Effect of salinity on the biosynthesis of n-3 long-chain polyunsaturated fatty 1 acids in silverside Chirostoma estor. 2 3 Fonseca-Madrigal J¹., Pineda-Delgado D¹., Martínez-Palacios C¹., Rodríguez C²., 4 Tocher D.R³. 5 6 ¹ Laboratorio de Acuicultura, Instituto de Investigaciones Agropecuarias y Forestales, 7 8 UMSNH, Morelia, Michoacán, México. ² Departamento de Biología Animal (U.D.I. Fisiología Animal), Facultad de Biología, 9 Universidad de La Laguna, La Laguna, Tenerife, España. 10 ³ Institute of Aquaculture, University of Stirling, U.K. 11 12 Corresponding author: Jorge Fonseca-Madrigal, Laboratorio de Acuicultura, Instituto 13 de Investigaciones Agropecuarias y Forestales, UMSNH. Av. San Juanito Itzícuaro 14 S/N, Col. San Juanito Itzícuaro, C.P. 58330, Morelia, Michoacán, México. 15 fonseca.madrigal@gmail.com 16 17 18 Key Words: Chirostoma estor, Menidia estor, long-chain polyunsaturated fatty acids, 19 biosynthesis, desaturation, elongation, salinity 20 21 22

22 Abstract

23

The genus Chirostoma (silversides) belongs to the family Atherinopsidae, which 24 contains around 150 species, most of which are marine. However, Mexican silverside 25 (Chirostoma estor) is one of the few representatives of freshwater atherinopsids and 26 is only found in some lakes of the Mexican Central Plateau. However, studies have 27 shown that *C. estor* has improved survival, growth and development when cultured in 28 water conditions with increased salinity. In addition, *C. estor* displays an unusual fatty 29 acid composition for a freshwater fish with high docosahexaenoic acid (DHA) : 30 eicosapentaenoic acid (EPA) ratios. Freshwater and marine fish species display very 31 different essential fatty acid metabolism and requirements and so the present study 32 investigated long-chain polyunsaturated fatty acid (LC-PUFA) biosynthesis to 33 34 determine the capacity of C. estor for endogenous production of EPA and DHA, and the effect that salinity has on these pathways. Briefly, C. estor were maintained at 35 three salinities (0, 5 and 15 ppt) and the metabolism of ¹⁴C-labelled 18:3n-3 36 determined in isolated hepatocyte and enterocyte cells. The results showed that C. 37 estor has the capacity for endogenous biosynthesis of LC-PUFA from 18-carbon fatty 38 acid precursors, but that the pathway was essentially only active in saline conditions 39 with virtually no activity in cells isolated from fish grown in freshwater. The activity of 40 the LC-PUFA biosynthesis pathway was also higher in cells isolated from fish at 15 41 ppt compared to fish at 5 ppt. The pathway was around 5-fold higher in hepatocytes 42 compared to enterocytes, although the majority of 18:3n-3 was converted to 18:4n-3 43 and 20:4n-3 in hepatocytes whereas the proportions of 18:3n-3 converted to EPA 44 and DHA were higher in enterocytes. The data were consistent with the hypothesis 45

- that conversion of EPA to DHA could contribute, at least in part, to the generally high
- 47 DHA:EPA ratios observed in the tissue lipids of *C. estor*.

48 Introduction

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Mexican silverside (Chirostoma estor, also reported as Menidia estor) from lake 50 Pátzcuaro is one of the most valued freshwater fish in Mexico. The species, locally 51 known as "pez blanco", has greatly influenced the cultural environment and economy 52 of the native people of the region. The species is now endangered because of a 53 range of factors including over-fishing, environmental degradation of the lake and 54 introduction of exotic species (Martínez-Palacios et al. 2008). However, silverside is 55 a species with high potential for aquaculture commanding a good price in regional 56 markets (\$40-80 USD kg⁻¹) (Martínez-Palacios et al. 2008). Recently, there have 57 been efforts to preserve the species through aquaculture techniques (Martínez-58 Palacios et al. 2002, 2003, 2004, 2006, 2007). The genus Chirostoma (silversides) 59 belongs to the family Atherinopsidae, which contains around 150 species, most of 60 which are marine. However, C. estor is one of the few representatives of freshwater 61 atherinopsids and is only found in some freshwater lakes of the Mexican Central 62 Plateau. Therefore, although silverside is a freshwater species, it shares many 63 characteristics in common with marine Atherinopsids because of their common 64 ancestry (Barbour 1973). Thus, aquaculture of Mexican silverside C. estor involves 65 the transfer of the fish to different salinities over the whole cycle of production from 66 incubation of the eggs to juvenile development. Specifically, silverside has better 67 growth and survival when cultured in saline conditions (Martínez-Palacios et al. 68 2004). 69

C. estor is also considered to be a carnivorous species and so their essential fatty acid (EFA) requirements were expected to be more similar to that of a marine carnivorous species than that of a typical freshwater fish (Martínez-Palacios et al.

