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Morini, M.; Peñaranda, D.; Vilchez Olivencia, MC.; Tveiten, H.; Lafont, A.; Dufour, S.; Pérez Igualada, LM.... (2017). The expression of nuclear and membrane estrogen receptors in the European eel throughout spermatogenesis. *Comparative Biochemistry and Physiology Part A Molecular & Integrative Physiology*. 203:91-99. doi:10.1016/j.cbpa.2016.08.020



The final publication is available at

<https://doi.org/10.1016/j.cbpa.2016.08.020>

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The expression of nuclear and membrane estrogen receptors in the European eel throughout spermatogenesis

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28 **Abstract**

29 Estradiol (E₂) can bind to nuclear estrogen receptors (ESR) or membrane estrogen
30 receptors (GPER). While mammals possess two nuclear ESRs and one membrane
31 GPER, the European eel, like most other teleosts, has three nuclear ESRs and two
32 membrane GPERs, as the result of a teleost specific genome duplication. In the current
33 study, the expression of the three nuclear ESRs (ESR1, ESR2a and ESR2b) and the two
34 membrane GPERs (GPERa and GPERb) in the brain-pituitary-gonad (BPG) axis of the
35 European eel was measured, throughout spermatogenesis.

36 The eels were first transferred from freshwater (FW) to seawater (SW), inducing
37 parallel increases in E₂ plasma levels and the expression of ESRs. This indicates that
38 salinity has a stimulatory effect on the E₂ signalling pathway along the BPG axis.

39 Stimulation of sexual maturation by weekly injections of human chorionic gonadotropin
40 (hCG) induced a progressive decrease in E₂ plasma levels, and different patterns of
41 expression of ESRs and GPERs in the BPG axis. The expression of nuclear ESRs
42 increased in some parts of the brain, suggesting a possible upregulation due to a local
43 production of E₂. In the testis, the highest expression levels of the nuclear ESRs were
44 observed at the beginning of spermatogenesis, possibly mediating the role of E₂ as
45 spermatogonia renewal factor, followed by a sharply decrease in the expression of
46 ESRs. Conversely, there was a marked increase observed in the expression of both
47 membrane GPERs throughout spermatogenesis, suggesting they play a major role in the
48 final stages of spermatogenesis.

49

50

51 **Keywords:** ESR, GPER, *Anguilla*, teleost, reproduction

52

53 **1. Introduction**

54 In male vertebrates, sex steroids, androgens, estrogens, and progestins, play significant
55 roles in the control of spermatogenesis (Schultz and Miura, 2002), process in which
56 diploid spermatogonia differentiate to mature haploid spermatozoa. Estrogens have
57 been shown to be indispensable for the early spermatogenic cycle, controlling the
58 spermatogonial stem cell renewal through its receptor (Miura et al., 1999, Miura and
59 Miura, 2011). Estradiol (E_2), as all sex steroids is a small lipophilic hormone, which can
60 diffuse through the cell membrane (Oren et al., 2004). E_2 can bind to intracellular
61 nuclear estrogen receptors (ESRs) and modulates gene transcription (Mangelsdorf et al.,
62 1995), which corresponds to the classic genomic mechanism of steroid action. Two
63 nuclear ESRs, ESR1 and ESR2 (also named $ER\alpha$ or NR3A1, and $ER\beta$ or NR3A2,
64 respectively), are present in mammals. They belong to the nuclear steroid receptor
65 superfamily, as well as androgen, progestin, gluco- and mineralocorticoid receptors
66 (Carson-Jurica et al., 1990; Laudet et al., 1992). Teleost species have at least three
67 distinct ESR subtypes, including ESR1, ESR2a and ESR2b (Hawkins et al., 2000; Ma et
68 al., 2000; Menuet et al., 2002), with ESR2a (also named $ER\beta_2$) and ESR2b (also named
69 $ER\beta_1$) resulting from the third whole genome duplication (3R) event that occurred in
70 teleost lineage (Hawkins et al., 2000; Lafont et al., in press).

71 In addition to the classic genomic functions, E_2 can bind itself to membrane receptors,
72 which activates intracellular signalling pathways through a fast, non-genomic action
73 (for review see: Thomas et al., 2012, or Nelson and Habibi, 2013). In mammals, the
74 former orphan receptor GPR30 was characterized as an E_2 membrane receptor, and is
75 also called G-protein coupled estrogen receptor GPER (Filardo and Thomas, 2005;
76 Filardo et al., 2007; for review see Prossnitz and Maggiolini, 2009). Two membrane
77 GPERs have recently been observed in most teleosts including the eel, likely resulting

78 from teleost 3R (Lafont et al., in press).

79 The European eel (*Anguilla anguilla*) has a complex catadromous life cycle which
80 includes a 5000-6000 km oceanic reproductive migration to reach its spawning site in
81 an unknown area of the Sargasso sea. Eels are euryhaline fish which are subjected to
82 high variations in salinity during their life cycle (Daverat et al., 2006). After their
83 juvenile growth period in continental waters, eels change from yellow eels to
84 prepubertal silver eels, future genitors that will undergo the transoceanic reproductive
85 migration. In captivity, the reproductive cycle is still not closed, and long-term
86 hormonal treatments (fish pituitary extracts for females, and human chorionic
87 gonadotropin, hCG, for males) are required to induce sexual maturation in silver eels
88 (Boëtius and Boëtius, 1967; Pérez et al., 2000; Asturiano et al., 2006; Gallego et al.,
89 2012). This, together with the dramatic reduction in the wild European eel population
90 (ICES, 2012) has increased the interest in deciphering the basic mechanisms controlling
91 the reproduction of this species. Furthermore, the phylogenetical position of the
92 European eel, branching at the base of teleosts, may provide insights into ancestral
93 regulatory functions in teleosts, the largest group of vertebrates (Henkel et al., 2012a,b).

94 As far as we know, this is the first study on male teleosts to look at the expression of the
95 three nuclear (ESR1, ESR2a and ESR2b) and two membrane (GPERa, GPERb)
96 estrogen receptors in the BPG axis throughout the spermatogenetic process.

97

98 **2. Material and methods**

99 **2.1. Fish maintenance, hormonal treatments and sampling**

100 Eighty male European eels (mean body weight 100 ± 6 g) were purchased from the fish
101 farm Valenciana de Acuicultura, S.A. (Puzol, Valencia, Spain) and transferred to the
102 Aquaculture Laboratory in the Polytechnic University of Valencia. The 80 males were

103 randomly distributed and kept at 20 °C in two freshwater 200-L aquaria equipped with
104 separated recirculation systems, thermostats/coolers, and covered to maintain constant
105 darkness.

106 One group of 8 eels was anaesthetized with benzocaine (60 ppm) and sacrificed by
107 decapitation in freshwater (FW). The rest of the fish were gradually acclimatized over
108 the course of one week to seawater (37±0.3‰ of salinity). Groups of 8 eels were
109 anaesthetized and sacrificed by decapitation in seawater conditions (SW). Once a week
110 for 8 weeks the rest of the fish were anesthetized, weighed and injected with hCG (1.5
111 IU g⁻¹ fish; Profasi, Serono, Italy), to induce the spermatogenesis as previously
112 described by Pérez et al. (2000). Groups of 8 eels were anaesthetized and sacrificed by
113 decapitation each week (W1-8) through the hormonal treatment. For the analysis of
114 ESR expression through the spermatogenesis, the 8 latter groups have been redistributed
115 to 4 groups based on their spermatogenic stage.

116 Total body weight and testis weight were recorded to calculate the gonadosomatic index
117 [GSI = (gonad weight/total body weight)*100]. Blood samples were collected,
118 centrifuged and stored at -20 °C until E₂ plasma level analysis. Testicular tissue samples
119 were fixed in 10% formalin buffered at pH 7.4 for histological analysis.

