1	Evaluation of toxicological endpoints in female zebrafish after
2	bisphenol A exposure
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25 ABSTRACT

Given the importance of bisphenol A (BPA) as a xenoestrogen and its potential effects 26 on human and animal health, we evaluated BPA exposure's short-term effects on 27 follicular development, volk protein vitellogenin (VTG) production and aromatase 28 expression in female zebrafish. Histological modifications were observed along with 29 30 increased presence of atretic follicles. Whole-body VTG concentration increased with the dose of BPA exposure. In contrast, expression of Cyp19a mRNA in the ovaries of BPA-31 exposed fish exhibited an apparent non-monotonic response curve, marked by 32 downregulation at 1 µg/L BPA, upregulation at 10 µg/L BPA, and a return to 33 downregulation at 100 µg/L BPA and higher doses. Ovaries only exhibited significant 34 increases in follicular atresia and VTG concentration after exposure to 100 µg/L BPA and 35 higher doses. Ovarian histopathology, aromatase Cyp19a transcript levels and whole-36 body VTG protein abundance may be good biomarkers for early detection of 37 38 environmental BPA exposure.

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40 Keywords: bisphenol A; zebrafish; aromatase; vitellogenin; atretic follicle

42 **1. Introduction**

43 Endocrine-disrupting chemicals (EDCs) have attracted considerable worldwide scientific and public attention due to their potential adverse effects on exposed organisms. 44 45 Xenoestrogenic endocrine-disrupting compounds are found in the environment as a result of industrial and manufacturing activities (Cheng et al., 2011). Among these compounds, 46 47 bisphenol A (BPA) is one of the most highly produced chemicals worldwide. BPA is ubiquitously present as an environmental contaminant in rivers and drinking water, most 48 likely due to the migration of plastic containers from industrial waste sites. In river water 49 BPA levels were reported of 0.01-21 µg/L while higher concentrations were also detected 50 near wastewater treatment plants or landfills (Crain et al., 2007; Kang et al., 2007; Naderi 51 52 et al., 2014). BPA is widely used in the manufacturing of various polycarbonate plastics 53 used in lacquer coatings of food cans and food and beverage containers. The widespread 54 distribution and environmental persistence of xenoestrogen BPA indicates a strong potential for human and animal exposure. Because humans are exposed to this compound 55 56 daily, the effects of BPA are relevant to public health (Graselli et al., 2010). This exposure primarily occurs via the hydrolysis of polycarbonate plastics and epoxy resins and results 57 in low concentrations of free BPA in food and liquids. The presence of endocrine-58 disrupting chemicals in the environment reportedly disturbs the normal endogenous 59 hormone pathway and interrupts reproductive development in wildlife species. These 60 compounds can affect the reproductive regulation of the neuroendocrine system, often by 61 mimicking or blocking endogenous hormones (Qin et al., 2013; Naderi et al., 2014). 62

BPA acts as an EDC by causing adverse biochemical and physiological changes that alter
the histological structure of cells and modify the function of tissues and organs,
interfering with reproductive efficiency. Within the reproductive system, ovarian
granulosa cells have been documented as a target of BPA action (Graselli et al., 2010). It

is well-known that granulosa cells play a crucial role in ovarian physiology through the 67 68 production of estrogens, which depends on androgen production in theca cells in addition to other factors that interact with the oocyte during its development. Therefore, the 69 disruption of their functional activities by BPA could have a significant impact on fertility 70 (Graselli et al., 2010). Therefore, BPA's effects in ovaries could be evaluated by 71 histology. However, molecular responses usually occur earlier than histological 72 73 perturbations, which are considered to be higher-level biological responses. The yolk protein vitellogenin (VTG) has been widely used as an endpoint for many exposure 74 studies on the estrogenic effect of BPA in fish (Holbech et al., 2006). Changes in VTG 75 76 levels in females have been suggested as a complementary biomarker of potential 77 reproductive disruption, as well as modifications in the sex hormone balance (Mandich et al., 2007). 78

