclohexylammonium) ascorbic 214 acid-2-phosphate (Sigma-Aldrich, Madrid, Spain), used as a reference substance.

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## 6 *Measurement of thiobarbituric acid reactive substances (TBARS)*

TBARS were measured in triplicate samples using a method adapted from that used by Burk *et al.* (47). Approximately 20-30 mg of larval tissues per sample were homogenized in 1.5 mL of 20% trichloroacetic acid (w/v) containing 0.05 mL of 1% BHT in methanol. To this 2.95 mL of freshly prepared 50mM thiobarbituric acid solution were added before mixing and heating for 10 minutes at 100°C. After cooling, protein precipitates were removed by centrifugation (Sigma 4K15, Osterode am Harz, Germany) at 2000 x g, the supernatant was read in a spectrophotometer (Evolution 300, Thermo Scientific, Cheshire, UK) at 532 nm. The absorbance was recorded against a blank at the same wavelength. The concentration of TBA-malondialdehyde (MDA) expressed as nmol MDA per g of tissue was calculated using the extinction coefficient  $0.156 \,\mu M^{-1} \,cm^{-1}$ .

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- 229 Histopathological sampling
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Thirty larvae from each tank were collected every seventh day from the beginning of the feeding trial, and fixed in 10% buffered formalin for 1 or 2 days, dehydrated through graded alcohols, then xylene and finally embedded in paraffin wax. Six paraffin blocks containing 5 larvae per tank were sectioned at 3 µm and stained with Hematoxilin and Eosin (H&E) for histopathologic evaluation (48).

236 Ten larvae per tank were fixed for 24 hours at 4°C in 2.5% glutaraldehyde in 0.2 237 M phosphate buffer (pH 7.2). Samples were then rinsed in phosphate buffer and post-238 fixed for 1 hour in 2% osmium tetraoxide in 0.2 M potassium ferrocyanide. Each larva 239 was then embedded in an Eppon/Araldite resin block. Serial transverse and longitudinal 240 larvae thick sections were cut at 1 µm, stained with toluidine blue and examined under 241 light microscopy (49). Thin sections were cut at 50 nm and stained with lead citrate 242 before observing with a ZEISS EM 910 transmission electron microscope (Germany) at 243 the Electron Microscope Service of University of Las Palmas de Gran Canaria.

244 Besides, 100 larvae from each tank at 35 dph were fixed in 10% buffered formalin 245 in order to perform deformity analyses. Prior to staining, larvae were measured under a

246 Profile Projector (Mitutoyo, PJ 3000, Japan), divided into three size classes (< 10.0 mm, 247 10.0-12.0 mm and >12.0 mm) and stained with Alizarin red (50) to determine bone 248 mineralization. Larvae from the different experimental groups were stained 249 simultaneously in order to prevent any technical variability. Deformities were classified 250 in three different groups according to their localization: cranial deformities, lordosis, 251 kyphosis, neural process alterations and others, including deformities such as vertebral 252 compression or scoliosis. Cranial deformities included those found in the jaws, such as 253 pugheadness or crossbite and opercular deformities. The axial skeleton deformities 254 includes spine curvatures such as lordosis (ventral curvature), kyphosis (dorsal 255 curvature) or scoliosis (lateral curvature); neural processes alterations like wrong 256 direction or twisting and vertebral compression that comprises two alterations, the 257 flattering of vertebral end plates and vertebral fusion. The surface corresponding to 258 bone in whole colored larvae was visualized and quantified using a computerized image analysis package (Image-Pro Plus<sup>®</sup>, Media Cybernetics, Maryland, USA). By selecting 259 260 ranges of pixel values in color images the pixels associated with red could be 261 distinguished. The number of selected pixels was then quantified using a particle 262 analysis operation and by counting the area of all bright objects (in pixels). Larval size 263 was estimated by calculating the surface areas (in pixels) covered by whole stained 264 larvae.

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## 266 **RNA extraction and quantitative RT-PCR**

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Molecular biology analyses were carried out at the University of Insubria (Varese,
Italy). Total RNA was extracted from sea bass larvae (≈200 mg; pool per tank), using
PureYield RNA Midiprep System (Promega, Italy). The quantity and purity of RNA

271 was assessed by spectrophotometer. Visualization on 1% agarose gel stained with 272 ethidium bromide showed that RNA was not degraded. Three micrograms of total RNA 273 was reverse transcribed into complementary DNA (cDNA) in a volume of 12 µl, 274 including 1 µl of oligo dT16 primer (50 pmol) and 1 µl of 10 mM deoxynucleotide 275 triphosphates (dNTPS). This mix was heated at 65°C for 5 minutes, chilled on ice and 276 then 4 µl of 5X reverse transcription buffer, 2 µl 0.1M DTT, 1 µl RNAse out and 1 µl of 277 Moloney murine leukemia virus (M-MLVRT) were added. After incubation at 37°C for 278 50 minutes, the reaction was stopped by heating at 75°C for 15 minutes.

279 PCR primers sequences used for the PCR amplification of the cDNAs of target 280 genes were CAT, SOD, GPX, IGF-I, IGF-II and MyHC. To perform PCR, an aliquot of 281 4  $\mu$ l of cDNA was amplified using 25  $\mu$ l GoTaq Green Master Mix (Promega, Italy) in 282 50  $\mu$ l of final volume and 50 pmol of each designed primer.

283 A total of 31 PCR amplification cycles (eight touchdown) were performed for all 284 primer sets, using an automated Thermal Cycler (MyCycler, BioRad, Italy). An aliquot 285 of each sample was then subject to electrophoresis on a 1% agarose gel in 1X TAE 286 buffer (Bio-Rad, Italy) and bands were detected by ethidium bromide. Samples were run 287 together with a 100 bp±1.5 kb DNA ladder to control molecular weight of the DNA. 288 The negative control (a reaction mixture without cDNA), confirmed the absence of genomic contamination. The PCR products from each primer set amplification were 289 then cloned using pGEM<sup>®</sup>-T easy vector (Promega, Italy) and subsequently sequenced 290 291 in both directions (T7 and SP6).