2008). It has been reported that most freshwater fish studied, in contrast to the 73 marine species studied, have the ability to elongate and desaturate 18-carbon 74 polyunsaturated fatty acids (PUFA) (18:2n-6, linoleic acid/LOA and 18:3n-3 α -75 linolenic acid/ALA) to long-chain PUFA (LC-PUFA) of 20 carbons (20:4n-6, 76 arachidonic acid/ARA and 20:5n-3, eicosapentanoic acid/EPA) and 22 carbons 77 (22:6n-3, docosahexaenoic acid/DHA) (Tocher 2010). Thus, it is generally assumed 78 that LOA and ALA can satisfy EFA requirements for freshwater species but ARA, 79 EPA and DHA are the required EFA for marine species (Sargent et al. 1995a). 80 However, the feeding habits of fish may also be determinants of precise EFA 81 requirements: carnivorous fish obtain the biologically active LC-PUFA directly from 82 their diet and consequently they now have only a low ability to desaturate and 83 elongate 18-carbon fatty acids whereas herbivorous fish have higher levels of C₁₈ 84 PUFA and lower LC-PUFA in their diet and so have retained the ability to convert C₁₈ 85 PUFA to LC-PUFA (Sargent et al. 1999). If the assertion that C. estor may have 86 characteristics of a marine species is correct, then ARA, EPA and DHA would need 87 to be included in the diet to satisfy their nutritional requirements. 88

There is scarce information of the lipid and fatty acid compositions and metabolism 89 in C. estor (Palacios et al. 2007). Wild fish contained high levels of DHA (20 - 32% of 90 total fatty acids) but surprisingly low levels of EPA (1 - 3%) in contrast with the fatty 91 acid profile found in samples of zooplankton, its natural diet (12% DHA, 13% EPA) 92 (Martínez-Palacios et al. 2003). There are two possible explanations for these 93 findings; firstly, that C. estor selectively accumulates DHA preferentially over other 94 fatty acids such as EPA depending on its own physiological requirements, or 95 secondly, that this species has the capacity to convert EPA and/or other n-3 series 96 fatty acids to DHA (Tocher 2003). The second explanation is also supported by the 97

presence of DHA in larvae fed rotifers with a low DHA / high ALA composition
(Martínez-Palacios et al. 2006).

Salinity has been shown to affect lipid and fatty acid composition in salmonids 100 although many effects occurred in advance of seawater transfer during parr-smolt 101 transformation (Bendiksen et al. 2003; Peng et al. 2003). It was also shown that the 102 activity of the LC-PUFA synthesis pathway was regulated by environmental cues in 103 Atlantic salmon (Salmo salar) and peaked around seawater transfer and was 104 considerably lower during the seawater phase (Bell et al. 1997; Tocher et al. 2000). 105 These changes in activity reflected changes in the expression of fatty acyl 106 desaturase genes in freshwater and seawater phases (Zheng et al. 2005). There are 107 also several studies reporting fatty acid compositions in fish reared at different 108 salinities (Cordier et al. 2002; Kheriji et al. 2003; Haliloglu et al. 2004; Martinez-109 110 Alvarez et al 2005; Dantagnan et al. 2007; Navarro et al. 2009; Xu et al. 2010; Hunt et al., 2011). Results are not consistent with increased associated with both reduced 111 (Ciordier et al. 2002; Kheriji et al. 2003) and increased (Xu et al. 2010; Hunt et al. 112 2011) levels of LC-PUFA including EPA and DHA. In contrast, the effects of salinity 113 on lipid and fatty acid biochemistry and metabolism have been little studied in non-114 salmonid fish although the modulation of $\Delta 6$ fatty acyl desaturase in teleosts was 115 recently reviewed, with the effects of salinity again being variable (Vagner and 116 Santigosa 2011). However, the expression of $\Delta 6$ desaturase was higher in liver of 117 rabbitfish (Siganus canaliculatus) and red sea bream (Pagrus major) reared at lower 118 salinity (10-15 ppt) compared to fish reared at higher salinity (32-33 ppt) (Li et al 119 2008; Sarker et al 2011). 120

121 The present study aims to investigate the two issues of salinity preference and 122 LC-PUFA metabolism in *C.estor* to determine if there is a relationship between them.