Conversely, BPA, similar to some other EDCs, could also act as anti-androgens by 79 binding to the androgen receptor and producing specific alterations in gene expression 80 (López-Casas et al., 2012). The cytochrome P₄₅₀ CYP19 (aromatase) is involved in the 81 generation of estradiol from testosterone. In this way, CYP19 is considered a potential 82 EDC target because the modulation of its expression and function can dramatically alter 83 the rate of estrogen production (Cheshenko et al., 2008). Two different Cyp19 genes, 84 Cyp19a and Cyp19b, are expressed in many teleost fish, preferentially in the ovary and 85 the brain, respectively. Recent in vitro studies showed that BPA downregulates Cyp19a 86 mRNA and protein expression in rats (Lee et al., 2013), which is relevant to the evaluation 87 of the expression of the aromatase genes in ovaries as a biomarker of BPA exposure. 88

89 The purpose of our study was to evaluate the estrogenic activity of BPA as a ubiquitous90 environmental contaminant through the assessment of different biomarkers in the

91	zebrafish ovary. The toxicological endpoints were selected at different levels of biological
92	organization and included histological modifications in ovaries affecting the number of
93	atretic follicles, VTG measurements, and determinations of the aromatase Cyp19a mRNA
94	levels in zebrafish exposed to BPA (1, 10, 100 and 1000 μ g/L) for 14 days.

96 **2. Materials and Methods**

97 2.1. Fish exposure and sampling protocol

Sixteen-week-old female zebrafish (Danio rerio) (n=150; standard length: 4.17 ± 0.24 98 cm; 0.57±0.14 g wet weight) were used. The treated groups were exposed (OECD 99 Guideline No. 204) to graded concentrations (1, 10, 100 and 1000 µg/L) of BPA (Sigma-100 Aldrich[®]) for 14 days. A control group exposed to only unchlorinated tap water completed 101 the exposure design. The research procedure was conducted in the Experimental Animal 102 103 Service of the University of Córdoba (Spain) after approval by the animal care committee of the University of Córdoba (Spain) and in accordance with the European Regulations 104 for the Protection of Experimental Animals (Directive 2010/63/EU). 105

106 After 2 weeks of exposure, the zebrafish were sacrificed by an overdose of an anesthetic solution of tricaine methanesulfonate (MS-222[®] 500 mg/L; Sigma-Aldrich) buffered with 107 sodium bicarbonate (300 mg/L; Sigma-Aldrich). Immediately afterward, the standard 108 109 length (SL) and body weight (BW) of each animal were measured. The gonads from 30 animals (n=6 per group) were dissected and fixed for histological analysis for qualitative 110 and quantitative evaluations. To complete these experiments, each fish was necropsied 111 by placing it in right lateral recumbency on the stage of a dissecting microscope. The 112 ovaries of 45 additional zebrafish (n=9 per group) were removed, immediately immersed 113 114 in liquid nitrogen, and stored at -80°C for qRT-PCR. For the vitellogenin analysis, another 115 45 fish were frozen and stored at -80°C until further analysis. The final 30 fish (n=6 per group) were dried with sterile gauze, frozen and stored at -80°C until use in the 116 117 toxicological analysis (analytical BPA determinations).

118 **2.2. Light and electron microscopy**

For light microscopy, the fixed ovaries of 6 animals from each experimental group were routinely processed for paraffin sectioning by fixation in 10% buffered formalin solution, dehydration in a graded series of ethanol, immersion in xylol and embedding in paraffin wax by routine techniques. Tissue sections of 4 μ m were mounted. After deparaffinization, the sections were rehydrated, stained with hematoxylin and eosin, and mounted on microscope slides with Cristal/Mount (Paraplast, Oxford Labware, St. Louis, MO).

For electron microscopy, randomly selected gonad samples were first fixed in a 2% 126 glutaraldehyde solution in 0.1 M phosphate buffer (pH 7.4, 4°C, overnight) and later fixed 127 in 1% osmium tetroxide in 0.1 M phosphate buffer (pH 7.4) for 30 min. After dehydration 128 in a graded ethanol series and embedding in Araldite, semi-thin and ultra-thin sections 129 were cut on an LKB ultramicrotome (LKB). The semi-thin sections were stained with 130 131 toluidine blue, whereas the ultra-thin sections were double-stained with uranyl acetate and lead citrate. Tissue sections were examined in a JEM 1400 transmission electron 132 microscope (TEM; JEOL, Ltd.). 133

134 **2.3. Morphometric study**

For the structural quantifications, the fixed gonads were cut into three sections. Each portion was later processed and embedded in paraffin, as for routine histology. The first section (4 μ m thick) of each block was stained with hematoxylin and eosin and used for the stereological study.