120 Samples of anterior brain (dissected into three parts: olfactory bulbs, telencephalon,
121 mes-/di-encephalon), pituitary and testis were stored in 0.5 ml of RNAlater (Ambion
122 Inc., Huntingdon, UK) at -20 °C until extraction of total RNA.

123 Because eels stop feeding at the silver stage and throughout sexual maturation the fish
124 were not fed throughout the experiment. They were handled in accordance with the
125 European Union regulations concerning the protection of experimental animals (Dir
126 86/609/EEC).

127

128 **2.2. Gonadal histology**

129 The formalin-fixed mid-part testis samples were dehydrated in ethanol, embedded in
130 paraffin, sectioned to 5-10 μm thickness with a Shandon Hypercut manual microtome
131 (Shandon, Southern Products Ltd., England), and stained using the haematoxylin and
132 eosin method of National Diagnostic
133 (www.nationaldiagnostics.com/histology/article/staining-procedures). Five slides per
134 fish were observed with a Nikon Eclipse E-400 microscope, and pictures were taken
135 with a Nikon DS-5M camera attached to the microscope (Nikon, Tokyo, Japan). The
136 stages of spermatogenesis were determined according to the germ cell types present in
137 the testis (Miura and Miura, 2001; Leal et al., 2009) their relative abundance, the degree
138 of development of the seminal tubules and the sperm production of the male at the time
139 of sacrifice (Morini et al., submitted). The stages considered were: Stage SPGA:
140 dominance of A spermatogonia, B spermatogonia present in low numbers; Stage
141 SPGB/SPC: dominance of B spermatogonia and spermatocytes, in some cases low
142 numbers of spermatids; Stage SD: dominance of spermatids, in some cases a small
143 number of spermatozoa; Stage SZ: dominance of spermatozoa (Fig. 1).

144

145 **2.3. Extraction and Reverse-Transcription**

146 Total RNA of the testis, anterior brain parts and pituitary were isolated using a Trizol
147 reagent (Life Technologies, Inc, Carlsbad, CA) as described by Peñaranda et al. (2013).
148 RNA concentration was evaluated using a NanoDrop 2000C Spectrophotometer (Fisher
149 Scientific SL, Spain). The testis RNA was treated using a DNase I of NucleoSpin RNA
150 XS kit (Macherey-Nagel, Düren, Germany). Twenty μl cDNA were synthesized from
151 500 ng of testis total RNA, using a qScript cDNA Synthesis Kit (Quanta Bioscience,
152 MD, USA). The brain parts and pituitary RNAs were treated using a DNase (gDNA

153 Wipeout Buffer, Qiagen, Hilden, Germany). Using a Quantiscript Reverse Transcriptase
154 (Qiagen, Hilden, Germany), 20 µl cDNA was synthesized from 500 ng of total RNA in
155 the case of the olfactory bulb and pituitary, and from 1 µg in the case of the
156 telencephalon and the mes-/diencephalon.

157

158 **2.4. Gene expression analyses by quantitative real-time PCR**

159 The quantitative real-time Polymerase Chain Reactions (qPCR) were carried out using
160 specific qPCR primers for each European eel estrogen nuclear and membrane receptor
161 (Lafont et al., in press) and the Acidic ribosomal phosphoprotein P0 (ARP) (Weltzien et
162 al., 2005) was used as the reference gene (Table I).

163

164 **2.4.1. Reference gene**

165 The stability of the reference gene was determined using the BestKeeper program
166 (Pfaffl et al., 2004), reporting a standard deviation (SD[±Cq]) lower than 1. The
167 BestKeeper calculated that variations in the reference gene are based on the arithmetic
168 mean of the Cq values. Genes with a SD value higher than 1 are defined as unstable. In
169 the testis: SD= 0.83; p<0.05 with a Cq geometric mean of 24.21±1.77; in the brain and
170 pituitary, olfactory bulb: SD= 0.81; telencephalon: SD= 0.48; mes-/diencephalon: SD=
171 0.58, pituitary: SD= 0.63; p<0.05 and the Cq geometric mean of the olfactory bulb:
172 23.39±1.76; telencephalon: 21.76±1.40; mes-/diencephalon: 21.89±1.49; pituitary:
173 22.34±1.55.

174

175 **2.4.2. SYBR Green assay**

176 To determine the expression of each ESR and GPER gene, qPCR assays were
177 performed using a model 7500 unit (Applied Biosystems; Foster City, CA, USA) with

178 Maxima SYBR Green/ROX qPCR Master Mix (Fermentas Corp. Glen Burnie, MD,
179 USA). The qPCR program used for all was an initial step of 50 °C for 2 min, followed
180 by 95 °C for 10 min, and 40 cycles of 95 °C for 1 s and 60 °C for 10 s and 72 °C for 7
181 s. To evaluate assay specificity, the machine performed a melting curve analysis
182 directly after PCR by slowly (0.1 °C/s) increasing the temperature from 68 to 95 °C,
183 with a continuous registration of any changes in fluorescent emission intensity.

184 The total volume for each qPCR reaction was 20 µl, with 5 µl of diluted cDNA
185 template, forward and reverse primers (250 nM each), and SYBR Green/ROX Master
186 Mix (12 µl). The transcript levels were determined by the efficiency-adjusted relative
187 quantification method described by Weltzien et al. (2005). Serial dilutions of the cDNA
188 pool of the gonad tissues were run in duplicate and used for the standard curve to
189 measure all ESRs and GPERs in the testis. Serial dilutions of the cDNA pool of the
190 brain and pituitary tissues were used for the standard curve from which to measure all
191 the ESRs and GPERs in the different parts of the brain and pituitary. A 1/32 dilution of
192 the standard curve was included in each run of the corresponding gene as a calibrator.

193 Target and reference genes in unknown samples were run in duplicate PCR reactions. A
194 non-template control (cDNA replaced by water) for each primer pair was run in
195 duplicate on all plates. All ESR and GPER data were normalised to eel reference gene
196 ARP. qPCR calculations were performed according to the Roche Applied Science
197 protocol, Technical Note No. LC 13/2001, part 4 “Calibrator normalized relative
198 quantification”.

199

200 **2.5. Immunoassays for E₂**

201 Plasma concentrations of E₂ were measured by means of radioimmunoassay (RIA),
202 according to the method described by Schulz (1984). Free (i.e. not conjugated) steroids

203 were extracted from 200 μ l plasma, with 4 ml diethyl ether after vigorously shaking for
204 4 min. The aqueous phase was frozen in liquid nitrogen, whereas the organic phase was
205 transferred to a glass tube, evaporated in a water bath at 45 °C and then reconstituted by
206 addition of 600 μ l assay buffer, and then assayed for E₂. The assay characteristics and
207 cross-reactivities of the E₂ antiserum were previously examined by Frantzen et al.
208 (2004) and further validated for eel plasma by Mazzeo et al. (2014). The limit for the
209 assay was 0.2 ng/ml. The inter- and intra-assay coefficients of variation (CV) for the E₂
210 assay were 9.40% (n = 4) and 8.39% (n = 11), respectively. The standard curve was
211 made up of nine different concentrations of non-radioactive steroid, and ranged from
212 0.15 to 40 ng/ml. It was prepared by eight (1:1) serial dilutions in RIA buffer, starting at
213 40 ng/ml.

214

215 **2.6. Statistics**

216 Statistical analyses were performed to study the expression of ESRs and GPERs in the
217 different tissues, as well as the E₂ plasma levels in FW, SW and throughout
218 spermatogenesis (SPGA, SPGB/SPC, SD or SZ).