TaqMan<sup>®</sup> real time reverse transcription PCR was performed on a StepOne Real Time PCR System (Applied Biosystems, Italy) using Assays-by-Design<sup>SM</sup> PCR primers (Applied Byosystems) and gene-specific fluorogenic probes. Primer sequences and TaqMan<sup>®</sup> probes of target genes were the following: 296

# 297 Target gene: Sea bass CAT

298	Forward primer: 5'- ATGGTGTGGGGACTTCTGGAG - 3'
299	Reverse primer: 5'- GCTGAACAAGAAAGACACCTGATG - 3'
300	TaqMan <sup>®</sup> probe: 5'- CAGACACTCAGGCCTCA - 3'
301	Target gene: Sea bass SOD
302	Forward primer: 5'- TGGAGACCTGGGAGATGTAACTG - 3'
303	Reverse primer: 5'- TCTTGTCCGTGATGTCGATCTTG - 3'
304	TaqMan <sup>®</sup> probe: 5'- CAGGAGGAGATAACATTG - 3'
305	Target gene: Sea bass GPX
306	Forward primer: 5'- AGTTAATCCGGAATTCGTGAG - 3'
307	Reverse primer: 5'- AGCTTAGCTGTCAGGTCGTAAAAC - 3'
308	TaqMan <sup>®</sup> probe: 5'- AATGGCTGGAAACGTG - 3'
309	Target gene: Sea bass IGF-I
310	Forward primer: 5'- GCAGTTTGTGTGTGGAGAGAGA- 3'
311	Reverse primer: 5'- GACCGCCGTGCATTGG - 3'
312	TaqMan <sup>®</sup> probe: 5'- CTGTAGGTTTACTGAAATAAAA - 3'
313	Target gene: Sea bass IGF-II
314	Forward primer: 5'- TGCAGAGACGCTGTGTGG - 3'
315	Reverse primer: 5'- GCCTA CTGAAATAGAAGCCTCTGT - 3'
316	TaqMan <sup>®</sup> probe: 5'- CAAACTGCAGCGCATCC - 3'

# 317 Target gene: Sea bass MyHC

318	Forward primer: 5' - TGGAGAAGATGTGCCGTACTCT - 3'
319	Reverse primer: 5'- CGTGTCATTGATTTGACGGACATTT - 3'
320	TaqMan <sup>®</sup> probe: 5'- AACTGAGTGAACTGAAGACC - 3'
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322	Data from TaqMan <sup>®</sup> PCR runs were collected with ABI's Sequence Detector
323	Program. Cycle threshold (Ct) values corresponded to the number of cycles at which the
324	fluorescence emission monitored in real time exceeded the threshold limit. The Ct
325	values were used to create standard curves to serve as a basis for calculating the
326	absolute amounts of mRNA in total RNA. To reduce pipetting errors, master mixes
327	were prepared to set up duplicate reactions (2 x 30 $\mu$ l) for each sample.
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329	Statistical analysis
329 330	Statistical analysis
329 330 331	Statistical analysis Results are given as mean $\pm$ SD. Survival, growth, fatty acid, ossification degree
<ul><li>329</li><li>330</li><li>331</li><li>332</li></ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances
<ul> <li>329</li> <li>330</li> <li>331</li> <li>332</li> <li>333</li> </ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances with Levene's test, not requiring any transformation. Chi-squared test was employed for
<ul> <li>329</li> <li>330</li> <li>331</li> <li>332</li> <li>333</li> <li>334</li> </ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances with Levene's test, not requiring any transformation. Chi-squared test was employed for incidence of muscular lesions, deformities and TBARS content. Survival, growth, fatty
<ul> <li>329</li> <li>330</li> <li>331</li> <li>332</li> <li>333</li> <li>334</li> <li>335</li> </ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances with Levene's test, not requiring any transformation. Chi-squared test was employed for incidence of muscular lesions, deformities and TBARS content. Survival, growth, fatty acid, ossification degree and molecular biology data were treated by one-way ANOVA.
<ul> <li>329</li> <li>330</li> <li>331</li> <li>332</li> <li>333</li> <li>334</li> <li>335</li> <li>336</li> </ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances with Levene's test, not requiring any transformation. Chi-squared test was employed for incidence of muscular lesions, deformities and TBARS content. Survival, growth, fatty acid, ossification degree and molecular biology data were treated by one-way ANOVA. Means were compared by Duncan's test (P<0.05) using SPSS software (SPSS for
<ul> <li>329</li> <li>330</li> <li>331</li> <li>332</li> <li>333</li> <li>334</li> <li>335</li> <li>336</li> <li>337</li> </ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances with Levene's test, not requiring any transformation. Chi-squared test was employed for incidence of muscular lesions, deformities and TBARS content. Survival, growth, fatty acid, ossification degree and molecular biology data were treated by one-way ANOVA. Means were compared by Duncan's test (P<0.05) using SPSS software (SPSS for Windows 14.0; SPSS Inc., Chicago, IL, USA, 2005). For analysis of one-way ANOVA
<ul> <li>329</li> <li>330</li> <li>331</li> <li>332</li> <li>333</li> <li>334</li> <li>335</li> <li>336</li> <li>337</li> <li>338</li> </ul>	Statistical analysis Results are given as mean ± SD. Survival, growth, fatty acid, ossification degree and molecular biology data were tested for normality and homogeneity of variances with Levene's test, not requiring any transformation. Chi-squared test was employed for incidence of muscular lesions, deformities and TBARS content. Survival, growth, fatty acid, ossification degree and molecular biology data were treated by one-way ANOVA. Means were compared by Duncan's test (P<0.05) using SPSS software (SPSS for Windows 14.0; SPSS Inc., Chicago, IL, USA, 2005). For analysis of one-way ANOVA the following general linear model was used:

340

341 where  $Y_{ij}$  is the mean value of the tank, m is the mean population,  $D_i$  is the fixed effect 342 of the diet and  $e_{ij}$  is the residual error.