139 The quantitative study was performed according to Molina et al. (2013). Each 140 microscopic image was processed using the Visilog 5[®] software (Noesis). Quantification 141 was performed in a blind manner by an observer experienced in the use of the analysis system (J.G.M.). The system was initially and regularly calibrated using a millimeterslide.

Atretic follicle identification was based on the determination of non-physiological alterations in four follicular components: oocyte, follicular cells, and pellucid zone. The numerical density (Q_A) of the atretic follicles in the plane was estimated using a test system consisting of sixteen rectangular counting frames superimposed onto each microscopic image. Thus, the number of profiles per area Q_A (*nucl/tis*) was estimated using the following equation:

150
$$estQ_A(nucl/tis) = \sum Q(nucl)/(\sum P(tis) \cdot a/p)$$

151 where Q_A (*nucl/tis*) is the numerical density of follicular nuclei per ovary, $\sum Q(nucl)$ is the 152 total number of nuclear profiles counted within the counting frames of the area obtained 153 from $\sum P(tis)$ (i.e., the total number of points in the tissue), and a/p is the area associated 154 with one point in the test system (in our study, $a/p = 125 \ \mu m^2$).

155 2.4. Whole-body vitellogenin measurements

The VTG levels in whole-body homogenates (n=9 per experimental group) were determined using a commercial enzyme-linked immunosorbent assay developed for *Danio rerio* (ELISA, Biosense Laboratories, Bergen, Norway) following the manufacturer's indications. A multiple range test was used to make multiple comparisons between groups.

161 2.5. *Cyp19a* transcript quantification by real-time qRT-PCR

162 2.5.1. RNA isolation

163 Nine zebrafish from each experimental condition were selected for this experiment. The
164 total RNA from the ovaries of each animal was isolated using the Isol-RNA Lysis Reagent
165 (5PRIME). Genomic DNA was removed by the Total RNA Cleanup with DNase

Digestion kit using the Qiagen RNeasy Protocol (Qiagen). The absence of any remaining gDNA contamination was confirmed by PCR amplification of the RNA samples without previous retrotranscription. An Agilent 2100 Bioanalyzer (Agilent Technologies) was used to determine the RNA integrity number (RIN) (Schroeder et al., 2006). The RNA purity and concentrations were determined by spectrophotometry. Only high-quality RNAs with RIN values > 8.5 and A260:A280 ratios close to 2.0 were used for the subsequent experiments (Fleige and Pfaffl 2006; Taylor and Mrkusich 2014).

cDNAs was generated from total RNA (1 µg/rx) by using the QuantiTect Reverse
Transcription Kit (Qiagen), according to the manufacturer's protocol.

175 2.5.2. Primer design and amplification efficiency

The Danio rerio sequences for Cyp19a and six candidate reference genes (ActB, B2m, 176 177 Hprt1, GusB. Nono and *Rpl13*) were obtained from GenBank 178 (http://www.ncbi.nlm.nih.gov/gene) to design the primers pairs. The primers (Supp. Inf. Table 1) were designed with Oligo 7 software (Molecular Biology Insights, Inc.), as 179 previously described (Pueyo et al., 2002). To obtain high specificity and performance, the 180 primers were required to have high Tm ($\geq 80^{\circ}$ C), optimal 3'- Δ G (\geq -5 kcal/mol) values 181 and no hairpin or duplex structures. 182

The amplification efficiency curves were obtained by preparing ten-fold serial dilutions from 2 x 10^5 to 0.02 pg of total RNA input, which were subsequently reverse transcribed and amplified by real-time PCR. The log of the RNA input versus the Ct value was plotted, and the efficiency value was estimated from the slope of the efficiency curve. All primer pairs amplified the selected targets with optimal ($\approx 100\%$) PCR efficiencies (Supp. Inf. Table 1 and Supp. Inf. Fig. 1) and produced specific PCR products, as confirmed by PCR product sequencing.