219 Two non-parametric tests were performed. Means between FW and SW conditions were
220 compared by Mann-Whitney *U* test. Means between the stage of spermatogenesis were
221 compared by Kruskal-Wallis ANOVA. Differences were considered significant when
222 $p < 0.05$.

223 All statistical procedures were performed using Statgraphics Plus 5.1 (Statistical
224 Graphics Corp., Rockville, MO, USA). Results are presented as mean \pm standard error
225 (SEM).

226

227

228 **3. Results**

229 Male eel ESRs (ESR1, ESR2a and ESR2b) and GPERs (GPERa and GPERb)
230 expressions were studied in the BPG axis after the transfer from FW to SW (before hCG
231 treatment) and throughout hormonally induced spermatogenesis (SPGA, SPGB/SPC,
232 SD and SZ stages). E₂ plasma levels were also measured.

233

234 **3.1. Histological observation**

235 The different spermatogenic stages were determined based on histological analyses of
236 European eel testis during hCG hormonal treatment. Mean GSI were calculated for each
237 spermatogenic stage: SPGA: GSI = 0.07±0.02; SPGB/SPC: GSI = 0.74±0.1; SD: GSI =
238 3.65±0.4; SZ: GSI = 7.89±0.4. Spermiating males were observed from the fifth week of
239 hCG treatment (W5).

240

241 **3.2. Brain estrogen receptor expressions**

242 In all the brain parts, the expressions of the three ESRs (ESR1, ESR2a, ESR2b)
243 increased from FW to SW (Figs. 2, 3 and 4) (9.75, 4.7 and 6.7-fold higher in SW in the
244 olfactory bulb, respectively; 8, 5, and 3.7-fold higher in SW in the telencephalon,
245 respectively; and 7.8, 3.8 and 3.5-fold higher in SW in the mes-/diencephalon,
246 respectively) (p<0.05).

247 During spermatogenesis, ESRs mRNA in the brain remained stable until the end of the
248 experiment, except in the mes-/diencephalon, where the expression of the three ESRs
249 increased (p<0.05), with them being 1.9, 2 and 1.7-fold higher respectively at SZ than at
250 SPGA.

251 Concerning the GPERs, the expression levels of GPERa in the telencephalon and
252 GPERb in the mes-/diencephalon decreased with the change from FW to SW (p<0.05),

253 with them being 12.5 and 4-fold higher respectively. GPERa and GPERb expression
254 levels then remained stable until the end of spermatogenesis.

255

256 **3.3. Pituitary estrogen receptor expressions**

257 In the pituitary, the expression of the three ESRs increased with the change from FW to
258 SW ($p<0.05$) (Fig. 5). The expression of ESR1, ESR2a and ESR2b was 7, 4 and 3-fold
259 higher respectively, in SW than in FW. During spermatogenesis, ESR1 and ESR2b
260 showed significant changes in the pituitary throughout testis development, while ESR2a
261 remained stable. ESR1 expression levels increased from stage SPGA to stage
262 SPGB/SPC ($p<0.05$), and then decreased from SPGB/SPC until the final maturation
263 stage SZ ($p<0.05$). ESR2b remained stable from stage SPGA to stage SPGB/SPC, and
264 then progressively decreased until the final maturation stage SZ.

265 Concerning the GPERs, the expression levels of both GPERa and GPERb increased
266 with the change from FW to SW ($p<0.05$), with them being 4 and 2-fold higher
267 respectively. GPERa and GPERb expression levels then remained stable until the end of
268 spermatogenesis.

269

270 **3.4. Testis estrogen receptor expressions**

271 In the testis, the expression levels of ESR1 and ESR2a increased with the change from
272 FW to SW ($p<0.05$), with them being 3 and 2-fold higher respectively (Fig. 6). In
273 contrast, the expression of ESR2b remained stable with the change of salinity. During
274 spermatogenesis, the expressions of the three ESRs decreased sharply from stage SPGA
275 to SPGB/SPC: the expressions of ESR1, ESR2a and ESR2b were 7, 9.5 and 9-fold
276 lower respectively, at stage SPGB/SPC compared to stage SPGA, then remained low
277 until stage SZ ($p<0.05$).

278 Neither GPERa nor GPERb expression levels altered significantly between FW and
279 SW, but they did increase progressively and markedly throughout spermatogenesis
280 ($p < 0.05$), with levels being 14.2 and 15.2-fold higher at stage SZ than at stage SPGA.

281

282 **3.5. Estradiol plasma levels**

283 E_2 plasma levels increased significantly with the change from FW to SW (Fig. 7), with
284 them being 4-fold higher in SW than in FW ($p < 0.05$). In hCG treated males, E_2 plasma
285 level decreased progressively until stage SZ ($p < 0.05$), and were 2-fold lower at stage SZ
286 than at stage SPGA. By the end of spermatogenesis, E_2 plasma thus returned to very
287 similar levels to those recorded in the initial FW stage.

288

289 **4. Discussion**

290 **4.1. Effect of salinity on estradiol and its receptors**

291 Before any hormonal treatment, the E_2 plasma levels of immature male European eels
292 increase sharply with the change from FW to SW (37 g/l), suggesting that salinity plays
293 a role in sex steroidogenesis. These results match those from previous studies, where the
294 increase in salinity to SW conditions augmented E_2 plasma levels in both male
295 (Peñaranda et al., 2016) and female eels (Quérat et al., 1987). According to Quérat et al.
296 (1987), the E_2 plasma level was higher in SW than in FW, in both hypophysectomized
297 as well as in intact female silver European eels, which suggests that an extra-pituitary
298 mechanism is at work modulating levels of E_2 with the transfer to SW. The increase of
299 E_2 in SW may be related to the osmoregulation, as relation between osmoregulation and
300 reproduction has been demonstrated. For instance in salmon, an anadromous species,
301 there is evidence of a negative relationship between sexual maturation and SW
302 adaptability (for review, see McCormick and Naiman, 1985; Lundqvist et al.,

303 1989; Staurnes et al., 1994; Madsen et al., 1997). E₂ is shown to be inversely correlated
304 with pretransfer gill Na⁺/K⁺-ATPase activity: E₂ may have an important role during the
305 development, as elevated plasma E₂ levels has a deleterious effect on hypo-
306 osmoregulatory physiology (Madsen et al., 1997). In contrary, the catadromous
307 European eel may respond in the opposite way to its oceanic salinity changes, attending
308 to its reproductive migration in SW.

309 The change from FW to SW induced variation of E₂ receptor expression through the
310 BPG axis. ESR1, ESR2a and ESR2b expression levels increased in the anterior brain
311 and in the pituitary, as well as GPERa and GPERb in the pituitary, and ESR1 and
312 ESR2a in the testis. In contrary, GPERa increased in the telencephalon and GPERb
313 increased in the mes-/diencephalon. These results suggest that the increase in salinity, in
314 the absence of any hormonal treatment, affects the expression of reproductive genes
315 along the gonadotropic axis of the male European eel.