- 343
- 344 **Results**

345 The diet containing about 1% DHA (diet 1/1500) showed higher amounts of 346 monoenoic fatty acids than diets containing 5% DHA (5/3000 and 5/3000+AA) due to 347 the higher oleic acid content in the former diet (Table 2). DHA contents in the diets 348 varied from 4.58% of total fatty acids in 1/1500 diet, 24.55% in 5/3000 diet and 27.54% 349 in 5/300+AA diet. Elevation of dietary DHA (5/3000 diets) increased n-3 PUFA and n-350 3 LC-PUFA fatty acids contents, as well as n-3/n-6 PUFA ratio. α-TOH levels were 351 more than 2 times higher in diets containing 3000 mg/kg compared to the control diet 352 (1/1500) (Table 3). AA contents were also higher in the diet supplemented with AA 353 than in the others.

All experimental diets were well accepted by larvae. The highest total length was found in 1/1500 larvae. Increasing DHA from 1 to 5% in 5/3000 larvae, significantly reduced larvae total length, despite the  $\alpha$ -TOH increase. However, AA levels in 5/3000+AA diet enhanced larval total length compared to 5/3000 larvae (*P*=0.005). Sea bass larvae survival or dry weight was not significantly different among the treatments (Table 4).

The level of lipid peroxidation-derive aldehydes, as indicated by TBARS content (nmol/g larval tissues) was lowest in larvae fed diet 1/1500 and was significantly higher in larvae fed the higher DHA content (5/3000 and 5/3000+AA diets). However, the inclusion of AA prevented the formation of hydroperoxides, observed by a decrease in TBARS levels. Thus, the lowest  $\alpha$ -TOH content was found in larvae fed 5/3000 diet. However, an increase in dietary AA contents increased  $\alpha$ -TOH levels in larvae fed 366 5/3000+AA diet. Regarding larval  $\alpha$ -TOH contents, the elevation of dietary vitamin E in 367 diet 5/3000, together with the increase in DHA, in comparison to diet 1/1500, did not 368 significantly affect the  $\alpha$ -TOH contents in the larvae (Table 4).

369 Fatty acid composition of the larvae (Table 5) generally reflected the fatty acid 370 composition of the diet. Accordingly, a higher content of 18:1n-9 was observed in 371 1/1500 larvae. However, its retention rate regarding dietary levels was much lower in 372 1/1500 larvae (47.30%) than in 5/3000 (67.22%) or 5/3000+AA larvae (67.82%), 373 balancing the monoenoic acids content among larvae fed the different dietary 374 treatments. Equally, EPA retention was low in all larvae, especially in those fed 5/3000 375 (43.93%) and 5/3000+AA (43.96%) diets. The highest content of total n-3 LC-PUFA 376 (P=0.006) was observed in larvae fed the diet supplemented with AA probably due to a 377 higher dietary content. However, regarding 22:6n-3 content, the highest retention rate 378 was observed in 1/1500 larvae. Similarly 20:4n-6 was highly retained in larvae fed 379 1/1500 diet, although higher contents were found in larvae fed diets 5/3000. In contrast 380 to the differences in the n-3/n-6 ratio observed in diets, no differences were observed in 381 larvae among the different treatments.

382 Histopathological examinations revealed the presence of lesions affecting larvae 383 axial musculature, showing the typical features of necrotic degeneration of muscle. The 384 incidence of muscular lesions increased with DHA dietary content (5/3000 diet; Table 385 4). However, inclusion of AA (5/3000+AA diet) proved to be effective in reducing 386 incidences to less than half of those in 5/3000 larvae. More detailed features of 387 muscular lesions could be observed on semithin and TEM sections, where muscle 388 degeneration with the presence of hydropic vacuoles and organelles swelling within 389 some of the affected fibers was found in larvae fed 5/3000 diet. Some fibres presented disarrangement of the myofilaments just like myelin figures, denoting intracellular lipidoxidation (Figure 1).

392 Regarding skeletal morphology, among larvae measuring 10-12 mm, 393 28.33±1.30% of the larvae fed 1/1500 diet showed skeletal deformities, and similar 394 values were found in those fed the 5/3000+AA diet (29.67±6.51% of the larvae 395 anlayzed). Larvae fed the 5/3000 diet presented 33.11±5.11% deformities, but no 396 statistical differences were detected between groups (P=0.10; Table 4). The ossification 397 degree of 35 dph sea bass larvae was determined in terms of surface of mineralized 398 bones per larval surface (Table 4). High dietary levels of DHA decreased the formation 399 of mineralized bone in larvae, whereas increase in AA did not affect mineralization.

Different types of deformities were observed at the end of the experimental period, depending on the level of dietary DHA. All of the experimental groups exhibited a statistically similar percentage of kyphosis, however, no lordosis was observed in larvae fed the 1/1500 diet (Figure 2). Skull deformities were especially high in fish fed high DHA levels, although AA increase reduced the incidence of this deformity (P=0.013).

406 The general pattern of gene expression, excepting IGF-II, in all groups of sea 407 bass larvae was characterized by a rapid increase between 14 and 26 dph (Figures 3 and 408 4). CAT gene expression was elevated in larvae fed diets containing a high content of 409 DHA, the highest number of mRNA copies being found in larvae fed diet 5/3000+AA 410 (P=0.027; Figure 3A). The SOD mRNA expression was also highest in 5/3000 groups 411 (P=0.048; Figure 3B). Accordingly, GPX gene expression was quite strong in larvae fed 412 diets containing a high level of DHA compared to larvae fed low levels (P=0.039; 413 Figure 3C).

The IGF-I gene expression increased from 14 to 26 dph in all treatments, at day 35 showing a decrease in 1/1500 and 5/3000+AA larvae and a continuous increase in 5/3000 fed larvae (Figure 4A). Regarding IGF-II, gene expression was higher in larvae fed 5/3000 and 5/3000+AA diets than in those fed diet 1/150 throughout the trial (Figure 4B).

419 MyHC gene expression was low in larvae fed diet 1/1500 and significantly 420 increased by the elevation of both  $\alpha$ -TOH and DHA in diet 5/3000 (Figure 4C). 421 However, increase in AA in diet 5/3000+AA significantly reduced MyHC expression to 422 levels similar to those of 1/1500 diet.