190 *2.5.3. qRT-PCR*

191 Real-time PCR reactions were performed in quadruplicate using 50 ng of cDNA template,

192 0.3 μM of each primer, 3 mM MgCl₂, 250 μM of each dNTP, 0.75 units of Platinum Taq

193 DNA polymerase, and 1:100,000 SYBR Green I dye (Roche) in a volume of 25 µl. The

194 reactions were analyzed on a Cycler Q Real-Time PCR System (Bio-Rad). The cycling

195 conditions were as follows: 2 min at 95°C for Platinum Taq activation and 40 cycles for

the melting (15 s, 95°C) and annealing/extension (30 s, 70°C) steps. Next, a melting curve

analysis was performed (60° C to 95° C) to verify the specificity of the amplicons.

- 198 For absolute quantification, the number of transcript molecules was calculated from the
- linear regression of a calibration curve (Jurado et al., 2003; Prieto-Alamo et al., 2003).

200

201 **2.6. Statistics**

The data were analyzed using the statistical program Statgraphic (Centurion XVI[®]) to 202 203 determine the effects of BPA in each of the treated groups. ANOVA (F-test) was used to determine significant differences between the mean values of each group. Fisher's LSD 204 post hoc test was used to perform multiple comparisons between groups. The results are 205 206 expressed as the mean \pm standard deviation (SD), and P < 0.05 was considered to be significant. In the transcriptional analysis, the statistically significant differences between 207 the control and experimental samples were determined using Student's t-test and were 208 expressed by * to denote P < 0.05; and ***, P < 0.001. The statistics program SigmaStat 209 210 5.1 (GraphPad Prism) was used throughout the study.

211 **3. Results**

None of the fish died during the study. There were no significant differences in the meanBW and SL between the control and BPA-exposed groups.

214 **3.1. Histopathology**

Follicular atresia (Fig. 1) was variable, depending on the type of affected follicle and the study group. In the control group, all follicles showed normal appearance under both the light and electron microscopes (Fig. 1 cA, cB). In the exposed groups, different levels of atresia were observed in the various follicular populations. Primordial follicles (Fovg) showed modifications, with atretic degeneration being visible at the intersection between follicular cells and oocytes. Moreover, both the optical and electronic microscope preparations showed total or partial vacuolization that may even be irregular (Fig. 1A, B).

The alveolar cortical follicles (Fc) exhibited a poor formation of the pellucid zone, leading 222 223 to egg-follicular cell separation due to vacuolization (Fig. 1C, D). In these follicles, a high amount of modifications was observed, especially under the light microscope, which 224 triggered follicular atresia with a total disintegration of the follicular components and 225 hyalinization in the oocyte cytoplasm. The vitellogenic follicles (Fv) exhibited full 226 vacuolization due to modifications affecting the oocyte, the pellucid zone and the 227 follicular cells. The poor formation of the pellucid zone caused the complete separation 228 of the oocyte from the follicle cells (Fig. 1E, F), similar to that observed in Fc. Finally, 229 mature follicles (Fm) showed a complete degeneration of all components (Fig. 1G, H), 230 231 yielding a breakdown of all organelles, primarily in the oocyte, as the cellular debris remained within the follicle until it further disintegrated. As the BPA concentration 232 increased, we observed a concomitant increase in the proportion of follicular atresia 233 234 affecting the different types of follicles. This trend of increasing follicular atresia was

235	observed in the Fc, the Fv and the Fm. Considering the total amount of follicular atresia
236	in each group, significant differences ($P < 0.05$) between the control group and other
237	study groups were observed. Similarly, significant differences ($P < 0.05$) were observed
238	between the groups treated with 1, 10, 100 and 1000 μ g/L BPA (Fig. 2).

- 240 3.2. Whole-body VTG levels
- 241

The whole-body VTG concentration in the control group was 21.13 ± 2.16 ng/g. These levels significantly increased (P < 0.05) in the exposed animals, even at the lowest BPA dose assayed (1 µg/L), in a dose-dependent manner (Fig. 3), reaching a 12-fold increase at the highest assayed dose.