The basic hypothesis investigated was that LC-PUFA synthesis in C. estor will be 123 influenced by ambient salinity, and that the improved performance at increased 124 salinity will be related to this interaction between salinity and LC-PUFA biosynthesis. 125 The specific objectives were to determine the pathways and activities of LC-PUFA 126 biosynthesis in C. estor in order to elucidate potential mechanisms underpinning its 127 uncommon fatty acid profile and if the pathway reflects its marine ancestry, and to 128 determine if salinity affects the biosynthesis of LC-PUFA in a way that can explain 129 the apparent preference of C. estor for saline conditions (Martínez-Palacios et al. 130 2004). 131

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133 Materials and methods

134 Experimental fish

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Forty-five juvenile silverside (Chirostoma estor) of average initial weight around 50g 136 were obtained from a research production plant (UMSNH, Michoacan, Mexico). Fish 137 were maintained in glass-fiber tanks of 40cm high x 60cm diameter and 100L 138 capacity with constant aeration and temperature control (25± 0.4°C). All the 139 experimental units were maintained in a 12:12 dark: light photoperiod. All fish were 140 fed a standard commercial pellet feed (see Palacios et al. 2007), and every three 141 days tanks were siphoned and 30% of the water was renewed in order to maintain 142 high water quality with dissolved oxygen, nitrites, nitrates, pH, and total ammonia 143 monitored at 3-day intervals. The experimental design consisted of three salinity 144 treatments: freshwater (0 ppt; i.e < 0.05), 5 ppt and 15 ppt of salinity, each in 145 triplicate with 5 fish per tank (15 per treatment). Different salinities were obtained by 146 using artificial seawater (Instant Ocean Synthetic sea salt, Aquarium Systems) and 147

UV filtered ground water. Fish were fed a diet consisting of *Artemia franciscana* and
 a commercial feed (1:3) four times per day over 15 days prior to experimentation.

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151 *Preparation of isolated hepatocytes and enterocytes*

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With some modifications, the method for the preparation of isolated hepatocytes and 153 enterocytes established for salmonids was followed (Bell et al. 1997; Tocher et al. 154 1997, 2002). Briefly, six fish from each salinity treatment (two per tank) were 155 sacrificed with an overdose of benzocaine (50-60mg L⁻¹) to minimize stress (Ross et 156 al. 2007) and the livers and intestinal tracts dissected immediately. The livers and 157 intestines of two fish (i.e. per tank) were pooled for each sample so that there were 3 158 liver and 3 intestinal samples per treatment. The gall bladder was removed carefully 159 160 from the liver, the main blood vessels trimmed, and the liver perfused via the hepatic vein with solution A (calcium and magnesium-free Hanks balanced salt solution 161 (HBSS) containing 10 mM HEPES) to clear blood from the tissue. The liver was 162 chopped finely and about 0.5 g was taken and incubated with shaking in 20 ml of 163 solution A containing 0.1% (w/v) collagenase in a temperature controlled incubator at 164 25 °C for 45 min. Digested liver tissue was filtered through 100 µm nylon gauze and 165 the cells collected by centrifugation at 300 x g for 2 min. The cell pellet was washed 166 with 20 ml of solution A containing 1% w/v fatty acid-free bovine serum albumin 167 (FAF-BSA) and re-centrifuged. The washing was repeated with a further 20 ml of 168 solution A without FAF-BSA. The hepatocytes were resuspended in 10 ml of Medium 169 199 containing 10 mM HEPES. One hundred µl of cell suspension was mixed with 170 400 µl of the vital stain, Trypan Blue, and hepatocytes counted and viability assessed 171 using a haemocytometer. 172

With relatively minor modification, the above method was used to isolate 173 enterocyte-enriched preparations from *C. estor* intestine as described previously for 174 caecal enterocytes from salmon (Fonseca-Madrigal et al. 2006). Briefly, entire 175 intestinal tracts were dissected, cleaned of adhering adipose tissue, and luminal 176 contents rinsed away with solution A before being chopped finely and incubated with 177 0.1% (w/v) collagenase as above. The digested intestinal tissue was filtered through 178 100 µm nylon gauze and the cells collected, washed, resuspended in medium (as 179 above), and viability checked as for hepatocytes. The enriched enterocyte 180 preparation was predominantly enterocytes although some secretory cells were also 181 present. 182

¹⁸³ Viability of both isolated cell preparations was > 95% at isolation and decreased by ¹⁸⁴ less than 5% over the period of the incubation. One hundred μ l of the hepatocyte and ¹⁸⁵ enterocyte suspensions were retained for protein determination according to the ¹⁸⁶ method of Lowry et al. (1951) after incubation with 0.4 ml of 0.25% (w/v) SDS/1M ¹⁸⁷ NaOH for 45 min at 60 °C.