423

#### 424 **Discussion**

425 Marine fish larvae are subjected to high levels of oxidative stress when using inert 426 diets due to the high content of LC-PUFA, particularly DHA, and pro-oxidants such as 427 minerals as well as the high surface to volume ratio of these feed particles (51). 428 Therefore, inclusion of high dietary levels of LC-PUFA to match the high requirements 429 of marine fish larvae may call for increased dietary supplementation with antioxidants 430 such as  $\alpha$ -TOH to prevent oxidative damage. For instance, increasing DHA by up to 5% 431 in diets for sea bass markedly reduced larval survival and growth and increased the 432 incidence of muscular lesions (38). Despite an increase in dietary  $\alpha$ -TOH from 1500 to 433 3000 mg/kg which partially reduced the occurrence of muscular alterations at DHA 434 dietary levels up to 3% (38), lesions caused by the further elevation of DHA up to 5% 435 could not be prevented by the increase in dietary  $\alpha$ -TOH (3000 mg/kg) (39). Similarly, 436 in the present study high levels of  $\alpha$ -TOH (3000 mg/kg) together with high DHA (5%) 437 were not able to counteract the adverse effects of lipid oxidation on the incidence of 438 muscular lesions. Accordingly, these larvae showed very high levels of TBARS

439 indicating that their oxidative status is altered when they are fed high DHA levels even 440 at such high dietary  $\alpha$ -TOH levels, in agreement with our previous studies (39). 441 Moreover, AOE expression was higher in those larvae, denoting a high antioxidant 442 response. A compensatory induction of these endogenous antioxidants is found in 443 animals exposed to dietary oxidative stress (52). Indeed,  $\alpha$ -TOH contents in these larvae 444 were not increased by the elevation of dietary  $\alpha$ -TOH levels, suggesting a depletion of 445 this vitamin to neutralize ROS and its limited capacity to prevent lipid peroxidation 446 under these conditions. Similarly, increased n-3 LC-PUFA did cause depletion of  $\alpha$ -447 TOH contents when this vitamin was supplemented to the diet in previous studies 448 (51,53, 54).

449 In contrast, the increase in AA dietary supplementation from 1800 to 3600 mg/kg, 450 markedly improved the protection against peroxidation, decreasing TBARS contents to 451 less than one third, sparing vitamin E that was significantly increased in larval tissues 452 and effectively reducing the incidence of muscular lesions. AA supplementation has 453 also been found to reduce TBARS formation in hybrid striped bass (Morone chrysops Q454 x *M. saxatilis*  $\mathcal{A}$ ) fed diets deficient in  $\alpha$ -TOH (21). The increased  $\alpha$ -TOH content found 455 in the present study when larvae were fed high levels of AA, is in agreement with 456 studies in other fish species (20.22,51) and denotes the sparing effect of AA on  $\alpha$ -TOH 457 in sea bass larvae, the first vitamin recycling the second one. Thus, under dietary 458 conditions of high LC-PUFA and  $\alpha$ -TOH, AA dietary contents of 1800 mg/kg may not 459 be sufficient to recycle  $\alpha$ -TOH and prevent the high rate of ROS formation and, 460 therefore, AA requirements may be higher than under low oxidation risk dietary 461 situations.

462 AA is known to be a powerful antioxidant, by efficiently trapping peroxyl radicals 463 in the aqueous phase before they can initiate lipid peroxidation, thereby protecting the

biomembranes (55). Thus, the decrease in the incidence of muscular lesions observed in 464 465 the present study, when high levels of AA were employed, could be due to the 466 protective effect of this antioxidant nutrient, quenching free radicals before they can 467 attack muscular membranes, in addition to recycling  $\alpha$ -TOH. Moreover, the species 468 formed after the loss of one electron are relatively stable and fairly unreactive compared 469 to tocopheroxyl radical (56), thus preventing a pro-oxidant action. The reduced 470 incidence of muscular lesions was also accompanied by a decrease in IGF-I expression, 471 the highest IGF mRNA copies occurring in larvae fed the highest DHA and  $\alpha$ -TOH 472 levels, which also showed the lowest growth and the highest incidence of muscular 473 lesions. These results are in line with the higher expression of IGF-I found in sea bream 474 larvae fed high DHA levels (57). On the contrary to the present study, in sea bream, 475 high IGF-I expression was correlated with high growth (57), suggesting the negative 476 effect of high DHA contents in sea bass growth in relation to the altered oxidative 477 status. A feasible explanation for the overexpression of IGF-I in larval groups with 478 higher TBARS values could be a compensatory mechanism in fish larvae to try to 479 counteract the adverse effects of ROS since IGF-I interferes with activation of apoptosis 480 in several cells and organ systems in mammals (58). For instance, an increase in IGF-I 481 has been found to suppress oxidative stress in atherosclerotic Apo-E deficient mice (59). 482 Furthermore, the IGF system can promote muscle growth and differentiation in fish, by 483 activating cell proliferation and DNA synthesis. Thus, the increase in mRNA copies of 484 IGF-I observed in larvae fed 5% DHA and 3000 mg/kg α-TOH could be also due to the 485 compensatory regeneration process carried out by satellite cells, and which was not 486 directly reflected on a growth improvement. This is supported by the results of MyHC 487 expression, as a high expression of myosin has been associated with regeneration 488 processes in sea bream after mechanical injury (60). In addition, in the present study,

IGF-I and MyHC expression follows a similar pattern, indicating that their biological functions may be interrelated. In this sense, it is known that IGF-I overexpression results in greater skeletal muscle mass in fine flounder (*Paralichthys adspersus*, 61) and in mice, in which IGF-I can activate MyHC as well as other transcriptional factors (62). Thus the parallel increase in IGF-I and MyHC expression observed in the present study confirms that when sea bass larvae are subjected to oxidative stress, a compensatory overexpression of genes related to cell/muscle proliferation occurs.