247

248 **3.3. Real-time PCR**

3.3.1. Identification of valid reference genes for the normalization of the qRT-PCR expression studies.

The expression levels of six candidate reference genes (*ActB*, *B2m*, *Hprt1*, *GusB*, *Nono* and *Rpl13*) were evaluated using the threshold cycle (Ct) values from nine biological and three technical replicates from the five different experimental conditions. The box plot of the Ct values obtained is shown in Suppl. Inf. Fig. 2.

To identify the most stable reference genes, we used the RefFinder software (http://fulxie.0fees.us), which integrates the currently available major computational programs—geNorm (Vandesompele et al, 2002), Normfinder (Andersen et al., 2004), BestKeeper (Pfaffl et al., 2004), and the comparative Δ Ct method (Silver et al, 2006) to compare, rank and assign an appropriate weight to an individual gene and calculate the 260 geometric mean of their weights for overall final ranking. The recommended 261 comprehensive rankings are shown in Suppl. Inf. Table 2. *Nono*, *Hprt1* and *GusB*, 262 identified as the most stably expressed genes, were included in qRT-PCR experiments.

263 3.3.2. Quantification of Cyp19a transcripts in the ovaries of BPA-treated zebrafish.

We analyzed the changes in the levels of transcripts for the aromatase *Cyp19a* gene in the ovaries of BPA-treated zebrafish by real-time qRT-PCR. The relative quantification of *Cyp19a* mRNA is shown in Fig. 4 using *GusB*, *Nono* or *Hprt1* as internal reference genes. The lowest variations were obtained when the aromatase Ct values were normalized against *GusB*, although no significant differences were observed among any of the three reference genes. The results confirm that *Cyp19a* is a detectable but non-abundant transcript in the *D. rerio* ovary.

The expression of Cyp19a mRNA in the ovaries of BPA-exposed fish was significantly 271 272 altered, following a possible non-monotonic curve, where the slope sign changed from negative to positive and back to negative as the dose increased (Fig. 4). The lowest BPA 273 dose (1 µg/L) caused a 3.2-fold decrease in Cyp19a transcript levels. In contrast, the 274 275 levels of this transcript increased significantly (a 1.6-fold increase compared to the control) in zebrafish treated with 10 µg/L but later decrease again with higher doses of 276 BPA in a dose-dependent manner (1.5- and 5-fold in the animals treated with the 100 and 277 1000 µg/L doses, respectively). 278

279 **4. Discussion**

BPA is one of the highest volume chemicals produced worldwide. As with many other endocrine-disrupting chemicals (EDCs), BPA interferes with the body's endocrine system by binding to α - and β -estrogen receptors (ERs) (Kuiper et al., 1998; Metcalfe et al., 2001; Qin et al., 2013) and may produce adverse developmental, reproductive,

neurological, and immune effects in both humans and wildlife. Given its prevalence in 284 285 the environment, identification of BPA exposure biomarkers is urgent for predicting its detrimental effects. The toxicological endpoints of this work in zebrafish exposed to BPA 286 were selected at different levels of biological organization and included histological 287 evaluation of the ovaries, determination of whole-body VTG concentration and analysis 288 of the ovarian aromatase Cyp19a mRNA abundance. We chose these parameters for the 289 290 following reasons: (i) environmental chemical exposure may result in changes in the histological structure of cells, and consequently, histopathological analysis is considered 291 292 an important biomarker in evaluating the toxicological pathology of different compounds 293 in fish; (ii) VTG has been used as a biomarker of exposure to (anti)estrogenic compounds in a number of in vivo and in vitro studies with fish (i.e., Rankouhi, 2002; Navas, 2006; 294 295 Sun, 2010); and (iii) it has been reported that exposure to low doses of BPA causes a 296 decrease in 17β-estradiol (E2) serum concentration linked to ovarian aromatase downregulation (Lee et al, 2013). 297

298 The primary route of BPA contamination is through ingestion. However, the transdermal 299 route could also contribute to BPA exposure when direct contact with BPA occurs. Hence, 300 this work exposed female zebrafish to different concentrations of BPA (1, 10, 100 and 301 1000 μ g/L) introduced into the water of the aquariums by using a continuous flow system. Doses were chosen after reviewing the literature, as they have been reported by previous 302 303 studies in diverse fish species (Ishibashi et al., 2005; Mandich et al., 2007; Villeneuve et al., 2012; Molina et al., 2013). Doses under 200 ng/mL (< 1 uM) are considered low-dose 304 305 for aquatic organisms (vom Saal, 2006) but still have important physiological effects 306 (Inagaki, 2016). Because a non-monotonic dose-response curve has been previously described for BPA exposure, we decided to evaluate the effect of a 5-fold higher dose 307 (1000 μ g/L) over the toxicological endpoints addressed in this work. No differences in 308

mortality, body weight or size were observed in the animals used in this study, in keeping
with previously reported data (Mandich et al., 2007; Hatef et al., 2012; Villaneuve et al.,
2012; Molina et al., 2013).