316 The parallel increases in E₂ plasma levels and E₂ receptors could reflect a positive
317 autoregulation by E₂ of the expression of its receptors. In teleosts, the ESRs appeared to
318 be differentially regulated by E₂, according to tissue, stage of maturation, gender and
319 species. In the Japanese female eel, E₂ treatment induces an up-regulation of ESR1 but
320 not of ESR2 (Jeng et al., 2012b). In the European eel, while ESR and E₂ showed the
321 same expression pattern with the change of salinity, ESR expression levels in brain and
322 pituitary remained high during the spermatogenesis whereas E₂ plasma levels sharply
323 decreased through the spermatogenesis. In the fathead minnow (*Pimephales promelas*),
324 E₂ treatment induces an up-regulation of ESR1 in the testis and a down-regulation in the
325 ovaries, a down-regulation of ESR2b in both male and female gonads, while ESR2a
326 does not appear to be affected. In the pituitary, the three ESRs are up-regulated in
327 females but no significant differences (ESR1, ESR2b) or decreases (ESR2a) have been

328 shown in males undergoing E₂ treatment. In the brain no significant change was
329 observed for any ESRs (Filby et al., 2006). In goldfish, the response of the ESRs to E₂
330 seems to depend on the stage of maturation (for review see Nelson and Habibi, 2013).
331 The increase in the E₂ receptor expression levels in SW observed in the male eels in this
332 experiment could also be the result of other hormones involved in SW tolerance. For
333 instance, the growth hormone (GH) is a pleiotropic hormone which regulates various
334 functions in teleosts, including SW acclimation (McCormick, 2001; Eckert et al., 2001).
335 In salmonids, growth hormone acts in synergy with cortisol to increase seawater
336 tolerance (McCormick, 2001). In the European eel, GH was shown to increase the
337 number of nuclear E₂ receptors, as seen in the liver (Messaouri et al., 1991; Peyon et al.,
338 1996). Future studies could investigate the potential effect of GH or other
339 osmoregulatory hormones on E₂ receptors in the BPG axis.

340

341 **4.2. Brain and pituitary estrogen receptor expression levels during induced** 342 **spermatogenesis**

343 Several studies have demonstrated that E₂ is an important regulatory factor in the brain,
344 due to the role it plays in the neuroendocrine system controlling reproductive functions
345 (for review see Beyer, 1999), but also because of its neurotrophic, neuroprotective and
346 organizational properties (Behl, 2002). In this study we observed a progressive decrease
347 in male European eel E₂ plasma levels during induced spermatogenesis, contrary to the
348 increase observed in experimentally matured female eels (Pérez et al., 2011).
349 Nevertheless, in both sexes, androgen (testosterone and 11-ketotestosterone) plasma
350 levels increase during induced maturation (Aroua et al., 2005; Peñaranda et al., 2010).
351 A local production of E₂ in the brain/pituitary could exert autocrine and/or paracrine
352 actions in these organs, throughout spermatogenesis. Although most teleosts possess

353 duplicated aromatase *cyp19a1* genes (enzyme responsible of the conversion of
354 androgens to estrogens), eels have a single *cyp19a1* expressed in the brain, pituitary and
355 gonads (Ijiri et al., 2003; Jeng et al., 2012a; Peñaranda et al., 2014). Jeng et al. (2012a),
356 showed that hCG in males strongly up-regulates aromatase immunoreactivity in the
357 brain and pituitary, leading to a local production of E₂. In rainbow trout (*Oncorhynchus*
358 *mykiss*), aromatase expression has been localized to the neuroendocrine regions which
359 also express ESRs, indicating that locally produced estrogens can affect neuroendocrine
360 functions through genomic effects in a paracrine way (Menuet et al., 2003; Diotel et al.,
361 2010). According to Pellegrini et al. (2005), in fish, E₂ may be also involved in the
362 regulation of neuro-glial communications in the hypothalamus and in the
363 neurohypophysis.

364 In this study, ESR2a expression levels in the pituitary remained stable throughout
365 spermatogenesis, whereas ESR1 and ESR2b expression levels decreased, until testis
366 development was complete. According to Lafont et al. (in press), ESR1 expression
367 levels in the pituitary of female European eels increased as the eels matured, showing a
368 difference in the expression of ESR1 between the sexes. ESR2a and ESR2b expression
369 levels remained unchanged in both sexes (Lafont et al., in press; this study).

370 In the brain, the expression levels of the three ESRs progressively increased in the mes-
371 /diencephalon until the spermatozoa stage. In the female European eel, only ESR1
372 levels increased in the forebrain with maturation, with ESR2a and ESR2b levels
373 remaining stable (Lafont et al., in press). These results again suggest a differential
374 regulation of the nuclear estrogen receptors during the maturation of male and female
375 eels. According to Jeng et al. (2012b), E₂ plasma levels may up-regulate the expression
376 of ESR1 in the brain of female Japanese eel. As E₂ plasma levels decreased throughout
377 spermatogenesis in male eels, the expression of ESRs does not seem to respond to a

378 peripheral E₂ production but rather to a local production in the brain. Nevertheless, other
379 factors may be involved in the increase in the expression of ESR1, ESR2a and ESR2b
380 in the brain. An interaction between androgen and estrogen in endocrine tissues has
381 already been demonstrated (Panet-Raymond et al., 2000), and a cross talk between
382 androgens and estrogens and their receptors has been highlighted in mice bone
383 (Kousteni et al., 2001). Finally, according to Larsson et al. (2002), both androgens and
384 estrogens are involved in the physiological regulation of brain androgen receptors in the
385 Atlantic croaker (*Micropogonias undulatus*), another teleost species, during the
386 reproductive cycle. In the male European eel, we cannot rule out an interaction between
387 androgens and the regulation ESR1, ESR2a and ESR2b expressions in the brain.

388 Concerning GPERs, they are both expressed in the brain and in the pituitary of male
389 European eel, just as previously seen in the female (Lafont et al., in press). These results
390 suggest GPERs in the eel brain are likely to play a role in the reproductive system of
391 both sexes. These two studies on eel are so far the only ones to look at duplicated
392 GPERs. Previous studies on zebrafish and humans have discovered GPER expressed in
393 different brain regions, including regions that control reproduction and sex behavior
394 (Liu et al., 2009; reviewed by Olde and Leeb-Lundberg, 2009). We did not observe any
395 major change in the expression of GPERs in the brain and pituitary during
396 spermatogenesis.

397 ESRs and GPERs are co-expressed in all the brain parts and in the pituitary of male as
398 well as female European eels. Further analyses are required but these results may
399 suggest potential interactions between nuclear and membrane E₂ receptors.

400

401 **4.3. Testis estrogen receptor expression during spermatogenesis**

402 The three ESRs were shown to be expressed in the eel testis, and with the same

403 expression pattern throughout spermatogenesis. The highest expression levels were
404 measured at stage SPGA. All three ESR expression levels then sharply decreased at
405 stage SPGB/SPC, and remained low until the end of spermatogenesis.

406 When studying Japanese eel, Miura et al. (1999) discovered that E₂ plays an important
407 role in spermatogonial renewal. They demonstrated that low concentrations of E₂ act in
408 the primary stages of spermatogonia through receptors present in Sertoli cells,
409 stimulating and maintaining spermatogonia proliferation prior to the progression of
410 further stages of spermatogenesis. The high expression levels of all ESRs at stage
411 SPGA that we observed corroborates the proposed role of estrogens as a spermatogonial
412 renewal factor. In the European eel, the parallel regulation of the three ESRs suggests
413 that the role of E₂ as a spermatogonial renewal factor is mediated by ESR1, ESR2a and
414 ESR2b.

415 In the testis, the expression pattern of GPERs during spermatogenesis was notably
416 different from that of ESRs. The expression levels of both GPERs were low at the
417 beginning of the spermatogenesis (SPGA stage), and increased sharply until the end of
418 spermatogenesis. These results suggest that both GPERa and GPERb may play a role in
419 the final sperm maturation process. The high expression levels of both GPERs at the
420 spermatozoa stage suggests that these receptors may be mainly localized in the germ
421 cells in the eel. Nevertheless, as GPER was shown to be localized in the Leydig and
422 Sertoli cells, spermatogonia, spermatocytes and spermatozoa of humans, in the
423 spermatogonia cell line of mice (*Mus musculus*), and in the Sertoli cells, pachytene
424 spermatocytes and round spermatids of rats (*Rattus norvegicus*) (Luconi et al., 2004;
425 reviewed by Correia et al., 2015), it cannot be excluded that this may be due to Sertoli
426 cell proliferation in the final sperm maturation process.