496 In the present study, AA elevation did not reduce the expression of AOE genes, 497 suggesting an antioxidant effect independently of these enzymes, but acting in parallel 498 with them to quench ROS. AA acts as a cofactor for at least eight enzymes involved in 499 the biosynthesis of collagen and carnitine, conversion of the neurotransmitter dopamine 500 to noradrenaline, metabolism of tyrosine and amidation of peptide hormones. In this 501 sense, AA acts with peptidyl-glycine alpha-amidating monooxygenase (PAM), an 502 enzyme that adds amide groups to peptide hormones, greatly increasing their stability 503 (63, 64). Thus, the antioxidant role of AA in fish might not only be reduced to trapping 504 peroxyl radicals from the aqueous phase or recycling  $\alpha$ -TOH, but also to support the fo 505 rmation of molecules with sound antioxidant potential. More studies are required to 506 clarify the interrelations between the different components of antioxidant defenses in 507 marine fish, as well as to corroborate if the AOE activity really reflects their gene expression. 508

509 In terms of skeletal deformities, it can be observed that a DHA increase up to 5% 510 raised the incidence of alterations in chondroid bone, such as that of the cranium, 511 whereas no differences were found in other deformities attaining intramembranous 512 bone, such as kyphosis. These results match with previous studies on sea bream larvae 513 fed high DHA rotifers (5.2% DW; 57), as ROS are known to actively destroy cartilage

514 tissue (65), therefore affecting chondroid bones with characteristics of cartilage rather 515 than directly affecting intramembranous bones. However, in the same study, in contrast 516 to the present one, the incidence of cranial deformities was reduced when high contents 517 of  $\alpha$ -TOH were included in the rotifer enrichment media, in relation to the reduced 518 TBARS and AOE expression. In another study from our research group (66) inclusion 519 of organic selenium to diets containing high DHA and  $\alpha$ -TOH, was enough to decrease 520 the TBARS values, but not to reduce cranial deformities, suggesting that the appearance 521 of these kind of deformity could be not only related directly to the DHA oxidation, but 522 also to the deficit of AA due to the pro-oxidant environment originated by the high 523 levels of α-TOH and LC-PUFA. Similarly, in the present study, an extra dosage of AA 524 proved to be efficient in reducing the incidence of cranial deformities when high levels 525 of DHA are included in the diets. Apart from being a potent antioxidant, AA acts with 526 three enzymes that participate in collagen hydroxylation by adding hydroxyl groups to 527 the aminoacids proline or lysine in the collagen molecules, greatly increasing stability 528 of the collagen (56). Thus, the protective effect of AA on chondroid bones could be due 529 both to its antioxidant activity and to a higher stability in the cartilage formation. Recent 530 works in sea bass larvae also showed a reduction in the incidence of cranial deformities 531 when enhanced levels of AA were included in the diet (50 mg/kg; 67). However, in the 532 same study, elevated levels of this nutrient (400 mg/kg) caused a similar percentage of 533 deformities to diets with a deficiency in AA. In the present work, the levels of AA 534 employed are much more elevated (1800 mg/kg) but it also has to be noted that levels of 535 AA higher than those required for growth are necessary to satisfy the demands of other 536 nutrients, in this case, to counteract the depletion in  $\alpha$ -TOH caused by ROS.

537 The appearance of lordosis in fish fed high DHA and  $\alpha$ -TOH content could be 538 related to the high IGF-I expression observed in these larvae caused by an imbalance in 539 the development of the musculoskeletal system, in agreement with previous studies 540 (68). On the one hand, the high incidence of muscular lesions occurring in these larvae, 541 may also contribute to increase the lordosis rate, as a result of the increased muscular 542 tensions created during tissue regeneration. In this sense, Madsen and Dalsgaard (69) 543 showed that the rainbow trout fry syndrome, characterized by muscular dystrophy 544 among other pathologies, was associated with an increased incidence of vertebral 545 deformities. On the other hand, IGF-II expression follows a different tendency within 546 each dietary treatment and also as compared to IGF-I expression. These differences may 547 support the idea that different hormonal signals and mechanisms of gene transcription 548 control the regulation of expression of both IGF forms (70,71).

549 Concluding, an increased dosage of AA in microdiets for sea bass larvae containing a 5% of DHA and 3000 mg/kg of  $\alpha$ -TOH was shown to be successful in 550 551 compensating, to some extent, the effect of lipid oxidation, thereby preventing the 552 appearance of muscular lesions, reducing cranial deformities and TBARS values, a 553 major indicator of oxidative stress. However no counteracting effect was found on AOE 554 expression, suggesting that other nutrients could be involved in enhancing the 555 antioxidant defenses at such levels. Moreover, a sparing effect between AA and  $\alpha$ -TOH 556 seems to occur in sea bass larvae. The implication of AA in regulating other antioxidant 557 components requires further investigation.

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565

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776	Figure legends:
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778	Figure 1: Longitudinal semithin (A) and transversal electro micrographs (B) of sea bass
779	larvae fed 5/300 diet. (A) Damaged muscle fibres showing breakage (arrow) as well as
780	darkening due to protein coagulation (*). (B) Affected fibre showing disarrangement of
781	the myofilaments (arrows), swollen sarcoplasmic reticulum (SR) and myelin figures (*).
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783	Figure 2: Skeletal deformities found in 35 dph larvae at the end of the experimental
784	period.
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786	Figure 3: Catalase (CAT), superoxide dismutase (SOD) and glutathione peroxidase
787	(GPX) expression levels measured by real-time PCR in Dicentrarchus labrax larvae fed
788	diets 1/150 ( $\bullet$ ), 5/300 ( $\blacksquare$ ) or 5/300+VitC ( $\blacktriangle$ ). mRNA copy number of each gene was
789	normalized as a ratio to 100 ng total RNA. Values are means, with standard deviations
790	represented by vertical bars. Mean values with unlike letters were significantly different
791	in gene expression among the treatments at a given sampling points.
792	
793	Figure 4: Insulin-like growth factors I and II (IGF-I and II) and myosin heavy chain
794	(MyHC) expression levels measured by real-time PCR in Dicentrarchus labrax larvae
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796	normalized as a ratio to 100 ng total RNA. Values are means, with standard deviations
797	represented by vertical bars. Mean values with unlike letters were significantly different
798	in gene expression among the treatments at a given sampling points.