312 Histopathological analysis of BPA-exposed zebrafish ovaries revealed an increase in the number of atretic follicles as BPA concentration rose (Fig. 1). Previtellogenic oocyte 313 atresia has reportedly affected approximately 10% of female carps exposed to 314 environmental concentrations (1 and 10 µg/L) of BPA (Mandich et al., 2007). These data 315 were partially consistent with those presented here, as they indicated that BPA exposure 316 did not equally affect all types of the follicle. The proportion of atretic *Fovg* (primordial) 317 318 follicles was less than 10% at lower concentrations (1 and 10 μ g/L) but increased to levels over 10% at higher concentrations (100 and 1000 μ g/L) (Fig. 2). In fact, more than 10% 319 of the Fovg, Fc (alveolar cortical), Fv (vitellogenic) and Fm (mature) follicles were atretic 320 in the study groups exposed to higher concentrations of BPA (100 and 1000 μ g/L), 321 showing a dose-dependent increase in the number of atretic follicles (Fig. 2). The fact that 322 323 the Fc showed significant differences (P < 0.05) with respect to the Fovg in all study 324 groups could indicate that BPA disrupted follicular development at the primary stages. Currently, there is no study that has evaluated this hypothesis. Moustafa et al. (2016) 325 326 reported follicles degeneration, congestion blood vessels, and hemorrhage in ovaries, after rats exposure to BPA higher doses in a two generational study. The onset of 327 328 follicular atresia in response to exposure to different concentrations of BPA has been studied in depth (Weber et al., 2003; Wolf et al., 2004; Mandich et al., 2007; Molina et 329 330 al., 2013; Moustafa et al., 2016), but none of these studies addressed which follicular 331 population was the most affected in response to BPA exposure dose. The proportion of atretic follicles increased in the groups that were exposed to higher BPA concentrations 332

333 (100 and 1000 μ g/L) compared to that in the control group (*P*< 0.05), which suggests that 334 BPA affects reproductive function at these doses.

Another biomarker that was evaluated during this study was VTG, which has been widely 335 336 used as an endpoint for estrogen exposure in fish (Orn et al., 2003; Spano et al., 2004; Mandich et al., 2007). VTG is secreted by hepatocytes in the liver. In mature females, 337 VTG is generally synthesized in response to endogenous estrogens, such as 17β-estradiol 338 (E2), which are released into the bloodstream and transported to the ovary, where they 339 are stored in the developing oocytes (Van der Belt et al., 2003; Matozzo et al., 2008). The 340 animals from the control group exhibited whole-body VTG levels of $\approx 20 \ \mu g/g$ (Fig. 2). 341 342 After two weeks of exposure, VTG concentrations in all study groups increased (P < 0.05) in a dose-dependent manner, reaching values 500-fold higher at the highest BPA dose 343 tested. This finding could confirm the estrogenic potential of BPA, as previous reports 344 have shown a dose-dependent increase in VTG production (Mandich et al., 2007; Naderi 345 et al., 2014). Even the group exposed to the lowest concentration of BPA (1 μ g/L) showed 346 347 a significant (P < 0.05) 2-fold increase in VTG concentration after 14 days of exposure. 348 This result indicates that zebrafish respond to low BPA concentrations by synthesizing VTG (Mandich et al., 2007; Naderi et al., 2014) and suggests that VTG is a very sensitive 349 350 biomarker of BPA exposure, even at environmentally relevant BPA concentrations.

The observed increase in follicular atresia and in VTG synthesis depending on BPA concentration, suggests that estrogenic action of BPA is mediated by an abnormal synthesis of estrogen and VTG by the follicles in a dose-dependent manner. The increase in VTG concentration in BPA-exposed zebrafish ovaries might be related to the significant increment of Fv with respect to the control group that we reported in a previous work (Molina et al., 2013).