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189 Assay of hepatocyte and enterocyte fatty acyl desaturation/elongation activities

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Six ml of each hepatocyte or enterocyte suspension were dispensed into 25 cm² tissue culture flasks and incubated at 20°C for 2h with 0.3 μ Ci (~ 1 μ M) [1-¹⁴C]18:3n-3 or [1-¹⁴C]20:5n-3, added as complexes with FAF-BSA in phosphate buffered saline as described previously (Ghioni et al. 1997). After incubation, the cell suspensions were transferred to glass conical test tubes and centrifuged at 500 x g for 2 min. The supernatants were discarded and the cell pellets washed with 5 ml of ice-cold HBSS/FAF-BSA. The supernatant was carefully discarded and total lipid extracted

from the cell pellets using ice-cold chloroform/methanol (2:1, v/v) containing 0.01% 198 (w/v) BHT as described in detail previously (Tocher and Harvie 1988). Fatty acid 199 methyl esters (FAME) were prepared from total lipid by acid-catalyzed 200 transesterification using 2 ml of 1% H₂SO₄ in methanol plus 1 ml toluene as 201 described by Christie (1993), and FAME extracted and purified as described 202 previously (Tocher and Harvie 1988). The methyl esters were redissolved in 100 µl 203 isohexane containing 0.01% BHT and applied as 2.5 cm streaks to TLC plates 204 impregnated by spraying with 2 g silver nitrate in 20 ml acetonitrile and pre-activated 205 at 110°C for 30 min. Plates were fully developed in toluene/acetonitrile (95:5, v/v) 206 (Wilson and Sargent 1992) and autoradiography performed with Kodak MR2 film for 207 6 days at room temperature. Areas of silica containing individual PUFA were scraped 208 into scintillation mini-vials containing 2.5 ml of scintillation fluid (Ultima Gold, Perkin 209 210 Elmer, Monterrey, Mexico) and radioactivity determined in a scintillation ß-counter (Beckman LS Analyzer, Beckman Coulter de Mexico SA, Mexico City). 211

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213 Statistical analysis

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All the data are presented as means \pm SD (n = 3) and all statistical analyses were performed using S-Plus 2000 Professional Release 2 (MathSoft, Inc., Cambridge, MA, USA). The effects of salinity on LC-PUFA synthesis was analyzed by one-way ANOVA followed, where appropriate, by Tukey's post-test to determine significant differences between individual treatments (Zar 1999).

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221 Materials

[1-¹⁴C]18:3n-3 and [1-¹⁴C]20:5n-3 (50-55 mCi/mmol) were obtained from American
Radiolabeled Chemicals Inc. (St. Louis, MO, USA). HBSS, Medium 199, HEPES
buffer, collagenase (type IV), FAF-BSA, BHT, silver nitrate and all solvents (HPLC
grade) were obtained from Sigma Chemical Co. (St. Louis, MO, USA.). Thin-layer
chromatography (TLC) plates, precoated with silica gel 60 (without fluorescent
indicator) were obtained from Merck (Whitehouse Station, NJ, USA).

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229 Results

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231 Desaturation and elongation of ALA, [1-¹⁴C]18:3n-3

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Irrespective of tissue, activity of the LC-PUFA synthesis pathway from 18:3n-3 was 233 234 very low in fish maintained in freshwater. Increasing salinity resulted in significantly increased LC-PUFA synthesis in both hepatocytes and enterocytes as measured by 235 236 the recovery of radioactivity in the summed desaturated products (18:4, 20:4, 20:5, 22:5 and 22:6) of [1-¹⁴C]18:3n-3 (Fig.1). In both cell types, the rate of LC-PUFA 237 synthesis was highest in fish cultured at 15 ppt salinity, with rates of 0.41 ± 0.10 and 238 0.09 ± 0.04 pmol/h/mg protein in hepatocytes and enterocytes, respectively. These 239 values were 50- and 5-fold higher in hepatocytes and enterocytes, respectively, than 240 the activity observed in fish in freshwater. In both tissues, LC-PUFA synthesis at the 241 5 ppt salinity was intermediate between the activities in freshwater and 15 ppt salinity 242 with values of 0.13 \pm 0.01 and 0.06 \pm 0.01 pmol/h/mg protein in hepatoctes and 243 enterocytes, respectively. The LC-PUFA synthesis activity was 2.3- and 4.6-fold 244 higher in hepatocytes than in enterocytes at 5 and 15 ppt, respectively (Fig.1). The 245 rank order for recovery of radioactivity in desaturated products of 18:3n-3 was 18:4 246

>20:4 >22:6 >22:5 > 20:5 in hepatocytes (Fig. 2) whereas in enterocytes it was 20:4
>20:5 > 22:6 > 22:5 > 18:4 (Fig 3). In hepatocytes, recovery of radioactivity in DHA
exceeded that recovered in EPA, with the recovery of radioactivity in EPA and DHA
combined amounting to around 25% of the total radioactivity recovered (Fig. 2). In
contrast around 50% of total radioactivity recovered in enterocytes was as EPA and
DHA combined. Furthermore, in enterocytes, the recovery of radioactivity in EPA
increased, and that in DHA decreased, with increasing salinity (Fig. 3).