Dietary DHA/vitamin E	1/1500	5/3000	5/3000+AA
Defatted squid powder *	69.00	68.85	68.32
$EPA^\dagger$	2.80	1.80	1.80
DHA $^{\dagger}$	0.20	6.70	6.70
Oleic acid <sup>‡</sup>	10.00	4.50	4.50
Soy lecithin <sup>•</sup>	2.00	2.00	2.00
Gelatin	3.00	3.00	3.00
Attractants <sup>‡</sup>	3.00	3.00	3.00
Taurin <sup>•</sup>	1.50	1.50	1.50
Vitamin premix <sup>§</sup>	6.00	6.00	6.00
Mineral premix <sup>**</sup>	2.50	2.50	2.50
Vitamin C <sup>††</sup>	-	-	0.53
Vitamin E <sup>‡‡</sup>	-	0.15	0.15

**Table 1.** Formulation (in % DW) of experimental diets containing two levels of DHA,

vitamin E and ascorbic acid and fed to sea bass larvae for 21 days.

DW, Dry weight; 1/1500, 1% DHA and 1500 g/kg vitamin E diet; 5/3000, 5% DHA and 3000 mg/kg vitamin E diet; 5/3000+AA, 5% DHA and 3000 mg/kg vitamin E diet with increased ascorbic acid content.

<sup>\*</sup> Riber and Son, Bergen, Norway.

<sup>†</sup>Croda, East Yorkshire, UK.

<sup>‡</sup>Merck, Darmstadt, Germany.

Acrofarma, Barcelona, Spain.

Panreac, Barcelona, Spain.

<sup>\*</sup>Attractants premix supplied per 100 g diet: Inosine-5-monophosphate, 500.0 mg; betaine, 660.0 mg; L-serine, 170.0 mg; L-phenilalanine, 250.0 mg; DL-alanine, 500.0 mg; L-sodium aspartate, 330.0 mg; L-valine, 250.0 mg; glycine, 170.0 mg. Sigma-Aldrich

<sup>§</sup> Vitamin premix supplied per 100 g diet: Cyanocobalamine, 0.030; Astaxanthin, 5.00; folic acid, 5.44; pyridoxine-HCI, 17.28; thiamine, 21.77; riboflavin, 72.53; Capantothenate, 101.59; p-aminobenzoic acid, 145.00; nicotinic acid, 290.16; *myo*-inositol, 1450.90; retinol acetate, 0.18; ergocalcipherol, 3.65; menadione, 17.28; α-tocopherol acetate, 150.00; ascorbyl monophosphate, 180.00.

\*\*Mineral premix supplied g per 100 g diet: NaCl, 215.133; MgSO<sub>4</sub>·7H<sub>2</sub>O, 677.545; NaH<sub>2</sub>PO<sub>4</sub>·H<sub>2</sub>O, 381.453; K<sub>2</sub>HPO<sub>4</sub>, 758.949; Ca(H<sub>2</sub>PO<sub>4</sub>)·2H<sub>2</sub>O, 671.610; FeC<sub>6</sub>H<sub>5</sub>O<sub>7</sub>, 146.884; C<sub>3</sub>H<sub>5</sub>O<sub>3</sub>·1/2Ca, 1617.210; Al<sub>2</sub>(SO<sub>4</sub>)<sub>3</sub>·6H<sub>2</sub>O, 0.693; ZnSO<sub>4</sub>·7H<sub>2</sub>O, 14.837; CuSO<sub>4</sub>·5H<sub>2</sub>O, 1.247; MnSO<sub>4</sub>·H<sub>2</sub>O, 2.998; KI, 0.742; CoSO<sub>4</sub>·7H<sub>2</sub>O, 10.706.

<sup>††</sup> Rovimix Stay-C 35, Roche, Paris, France.

<sup>‡‡</sup> DL-α-tocopherol acetate,Sigma-Aldrich, Madrid, Spain.

		Dist	
	1/1500	5/3000	5/3000+AA
14.0	1 54	1 26	0.78
14.0 $14.1n_7$	0.15	0.25	0.76
14.1n-7 1/1.1n-5	0.13	0.25	0.00
15.00	0.22	0.33	0.15
15:1n-5	0.02	0.43	0.15 n.d
16:0180	0.02	0.14	0.07
16:0	7.86	5 59	5.08
16:1n 7	7.80	2.59	2.01
10:1n-7	0.10	0.23	0.11
10.111-5 16:2n 6	0.19 n.d	0.23	0.11
16:2n 4	0.32	0.17	0.00
10.211-4	0.32	0.39	0.20
17.0 16.2m 2	0.08	0.82	0.00
10:311-3 16:4n 2	0.08	0.12	0.07
10:411-5	0.09	0.15	0.10
18:0	1.29	2.29	2.18
18:1n-9+n-/	55.70	31.12	30.33
18:1n-5	0.72	0.46	0.36
18:2n-9	0.25	0.13	0.02
18:2n-6	7.40	6.99	6.71
18:2n-4	0.46	0.28	0.23
18:3n-6	0.11	0.11	0.10
18:3n-4	0.13	0.10	0.09
18:3n-3	0.72	0.83	0.82
18:3n-1	n.d.	0.04	0.04
18:4n-3	0.83	0.94	0.98
18:4n-1	0.08	0.08	0.09
20:0	0.10	0.31	0.30
20:1n-9+n-7	1.10	1.53	1.50
20:1n-5	0.05	0.12	0.11
20:2n-9	0.05	0.04	0.04
20:2n-6	0.09	0.21	0.21
20:3n-6	0.09	0.13	0.14
20:4n-6	0.71	1.57	1.62
20:3n-3	0.07	0.18	0.18
20:4n-3	0.32	0.52	0.54
20:5n-3	8.66	11.04	12.08
22:1n-11	0.17	0.51	0.53
22:1n-9	0.08	0.25	0.24
22:4n-6	0.02	0.19	0.20
22:5n-6	0.19	1.75	1.86
22:5n-3	0.32	1.29	1.42
22:6n-3	4.58	24.55	27.54
SAFA	12.28	10.70	9.16
MUFA	61.99	37.23	35.34
n-3 PUFA	15.68	39.61	43.73
n-6 PUFA	8.61	11.14	10.90
n-9 PUFA	57.19	33.10	32.17
n-3 LC-PUFA	13.96	37.58	41.76
n-3/n-6 PUFA	1.82	3.56	4.01
EPA/DHA	1.89	0.45	0.44
ARA/DHA	0.15	0.06	0.06
ARA/EPA	0.08	0.14	0.13

**Table 2.** Main fatty acids (% total of fatty acids) of the experimental diets fed to European sea bass for three weeks.