427 It is well known that in female fish GPER mediates the E₂-induced meiotic arrest of

428 oocytes (Pang et al., 2008; Pang and Thomas, 2009). Nevertheless, its role in male fish
429 has still yet to be explored. In adult male zebrafish (*Danio rerio*), a RT-PCR analysis of
430 gene expression in the isolated purified early and late germ cells revealed that GPER is
431 mainly expressed in early germ cells of the testis, including the spermatogonia and
432 spermatocytes, suggesting GPER may play a part in mediating estrogen action early on
433 in spermatogenesis (Liu et al., 2009). In contrast, according to their expression profile
434 during spermatogenesis, European eel GPERs (GPERa and GPERb) may have a major
435 role in the final sperm maturation process. This suggests that GPER have differential
436 functions during spermatogenesis depending on the teleost species. In humans, although
437 the role of estrogens in spermatogenesis is still unclear, decreased spermatozoa numbers
438 and motility in men who were genetically deficient in aromatase were observed,
439 suggesting aromatase/estrogens play a role not only during the development and
440 maintenance of spermatogenesis, but also in the final maturation of spermatozoa
441 (Carreau et al., 2010; reviewed by Correia et al., 2015). Accordingly, studies have
442 demonstrated the presence of sex steroid membrane receptors in human spermatozoa
443 and have shown that the effects of estrogens and progesterone on human ejaculated
444 spermatozoa may be mediated by GPERs and membrane progestin receptors (mPRs)
445 respectively (Revelli et al., 1998; Luconi et al., 2004; Carreau et al., 2010). Similarly, in
446 the European eel, sex steroid membrane receptors may be involved in final sperm
447 maturation. We recently found high expression levels of mPRs in the final stage of
448 spermatogenesis in male eel testis (Morini et al., submitted). This study suggests that,
449 both in the eel as well as in humans, progestin and estrogen membrane receptors have
450 are implicated in the completion of spermatogenesis.

451 In conclusion, this is the first study to describe the expression levels of five estrogen
452 receptors (three nuclear ESRs and two membrane GPERs) along the BPG axis of a male

453 teleost. The presence of all these estrogen receptors in theBPG axis suggests an
454 implication in the control of male eel reproduction. Our results support the evidence that
455 the regulation of eel ESRs and GPERs expressions are tissue and stage-specific, as
456 shown by the different expression profiles found in the different BPG tissues throughout
457 spermatogenesis. It appears likely that testes ESRs play a role in spermatogonia
458 renewal, while testes GPERs are mainly involved in the end of spermatogenesis. The
459 three ESRs and both GPERs in the brain and pituitary may control neuroendocrine
460 functions, mediating the autocrine or paracrine actions of locally-produced estrogen,
461 during the entire process of eel spermatogenesis. Finally, the expression of both ESRs
462 and GPERs in the BPG axis indicates a possible cooperation between genomic and non-
463 genomic estrogen actions in the control of reproduction.

464

465 **Acknowledgements**

466 Funded by the Spanish Ministry of Science and Innovation (REPRO-TEMP project;
467 AGL2013-41646-R) and IMPRESS (Marie Skłodowska-Curie Actions; Grant
468 agreement n°: 642893). M.C. Vílchez has a predoctoral grant from UPV PAID
469 Programme (2011-S2-02-6521), M. Morini has a predoctoral grant from Generalitat
470 Valenciana (Programa Grisolia). D.S. Peñaranda was supported by MICINN and UPV
471 (PTA2011-4948-I)..

472

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684 **Table legend**

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686 **Table I. Quantitative PCR primer sequences for nuclear estrogen receptors (*ESR1*,**
687 ***ESR2a*, *ESR2b*) and membrane progesterin receptors (*GPERa* and *GPERb*).**

688

689

690 **Figure legends**

691

692 **Figure 1. Histological sections of European eel testis at different developmental**
693 **stages during human chorionic gonadotropin (hCG) hormonal treatment. A: SPGA**
694 **(spermatogonia A); B: SPGB/SPC (spermatogonia B/spermatocyte); C: SD (spermatid),**
695 **D: SZ (spermiation). Scale bar: A=100 μ m; B= 10 μ m, C, D= 50 μ m; Cell types: SPG=**
696 **spermatogonia; SPC: spermatocytes; SD: spermatids; SZ: spermatozoa**

697

698 **Figure 2. Expression in the olfactory bulb of the male European eel ERs (*ESR1*,**
699 ***ESR2a* and *ESRb2*) and GPERs (*GPERa* and *GPERb*) in freshwater (FW, n=6) and**
700 **sea water conditions (SW, n=5), and through the stages of testis development.**
701 Means are given \pm SEM. Differences were considered significant when $p < 0.05$.
702 Asterisks indicate significant differences between FW and SW condition. SPGA=
703 Spermatogonia A stage (n=8), SPG/SPC= Spermatogonia B/Spermatocyte stage (n=11),
704 SD= Spermatid stage (n=9), SZ= Spermatozoa stage (n=18). See main text for
705 description of development stages.

706

707 **Figure 3. Expression in the telencephalon of the male European eel ERs (*ESR1*,**
708 ***ESR2a* and *ESRb2*) and GPERs (*GPERa* and *GPERb*) in freshwater (FW, n=7) and**
709 **sea water conditions (SW, n=6), and through the stages of testis development.**
710 Means are given \pm SEM. Differences were considered significant when $p < 0.05$.
711 Asterisks indicate significant differences between FW and SW condition. SPGA=
712 Spermatogonia A stage (n=9), SPGB/SPC= Spermatogonia B/Spermatocyte stage
713 (n=11), SD= Spermatid stage (n=10), SZ= Spermatozoa stage (n=21). See main text for
714 description of development stages.

715

716 **Figure 4. Expression in the mes/-diencephalon of the male European eel ERs**
717 **(*ESR1*, *ESR2a* and *ESRb2*) and GPERs (*GPERa* and *GPERb*) in freshwater (FW,**

718 **n=7) and sea water conditions (SW, n=6), through the stages of testis development.**
719 Means are given \pm SEM. Differences were considered significant when $p < 0.05$.
720 Asterisks indicate significant differences between FW and SW condition; small letters
721 indicate significant differences through the stages of development (hCG treated). SPGA=
722 Spermatogonia A stage (n=9), SPGB/SPC= Spermatogonia B/Spermatocyte stage
723 (n=12), SD= Spermatid stage (n=10), SZ= Spermatozoa stage (n=23). See main text for
724 description of development stages.

725

726 **Figure 5. Expression in the pituitary of the male European eel ERs (*ESR1*, *ESR2a***
727 **and *ESRb2*) and GPERs (*GPERa* and *GPERb*) in freshwater (FW, n=5) and sea**
728 **water conditions (SW, n=6), and through the stages of testis development.** Means
729 are given \pm SEM. Differences were considered significant when $p < 0.05$. Asterisks
730 indicate significant differences between FW and SW condition; small letters indicate
731 significant differences through the stages of development (hCG treated). SPGA=
732 Spermatogonia A stage (n=7), SPGB/SPC= Spermatogonia B/Spermatocyte stage
733 (n=12), SD= Spermatid stage (n=8), SZ= Spermatozoa stage (n=19). See main text for
734 description of development stages.