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255 Desaturation and elongation of EPA, [1-¹⁴C]20:5n-3

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As with [1-¹⁴C]18:3n-3, desaturation/elongation activity towards EPA in tissues from 257 fish maintained in freshwater was very low and increasing salinity significantly 258 259 increased desaturation/elongation activity in both hepatocytes and enterocytes as measured by the recovery of radioactivity in the summed products (22:5 and 22:6) of 260 [1-¹⁴C]20:5n-3 metabolism (Fig. 4). However, in contrast to LC-PUFA synthesis from 261 [1-¹⁴C]18:3n-3, the activity in hepatocytes was similar in fish at both 5 and 15 ppt 262 salinity with values of 0.37 ± 0.16 and 0.36 ± 0.15 pmol/h/mg protein, respectively. In 263 enterocytes, highest activity was obtained in fish at 5 ppt salinity, with a value of 0.13 264 265 \pm 0.01 pmol/h/mg protein compared to 0.05 \pm 0.02 in fish reared at 15 ppt (Fig. 4). Similar to the data obtained with [1-¹⁴C]18:3n-3, the LC-PUFA synthesis activity from 266 [1-¹⁴C]20:5n-3 was 2.9- and 7.0-fold higher in hepatocytes than in enterocytes from 267 fish reared at 5 ppt and 15 ppt, respectively (Fig.4). There was also a significant 268 difference in the products of [1-¹⁴C]20:5n-3 metabolism between hepatocytes and 269 enterocytes irrespective of treatment. The rank order for recovery of radioactivity in 270 products of 20:5n-3 metabolism was 22:6 >22:5 in hepatocytes, with the recovery of 271

radioactivity in DHA increasing with salinity with percentages of 57%, 65% and 78%
at 0, 5 and 15 ppt, respectively (Fig. 5). In enterocytes, the recovery of radioactivity in
274 22:5 exceeded the recovery in 22:6, with approximately 25%, 10% and 37% of
radioactivity recovered in DHA at 0, 5 and 15 ppt salinity, respectively (Fig 6).

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277 Discussion

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The primary objectives of the present work were to establish the extent and activity of 279 the LC-PUFA synthesis pathway in enterocytes and hepatocytes of C. estor and, 280 furthermore, to determine whether these activities were influenced by salinity. The 281 results demonstrated that both hepatocytes and enterocytes of C. estor displayed 282 physiologically relevant activities of LC-PUFA synthesis from ALA, particularly in 283 284 saline conditions. Thus, the values in hepatocytes and enterocytes from C. estor at 15 ppt salinity (0.41 and 0.09 pmol/h/mg protein, respectively) were lower than those 285 obtained in Atlantic salmon hepatocytes and enterocytes (0.9 and 1.2 pmol.h/mg 286 protein, respectively) (Zheng et al 2005), but higher than those obtained in 287 hepatocytes from the marine teleost Atlantic cod (Gadus morhua) (0.02 pmol/h/mg 288 protein) and similar to values from cod enterocytes (0.15 pmol/h/mg protein) (Tocher 289 et al. 2006). In addition, interest in the LC-PUFA synthesis pathway in C. estor is 290 partly due in its tissue fatty acid composition that, unusually for a freshwater species, 291 shows a very high DHA:EPA ratio (Martínez-Palacios et al. 2006). In marine and 292 freshwater fish tissue DHA:EPA ratios are most commonly in the range of 1:1 to 2:1 293 although ratios lower than this are also found in some species, particularly in 294 Southern oceans (Sargent et al. 1989). In contrast, C. estor has a fatty acid profile 295 with a DHA:EPA ratio that can vary from 10:1 to 20:1 (Martínez-Palacios et al. 2006). 296

This is generally unusual, even in marine fish, and very uncommon in freshwater 297 species (Ackman 1980). A few marine species show high DHA:EPA ratios, most 298 notable tuna species that can display ratios between 4 and 11, depending upon 299 tissue and species (Tocher 2003). In tuna the high tissue DHA:EPA ratios appear to 300 be due to generally higher DHA levels combined with relatively low EPA levels 301 (Tocher 2003). However, in *C. estor*, it appears that the main cause of the high 302 DHA:EPA ratios is the latter factor, that is, relatively low EPA levels in tissues, rather 303 than exceptionally high DHA (Martínez-Palacios et al. 2006). The present study has 304 confirmed that enterocytes and, especially, hepatocytes of C. estor demonstrated 305 significant DHA synthesis from EPA, particularly in saline conditions. Therefore, the 306 data are consistent with the hypothesis that conversion of EPA to DHA, particularly in 307 the liver, but also in the intestine, could contribute, at least in part, to the generally 308 309 high DHA:EPA ratios observed in the tissue lipids of C. estor (Martínez-Palacios et al. 2006). 310