1/1500, 1% DHA and 1500 g/kg vitamin E diet; 5/3000, 5% DHA and 3000 mg/kg vitamin E diet; 5/3000+AA, 5% DHA and 3000 mg/kg vitamin E diet with increased ascorbic acid content.

		Diets	
	1/1500	5/3000	5/3000+AA
Protein (% DW)	74.46±0.58	76.13±0.09	72.36±0.55
Ash (% DW)	5.01±0.12	5.38±0.14	5.39±0.14
Moisture (%)	10.31±0.46	$9.99 \pm 0.28$	9.48±0.11
Lipids (% DW)	14.98±0.31	15.80±0.02	$15.94{\pm}1.05$
α-tocopherol (mg/kg DW)	1410.12±38.77	3033.01±43.33	3179.72±75.69
Ascorbic acid ( mg/kg DW)	1495.88±5.54	1477.44±3.29	2998.56±7.46

**Table 3.-** Proximate composition,  $\alpha$ -tocopherol and ascorbic acid content (mean $\pm$  SD) in experimental diets fed to sea bass larvae for three weeks.

DW, Dry weight; 1/1500, 1% DHA and 1500 g/kg vitamin E diet; 5/3000, 5% DHA and 3000 mg/kg vitamin E diet; 5/3000+AA, 5% DHA and 3000 mg/kg vitamin E diet with increased ascorbic acid content.

**Table 4.-** Sea bass larvae performance and levels of lipid peroxidation products (TBARS) and vitamin E ( $\alpha$ -tocopherol) content in sea bass larvae at the beginning and after eating the experimental diets with two levels of DHA (1 and 5%) and  $\alpha$ -tocopherol (1500 or 3000  $\mu$ g/g) and supplemented or not with ascorbic acid (1800  $\mu$ g/g) for three weeks.

	Diets			
-	Initial	1/1500	5/3000	5/3000+AA
Results of dietary trial				
Larval total length (mm)	$8.58\pm0.64$	$12.60\pm0.93^{\rm a}$	$10.89 \pm 1.24^{\circ}$	$11.24\pm1.08^{\rm b}$
Larval dry weight (mg)	$0.36\pm0.00$	$1.33\pm0.46$	$0.94\pm0.05$	$1.01\pm0.07$
Survival (%)	-	$60.51 \pm 9.10$	$48.42 \pm 4.00$	$47.43 \pm 10.50$
Incidence of muscular lesions (%)	-	$17.50 \pm 14.14^{b}$	$52.63 \pm 15.93^{a}$	$20.70 \pm 15.62^{b}$
Incidence of skeletal malformations (%)	-	28.33±3.09	33.11±3.89	29.67±4.12
Ossification degree rate (%)	-	19.22±3.60 <sup>a</sup>	$15.24 \pm 4.18^{b}$	$15.07 \pm 4.60^{b}$
TBARS				
NMol MDA/g dry mass	62.85±0.61	166.62±25.08 <sup>c</sup>	2402.15±67.91 <sup>a</sup>	$846.87 \pm 94.74^{b}$
Vitamin F (a-tocopherol)				
mg/kg dry mass	111.45±43.26	$630.24{\pm}12.39^{b}$	542.10±80.51 <sup>b</sup>	$757.12 \pm 44.55^{a}$

DW, Dry weight; 1/1500, 1% DHA and 1500 g/kg vitamin E diet; 5/3000, 5% DHA and 3000 mg/kg vitamin E diet; 5/3000+AA, 5% DHA and 3000 mg/kg vitamin E diet with increased ascorbic acid content.

Data are means  $\pm$  SD. Values within the same row bearing different superscript letter are significantly different (*P*< 0.05)