735

736 **Figure 6. Expression in the testis of the male European eel ERs (*ESR1*, *ESR2a* and**
737 ***ESRb2*) and GPERs (*GPERa* and *GPERb*) in freshwater (FW, 5) and sea water**
738 **conditions (SW, 7), and through the stages of testis development.** Means are given \pm
739 SEM. Differences were considered significant when $p < 0.05$. Asterisks indicate
740 significant differences between FW and SW condition; small letters indicate significant
741 differences through the stages of development (hCG treated). SPGA= Spermatogonia A
742 stage (n=11), SPGB/SPC= Spermatogonia B/Spermatocyte stage (n=14), SD=
743 Spermatid stage (n=10), SZ= Spermatozoa stage (n=24). See main text for description
744 of development stages.

745

746 **Figure 7. Estradiol (E_2) plasma level of male European eel in freshwater (FW, n=6)**
747 **and sea water conditions (SW, n=5), and through testis of development stage.**
748 Means are given \pm SEM. Differences were considered significant when $p < 0.05$.
749 Asterisks indicate significant differences between FW and SW condition; small letters
750 indicate significant differences through the stages of development (hCG treated). SPGA=
751 Spermatogonia A stage (n=10), SPGB/SPC= Spermatogonia B/Spermatocyte stage

752 (n=15), SD= Spermatid stage (n=10), SZ= Spermatozoa stage (n=25). See main text for
753 description of development stages.
754

755 **Table I**

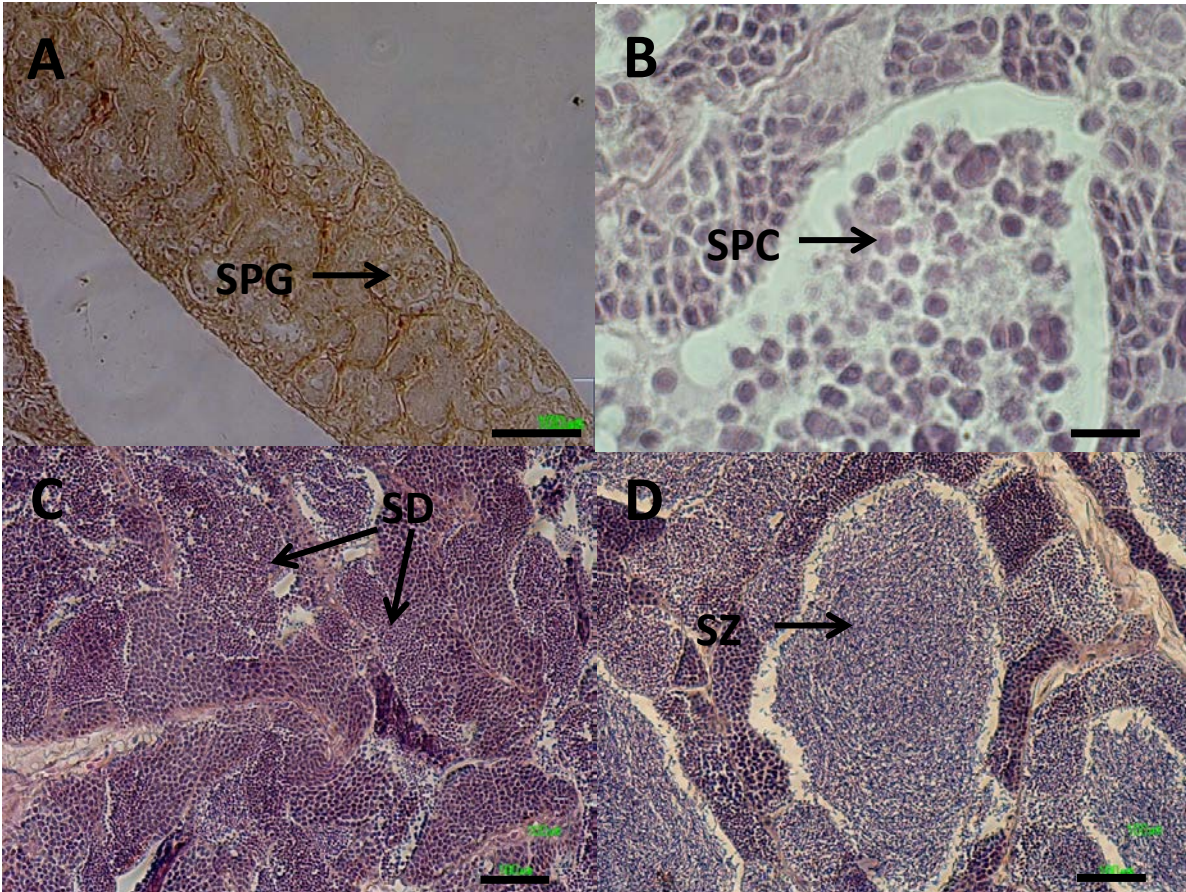
Name	Sequence (5'-3')	Orientation	Tm	Reference	Accession number	Primer efficiency	Amplicon size (bp)
<i>ARP</i>	GTGCCAGCTCAGAACAACCTG	Forward	56.36	Weltzien et al., 2005	AAV32820	2.14	107
	ACATCGCTCAAGACTTCAATGG	Reverse	60.09				
<i>ESR1</i>	GCCATCATACTGCTCAACTCC	Forward	58.20	Lafont et al., in press	CUH82767	2.04	75
	CCGTAAAGCTGTCGTTTCAGG	Reverse	59.32				
<i>ESR2a</i>	TGTGTGCCTCAAAGCCATTA	Forward	58.71	Lafont et al., in press	CUH82768	2	68
	AGACTGCTGCTGAAAGGTC	Reverse	57.16				
<i>ESR2b</i>	TGCTGGAATGCTGCTGGT	Forward	59.93	Lafont et al., in press	CUH82769	1.95	120
	CCACACAGTTGCCCTCATC	Reverse	58.44				
<i>GPERa</i>	CAACTCAACCACCGGGAGA	Forward	61.81	Lafont et al., in press	CUH82770	1.99	165
	TGACCTGGAGGAAGAGGGACA	Reverse	62.86				
<i>GPERb</i>	CAACCTGAACCACACGGAAA	Forward	60.36	Lafont et al., in press	CUH82771	1.97	162
	TGACCTGGAAGAAGAGGGACA	Reverse	60.59				

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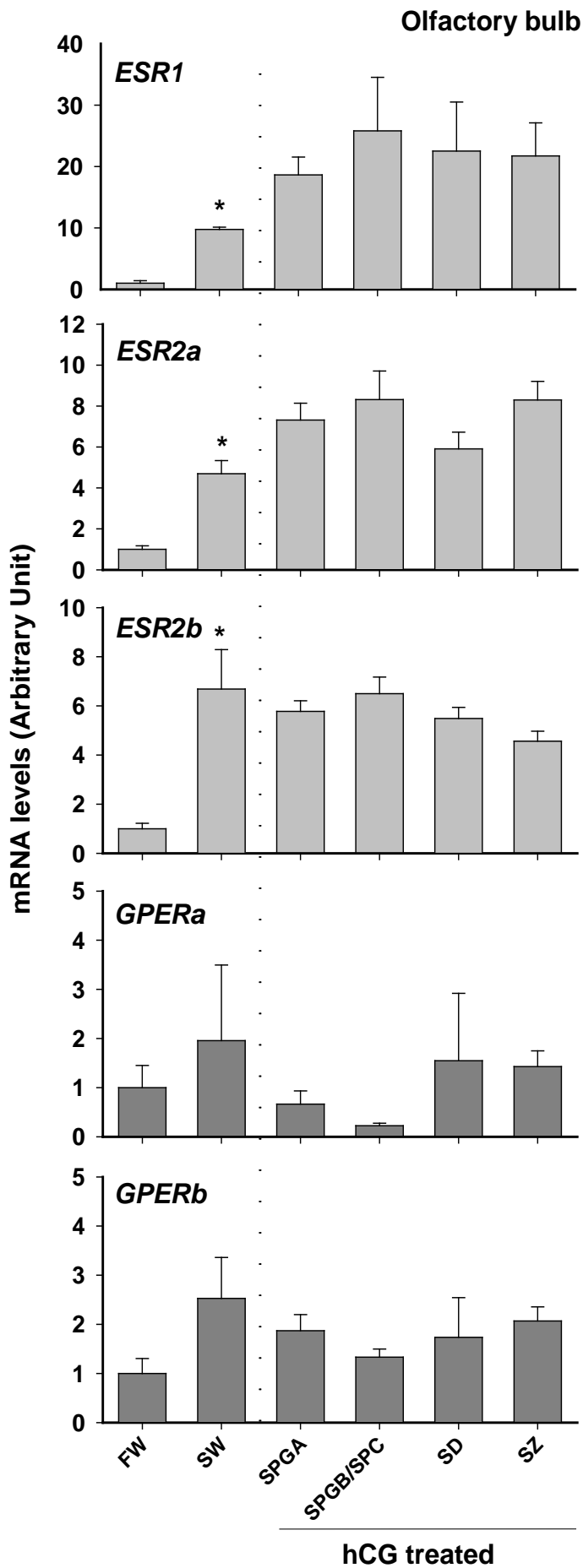
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Figure 1