Previously, intestine and pyloric caeca were shown to be tissue sites of substantial 311 LC-PUFA biosynthesis in salmonids (Atlantic salmon and trout) (Fonseca-Madrigal et 312 al. 2005, 2006), and this is why the capacity of enterocytes in C. estor for LC-PUFA 313 production was also investigated in the present study. The present study has 314 demonstrated that intestine in *C. estor* had the capability for LC-PUFA biosynthesis 315 but at significantly lower level than liver. The lower capacity of intestine for LC-PUFA 316 biosynthesis in *C. estor* in comparison to salmonids could be related to the feeding 317 habits of the species as C. estor is a zooplanktivorous fish with a short intestinal tract 318 and agastric digestive system (1:0.7 size of fish:size of intestine) with no pyloric 319 caeca (Martínez-Palacios et al. 2006). This is, of course, completely different to the 320 digestive tract of carnivorous species such as salmonids, which have a considerably 321

longer digestive system including a stomach and multiple caeca (Olsen and Ringø 322 1997). The results therefore suggest that the enterocytes in the much smaller 323 intestinal tract in C. estor and planktonivorous fish in general may not express the 324 same range of activities for processing absorbed nutrients as carnivorous fish 325 species, and may be focused more on the digestive and absorption roles. For 326 example, the activity of the LC-PUFA biosynthesis pathway is an order of magnitude 327 lower in enterocytes from C. estor compared to enterocytes from Atlantic salmon 328 (Zheng et al. 2005). Hepatocytes from C. estor showed much higher LC-PUFA 329 synthesis activity, which is expected due to the liver generally being the most 330 important organ in fatty acid and lipid metabolism in most fish species (Henderson 331 1996; Grum et al. 2002; Tocher 2003; Fonseca-Madrigal et al. 2005, 2006). 332

Studies on the development of C. estor aquaculture showed that this species 333 334 displays improved survival, growth and development when cultured in water conditions with increased salinity (Martinez-Palacios et al. 2004). Generally, egg 335 fertilization and incubation as well as many physiological processes including lipid 336 metabolism are dependent on, or influenced by, salinity (Bœuf and Payan 2001). For 337 example, changes in the fatty acid composition of tissue lipids associated with 338 changes in salinity have been reported previously in a number of fish species 339 including guppy (*Poecilia reticulata*) (Daikoku et al. 1982), milkfish (*Chanos chanos*) 340 (Borlogan and Benítez 1992) and turbot (Psetta maxima) (Tocher et al. 1994, 1995). 341 These adaptations in response to salinity include altered proportions of total 342 phospholipids and individual phospholipid classes, as well as changes in fatty acid 343 composition including levels of LC-PUFA and n-3/n-6 PUFA ratio. However, the data 344 are variable depending upon species and whether low or high salinity is the actual 345 challenge for that species. For instance, in marine fish, reduced salinity increased 346

percentages of DHA and ARA in mullet (*Mugil cephalus*) (Kheriji et al. 2003), but
reduced proportions of EPA and DHA in Japanese sea bass (*Lateolabrax japonicus*)
and European sea bass (*Dicentrarchus labrax*) (Xu et al. 2010; Hunt et al. 2011).

In the present study, there was a clear relationship between the salinity of the water 350 the fish were maintained and synthesis of LC-PUFA, independent of cell type, with 351 higher LC-PUFA synthesis activity in cells of fish cultured in water with higher salinity 352 compared to fish cultured in freshwater. However, it is perhaps more appropriate and 353 noteworthy to highlight the fact that the activity of the LC-PUFA synthesis pathway 354 was very low in freshwater. This was actually the most unusual feature of the 355 356 pathway in C. estor, rather the activities observed at higher salinity. Clearly, there was very little activity in freshwater and this was largely unprecedented as all the 357 freshwater fish species examined to date have generally shown appreciable LC-358 359 PUFA synthesis activity (Tocher 2010), such that 18:3n-3 and/or 18:2n-6 can satisfy their essential fatty acid requirements (NRC 2011). 360

The adaptation processes in response to a saline environment are primarily a series 361 of physiological changes involved osmoregulation, the regulation of ion balances 362 between the external medium and the corporal fluids (Morgan 1997; Laiz-Carrión et 363 al. 2004). Many of adaptations depend upon membrane processes and so changes 364 in lipid and, especially, fatty acid metabolism can be linked to the capacity of the fish 365 to adapt to salinity through changes in lipid and fatty acid compositions of 366 membranes that, in turn, affect membrane-associated proteins (receptors, enzymes 367 etc). Therefore, the influence of salinity on LC-PUFA production may be related to 368 the osmoregulatory response required for adaptation to higher salinity. The effects of 369 salinity on fatty acid compositions have been investigated (Tocher et al. 1994, 1995) 370 and the effects of salinity on LC-PUFA synthesis in hepatocytes have been indirectly 371