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	1/1500	5/3000	5/3000+AA
14:0	0.93±0.06	0.76±0.03	0.51±0.28
14:1n-7	$0.08 \pm 0.01$	$0.08 \pm 0.03$	0.34±0.37
14:1n-5	$0.07 \pm 0.02$	$0.04 \pm 0.01$	0.34±0.43
15:0	0.63±0.29	0.98±1.22	0.38±0.26
15:1n-5	0.12±0.10	$0.09 \pm 0.06$	0.79±0.23
16:0ISO	n.d.	$0.11 \pm 0.01$	0.28±0.25
16:0	17.51±2.44	17.61±0.25	16.32±0.26
16:1n-7	2.02±0.14	$1.60\pm0.06$	1.63±0.16
16:1n-5	$0.25 \pm 0.01$	$0.25 \pm 0.01$	$0.40\pm0.18$
16:2n-6	$0.30\pm0.00$	$0.36 \pm 0.04$	0.46±0.15
16:2n-4	$0.93 \pm 0.40$	0.96±0.15	0.89±0.12
17:0	0.91±0.10	$0.80 \pm 0.03$	$0.81 \pm 0.08$
16:3n-3	$0.12 \pm 0.01$	0.14±0.03	0.09±0.03
16:3n-1	0.10±0.02	$0.54 \pm 0.10$	0.54±0.09
16:4n-3	$0.62 \pm 0.35$	$0.44 \pm 0.10$	$0.48\pm0.01$
16:4n-1	n.d.	$0.17 \pm 0.01$	0.16±0.00
18:0	11.66±3.41	12.29±0.36	$11.07 \pm 1.04$
18:1n-9	$26.35 \pm 4.87^{a}$	20.92±0.51 <sup>b</sup>	20.57±0.13 <sup>b</sup>
18:1n-7	4.85±0.29	4.44±0.39	3.99±0.23
18:1n-5	$0.62 \pm 0.24$	$0.49 \pm 0.07$	0.45±0.12
18:2n-9	n.d.	0.13±0.11	0.13±0.10
18:2n-6	4.23±0.08	3.90±0.17	3.85±0.14
18:2n-4	$0.04 \pm 0.05$	$0.06 \pm 0.01$	$0.07 \pm 0.00$
18:3n-6	0.43±0.01	0.38±0.04	0.33±0.01
18:3n-4	$0.07 \pm 0.04$	$0.06\pm0.02$	$0.06\pm0.02$
18:3n-3	$0.32 \pm 0.05$	$0.44 \pm 0.04$	$0.46\pm0.05$
18:4n-3	$0.29 \pm 0.06$	0.29±0.13	0.29±0.01
20:0	0.38±0.19	$0.47 \pm 0.01$	$0.48\pm0.04$
20:1n-9+n-7	1.83±0.0	1.77±0.06	$1.80\pm0.02$
20:1n-5	0.26±0.15	0.13±0.01	0.13±0.02
20:2n-6	0.50±0.13	$0.65 \pm 0.08$	$0.60\pm0.07$
20:3n-6	$0.08 \pm 0.01$	$0.08 \pm 0.00$	$0.08\pm0.01$
20:4n-6	$2.38 \pm 0.04^{b}$	$3.07 \pm 0.22^{a}$	3.13±0.24 <sup>a</sup>
20:3n-3	0.12±0.09	$0.15 \pm 0.04$	0.13±0.00
20:4n-3	$0.14 \pm 0.00$	$0.16 \pm 0.01$	0.17±0.01
20:5n-3	5.91±1.18	4.85±0.20	5.31±0.33
22:1n-11	$0.46 \pm 0.30$	$0.19 \pm 0.08$	0.26±0.02
22:1n-9	0.26±0.15	$0.26 \pm 0.06$	$0.27 \pm 0.05$
22:4n-6	n.d.	$0.11 \pm 0.02$	0.13±0.04
22:5n-6	$1.09 \pm 0.09$	$1.26 \pm 0.07$	1.33±0.05
22:5n-3	$0.64 \pm 0.26$	$0.57 \pm 0.06$	$0.64 \pm 0.08$
22:6n-3	12.79±0.37 <sup>b</sup>	$18.04{\pm}1.19^{a}$	$20.26 \pm 0.25^{a}$
Saturated	32.01±6.29	32.91±1.54	29.56±1.20
Monoenoics	36.73±4.51 <sup>a</sup>	$30.20 \pm 0.55^{b}$	30.56±1.44 <sup>b</sup>
n-3 PUFA	$20.94 \pm 2.23^{b}$	$25.11 \pm 1.43^{a}$	$27.83 \pm 0.57^{a}$
n-6 PUFA	8.45±0.97	9.80±0.47	9.91±0.09
n-9 PUFA	$28.44 \pm 4.72^{a}$	$23.08 \pm 0.39^{b}$	$22.76 \pm 0.05^{b}$
n-3 LC-PUFA	12.92±7.45 <sup>c</sup>	$23.78 \pm 1.42^{b}$	26.51±0.67 <sup>a</sup>
n-3/n-6 PUFA	2.51±0.55	$2.56 \pm 0.05$	2.81±0.03
EPA/DHA	$0.33 \pm 0.07^{b}$	$1.16\pm0.10^{a}$	$1.11\pm0.17^{a}$
ARA/DHA	$0.18 \pm 0.01^{b}$	$0.88{\pm}0.07^{a}$	$0.80{\pm}0.04^{a}$
ARA/EPA	$0.54{\pm}0.04^{b}$	2.56±0.05 <sup>a</sup>	3.00±0.33 <sup>a</sup>

**Table 5.** Main fatty acid composition of total lipids from sea bass larvae after three weeks of feeding the experimental diets (% total fatty acid).

1/1500, 1% DHA and 1500 g/kg vitamin E diet; 5/3000, 5% DHA and 3000 mg/kg vitamin E diet; 5/3000+AA, 5% DHA and 3000 mg/kg vitamin E diet with increased

ascorbic acid content. Each value represents mean±SD. Values within the same row bearing different superscript letter are significantly different (P<0.05); n.d., not detected



**Figure 1.-** Longitudinal semithin (A) and transversal electro micrographs (B) of sea bass larvae fed 5/3000 diet. (A) Damaged muscle fibres showing breakage (arrow) as well as darkening due to protein coagulation (\*). (B) Affected fibre showing disarrangement of the myofilaments (arrows), swollen sarcoplasmic reticulum (SR) and myelin figures (\*).



**Figure 2.-** Skeletal deformities found in 35 dph larvae at the end of the experimental period. Different superscript letters mean significant differences for the same type of deformity among treatments.



**Figure 3.-** Catalase (CAT), superoxide dismutase (SOD) and glutathione peroxidase (GPX) gene expression levels measured by real-time PCR in *Dicentrarchus labrax* 

larvae fed diets 1/1500 ( $\bullet$ ;1% DHA and 1500 g/kg vitamin E), 5/3000 ( $\bullet$ ; 5% DHA and 3000 mg/kg vitamin E) or 5/3000+VitC ( $\blacktriangle$ ; 5% DHA and 3000 mg/kg vitamin E diet with increased ascorbic acid content). mRNA copy number of each gene was normalized as a ratio to 100 ng total RNA. Values are means, with standard deviations represented by vertical bars. Mean values with unlike letters were significantly different in gene expression among the treatments at a given sampling points.



**Figure 4.-** Insulin-like growth factors I and II (IGF-I and II) and myosin heavy chain (MyHC) gene expression levels measured by real-time PCR in *Dicentrarchus labrax* larvae fed diets 1/1500 (•;1% DHA and 1500 g/kg vitamin E), 5/3000 (•; 5% DHA and 3000 mg/kg vitamin E) or 5/3000+VitC (•; 5% DHA and 3000 mg/kg vitamin E diet with increased ascorbic acid content). Values are means, with standard deviations represented by vertical bars. Mean values with unlike letters were significantly different in gene expression among the treatments at a given sampling points.