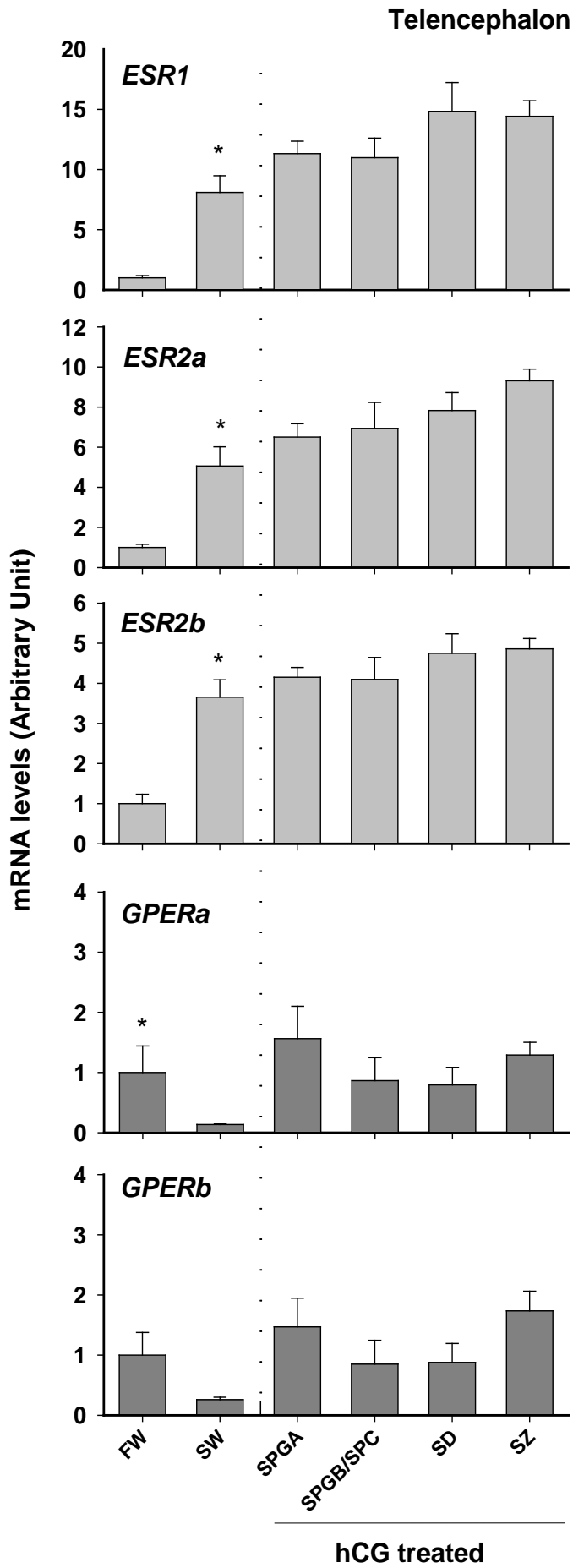
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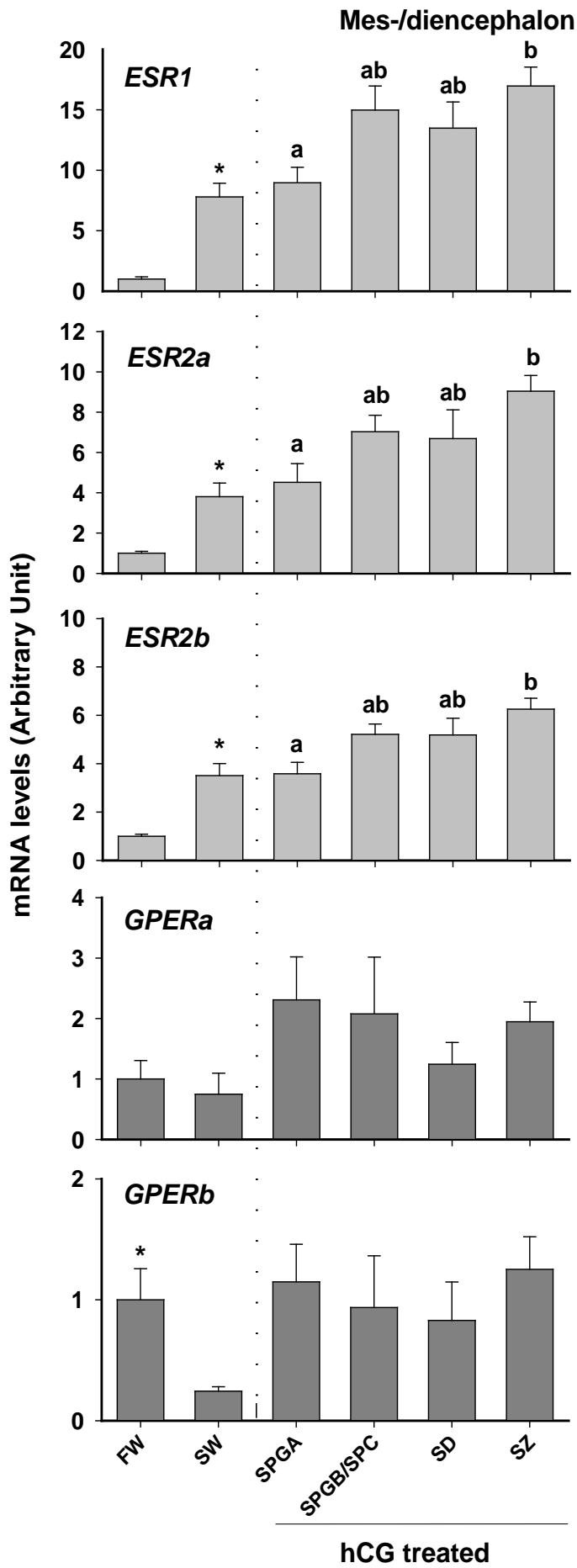
Figure 2