investigated in studies on the process of smoltification in Atlantic salmon (Bell et al. 372 1997; Tocher et al. 2000, 2002). In a trial investigating LC-PUFA synthesis in both 373 hepatocytes and enterocytes in farmed salmon, a peak of LC-PUFA production 374 occurred around the time the fish were transferred from freshwater to seawater, with 375 synthetic activity declining rapidly in the seawater phase to minimum levels (Tocher 376 et al. 2002). Although the effects of salinity on lipid and fatty acid biochemistry and 377 metabolism have been little studied in non-salmonid fish, the expression of $\Delta 6$ fatty 378 acyl desaturase in liver of the marine teleosts, rabbitfish and red sea bream, was 379 higher in fish maintained at low salinity compared to fish reared at high salinity (Li et 380 al 2008; Sarker et al 2011). This association between salinity and LC-PUFA 381 biosynthesis observed in fish was one of the factors underpinning the hypothesis 382 tested in the present study and the specific objectives were developed in this context. 383 384 However, the precise links between salinity changes, and LC-PUFA synthesis fatty acid composition in fish including C. estor require further investigation. 385

Irrespective of the precise mechanistic links, the results presented, showing very low 386 levels of activity in fish reared in freshwater and increased capacity for LC-PUFA 387 synthesis essentially in hepatocytes as salinity increased may be related with the fact 388 that this species displays better growth performance and development when cultured 389 in saline water. At the most simplistic level, increased capability for endogenous 390 synthesis of the biologically and physiologically essential LC-PUFA would be 391 potentially beneficial to the fish in comparison to the situation in freshwater where the 392 pathways appear almost totally suppressed. Therefore, it is tempting to speculate 393 that the differing activity of the LC-PUFA synthesis pathway is an underpinning factor 394 on the effect of salinity on growth performance of *C. estor*. However, it is not so clear 395 how the effect of salinity on the activity of LC-PUFA synthesis pathway in C. estor 396

relates to current knowledge of LC-PUFA and environmental salinity or to discuss of 397 the possible marine origin of this species. Marine species generally have a reduced 398 ability to produce LC-PUFA compared to freshwater species (Tocher 2010). This has 399 been explained as a possible evolutionary adaptation to the generally higher levels of 400 DHA in the marine environment (Sargent et al. 1995b), and so marine species have 401 had less evolutionary pressure to retain the ability to endogenously produce LC-402 PUFA; in contrast, freshwater food webs are generally characterized by lower levels 403 of DHA (Sargent et al. 1995b) and so evolutionary pressure for endogenous 404 production of LC-PUFA has been retained in freshwater species (Tocher 2010). 405 406 Therefore, this hypothesis would suggests it would be more advantageous for C. estor to have higher LC-PUFA biosynthesis in freshwater where the supply of EPA 407 and, especially, DHA would likely be lower. 408

409 Although the data obtained to date with over 30 species, generally support this hypothesis linking LC-PUFA levels and, especially, DHA levels in the different food 410 411 webs to evolutionary pressure for endogenous production of LC-PUFA, there are several potential confounding factors including precise feeding habit of different 412 species (herbivorous vs. carnivorous/piscivorous) as well as phylogenetic issues. For 413 instance, defining fish species simply as marine or freshwater is often not ideal 414 considering the large number of euryhaline and diadromous species. Furthermore, 415 the effect of feeding habit can also be generalized with the ability for endogenously 416 LC-PUFA biosynthesis being retained in herbivorous fish, but not in omnivorous, 417 carnivorous, or piscivorous fish. As alluded to above, most fish species studied to 418 date could fit either of these generalizations (environment or feeding habit). However, 419 recent studies have contributed directly to this debate. A feeding study with rabbitfish, 420 Siganus canaliculatus, which consumes benthic algae and seagrasses and is thus a 421

422 rare example of a herbivorous marine fish, suggested that it was able to 423 biosynthesize EPA and DHA (Li et al. 2008). Very recently, it was shown that 424 rabbitfish possess all the fatty acyl desaturase activities required for endogenous 425 synthesis of LC-PUFA (Li et al. 2010). These data suggest that trophic level and/or 426 feeding habit are indeed important factors associated with or determining a species' 427 ability for endogenous LC-PUFA synthesis.

The genus Chirostoma (silversides also known as Menidia) belongs to the family 428 Atherinopsidae, which contains around 150 species, most of which are marine. Thus, 429 C. estor is among the few representatives of totally freshwater atherinopsids and, 430 although it is only found in some lakes of the Mexican Central Plateau, it shares 431 common ancestry with marine Atherinopsids (Barbour 1973). However, the 432 evolutionary pathway for C. estor is not entirely clear (Barbour 1973), although 433 434 relationships determined by classical phylogeny can give clues to evolutionary history (Nelson 2006). Significant advances in determining the molecular mechanisms of LC-435 436 PUFA biosynthesis in fish have been made in the last decade with the cloning and functional characterization of fatty acyl desaturases and elongases from many fish 437 including freshwater, diadromous and marine species (Tocher 2010). Phylogenetic 438 analyses of the desaturase and elongase sequences have revealed some insights 439 into the possible evolutionary history of LC-PUFA biosynthesis in fish species 440 (Hastings et al. 2001; Zheng et al. 2004, 2009; Morais et al. 2009). The phylogenetic 441 sequence analysis generally reflected classical phylogeny, and grouped fish 442 desaturases in three distinct clusters (Leaver et al 2008). The Ostariophysi (common 443 carp and zebrafish), the Salmoniformes (trout and salmon), and the Acanthopterygia 444 (tilapia, sea bream, turbot, stickleback and medaka), with the 445 cod (Paracanthopterygii) branching from the Acanthopterygia line. However, many 446

questions still remain (Leaver et al. 2008; Li et al. 2010; Monroig et al. 2010) and *C. estor* represent an interesting species to study in this respect. The interesting
ancestry, pattern of LC-PUFA biosynthesis activity and the effects of salinity, which
conflicts with the existing paradigm, make *C. estor* a choice candidate for molecular
studies with the isolation, cloning and characterization of fatty acyl desaturases and
elongases being important goals for future studies.

The results of the present study have provided data that contribute to our 453 understanding of the unusual fatty acid profile found in tissues of *C. estor* indicating 454 that it could be explained, at least partly, by endogenous metabolic activity resulting 455 in elongation and desaturation of EPA to DHA. Moreover, with respect to the well-456 known beneficial effect of n-3 LC-PUFA on human health, it is noteworthy that the 457 results demonstrate that the increased ambient salinity used as part of the 458 459 management of this species in aquaculture farming should positively affect the nutritional quality of the flesh in terms of fatty acid composition. However, a complete 460 understanding of fatty acid metabolism in C. estor requires further more extensive 461 analysis to determine the potential roles of selective β -oxidation, acylation and 462 incorporation of fatty acids into lipid classes, and lipid and fatty acid transport 463 between tissues. 464

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670 Legends to Figures:

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FIG. 1. Total LC-PUFA biosynthesis (desaturation/elongation) activity in hepatocytes and enterocytes of *C. estor* cultured at different salinities. Results are means \pm S.D. (n= 3) and represent the rate of conversion (pmol.h⁻¹.mg protein⁻¹) of [1-¹⁴C]18:3n-3 to all desaturated products (sum of radioactivity recovered as 18:4n-3, 20:4n-3, 20:5n-3, 22:5n-3 and 22:6n-3). Columns assigned to a specific cell type with different superscript letters are significantly different as determined by one-way ANOVA followed by the Tukey post test (P < 0.05).

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FIG. 2. Individual fatty acid products of the desaturation and elongation of [1-¹⁴C]18:3n-3 in *C. estor* hepatocytes. Results are means \pm S.D. (n = 3) and represent the rate of production (pmol.h⁻¹.mg protein⁻¹) of individual fatty acids as determined by the recovery of radioactivity in each fatty acid fraction. Columns referring to a specific fatty acid having different superscript letters are significantly different as determined by one-way ANOVA followed by the Tukey post test (P < 0.05).

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FIG. 3. Individual fatty acid products of the desaturation and elongation of [1-¹⁴C]18:3n-3 in *C. estor* enterocytes. Results are means \pm S.D. (n = 3) and represent the rate of production (pmol.h⁻¹.mg protein⁻¹) of individual fatty acids as determined by the recovery of radioactivity in each fatty acid fraction. Columns referring to a specific fatty acid having different superscript letters are significantly different as determined by one-way ANOVA followed by the Tukey post test (P < 0.05).

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FIG. 4. Production of desaturation/elongation products from labeled EPA in hepatocytes and enterocytes of *C. estor* cultured at different salinities. Results are means \pm S.D. (n= 3) and represent the rate of conversion (pmol.h⁻¹.mg protein⁻¹) of [1-¹⁴C] 20:5n-3 to metabolised products (sum of radioactivity recovered as 22:5n-3 and 22:6n-3). Columns representing a specific cell type with different superscript letters are significantly different as determined by one-way ANOVA followed by the Tukey post test (P < 0.05).

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FIG. 5. Individual fatty acid products of the desaturation and elongation of $[1-^{14}C]$ 20:5n-3 in *C. estor* hepatocytes. Results are means ± S.D. (n = 3) and represent the rate of production (pmol.h⁻¹.mg protein⁻¹) of individual fatty acids as determined by the recovery of radioactivity in each fatty acid fraction. Columns referring to a specific fatty acid having different superscript letters are significantly different as determined by one-way ANOVA followed by the Tukey post test (P < 0.05).

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FIG. 6. Individual fatty acid products of the desaturation and elongation of $[1-^{14}C]$ 20:5n-3 in *C. estor* enterocytes. Results are means ± S.D. (n = 3) and represent the rate of production (pmol.h⁻¹.mg protein⁻¹) of individual fatty acids as determined by the recovery of radioactivity in each fatty acid fraction. Columns referring to a specific fatty acid having different superscript letters are significantly different as determined by one-way ANOVA followed by the Tukey post test (P < 0.05).

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