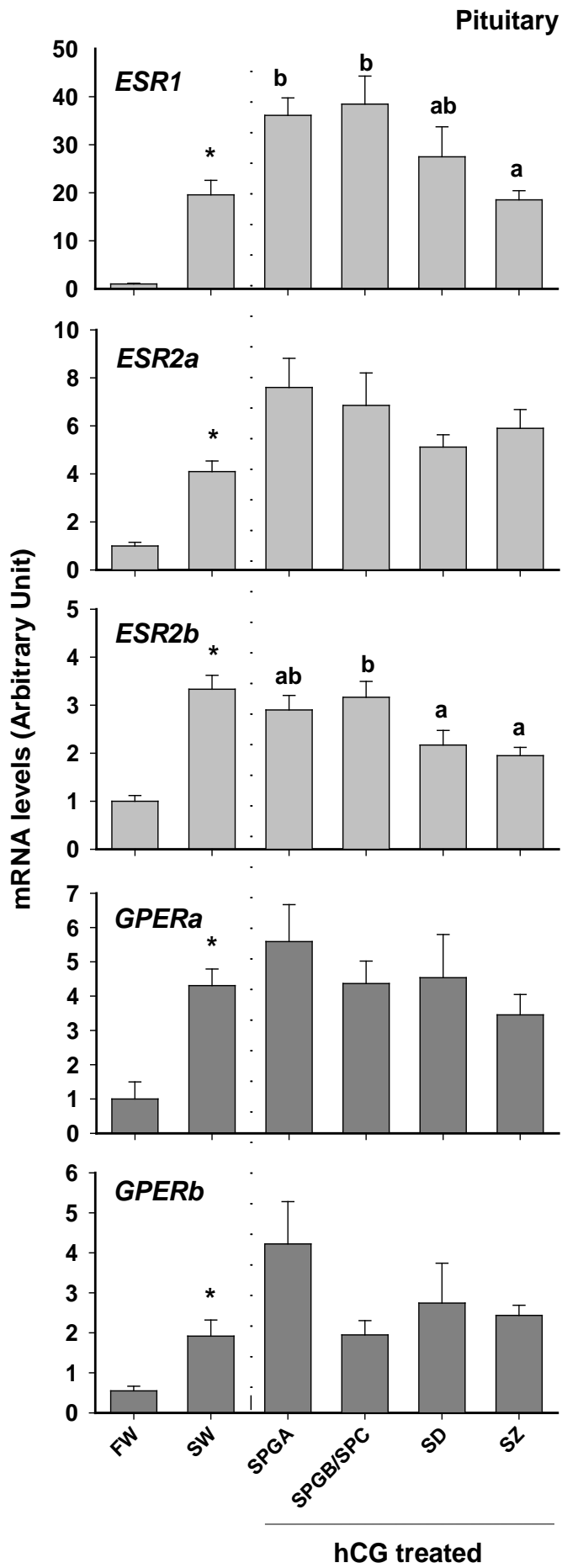


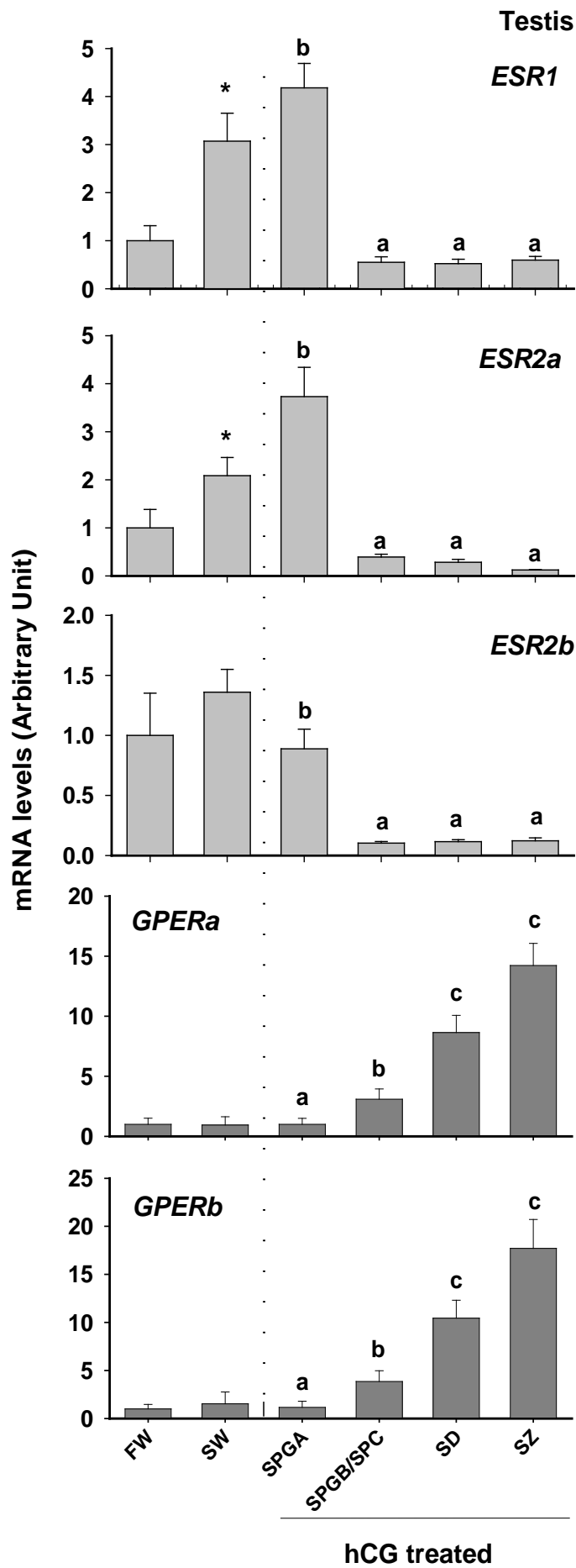
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Figure 3

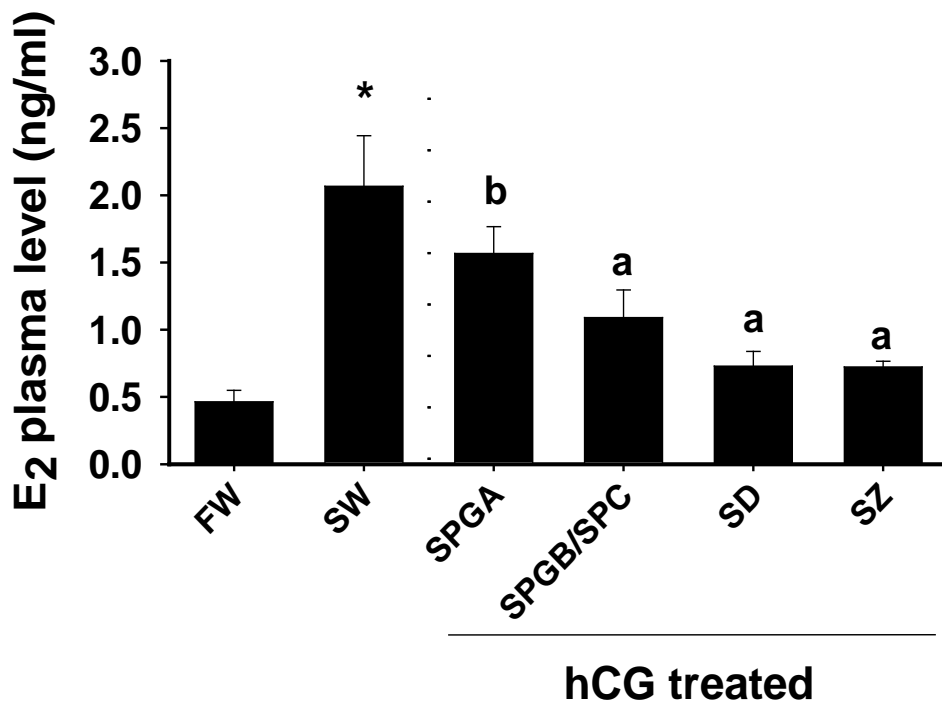








774 Figure 7



775