The most accepted hypothesis states that EDCs interfere with steroid hormone action by 357 358 disrupting steroid biosynthesis with consequences for downstream signaling pathways (López-Casas et al., 2012). In the ovary, androgens synthesized in theca cells are 359 converted into estrogens by ovarian aromatase CYP19a in the local granulosa cells 360 (Cheshenko et al., 2008). Endocrine disruptors have challenged the toxicological 361 assumptions related to dose-response relationships. Non-linear relationships between the 362 363 dose and the response have been observed for certain EDCs. Using a quantitative RT-PCR assay, the kinetics of Cyp19a transcript abundance variation was investigated in 364 zebrafish ovaries after BPA exposure. Data in Fig. 4 show that Cyp19a transcript 365 366 abundance was significantly diminished (P < 0.001) in response to exposure at the lowest BPA concentration (1 μ g/L), a dose that produced no important histological changes or 367 VTG increases. The peak abundance of Cyp19a mRNA was detected at the dose of 1 368 369 μ g/L BPA and decreased afterward. This type of non-monotonic dose-response curve has been reported for several endocrine-disrupting compounds, including BPA and many 370 371 other hormonally active compounds, and has been attributed to multiple (yet undetermined) mechanisms stimulated when the assimilated dose is low but able to inhibit 372 373 toxic effects of ulterior higher doses of the compound (see Myers et al., 2009; Vandenberg 374 et al., 2009, for reviews; Renieri et al., 2017). However, it must be taken in consideration that the apparent non-monotonic dose response is based on a single low dose, and testing 375 more low doses is needed to confirm this trend. Data presented in this study corroborated 376 that measurement of transcriptional profiles variations are very suitable early exposure 377 biomarkers, as transcription is affected prior to protein synthesis. These results are in 378 agreement with previous reports indicating changes in Cyp19a expression by even very 379 low concentrations of EDCs (Flint et al., 2012). However, in our experimental model, the 380 decrease in Cyp19a transcripts at the 1 µg /L BPA dose had no effect on histological 381

organization of the ovary (Fig. 1). Only when exposed to 100 μ g/L BPA or higher doses did the ovaries show significant histological changes with significant increases in the follicular atresia (over 10% compared to the control group, Molina et al., 2013). This result coincides with the inhibition of *Cyp19a* expression, which was accentuated in response to exposure to higher BPA concentrations (100 and 1000 μ g/L).

Although the precise cellular and biochemical mechanism(s) underlying these changes in 387 388 Cyp19a transcript levels after BPA exposure are currently unknown, low BPA doses presumably exert a direct negative action on the transcriptional regulation of Cyp19a 389 expression in granulosa cells. This effect would lead to a reduced number of Cyp9a 390 391 mRNA molecules but also to the onset of a stress response that stimulates Cyp19a transcript synthesis. In accord with Lee et al. (2013), the results presented in this study 392 suggest that the low-dose BPA exposure in zebrafish females initially reduces E2 393 synthesis and thus disrupts ovarian steroidogenesis by inhibiting Cvp19a. BPA has been 394 395 reported to regulate the expression of the Aryl hydrocarbon receptor (AhR) mRNA 396 (Kharrazian, 2014), a key transcription factor (TF) in the xenobiotic response that 397 activates the transcription of a battery of genes, including Cyp19a (Baba et al., 2005). Similar to aromatase, VTG synthesis is induced by estrogen-dependent stimulation of 398 399 gene expression via AhR cross-talk. However, the estrogenic potency of BPA is significantly increased by bioactivation, depending on the microsomal and cytosolic 400 401 constituents. It is likely, in contrast with the direct BPA inhibition of Cyp19a expression, that BPA doses over 1 µg/L are required to induce the production of biotransformation 402 403 enzymes and significantly increase the VTG levels in the zebrafish ovary.

404

405 **5.** Conclusions

We observed a good correlation between BPA exposure and the possibility of 406 407 reproductive dysfunction. The results of this work indicate that aromatase Cyp19a mRNA quantifications, whole-body VTG measurements and morphological endpoints are 408 sufficiently sensitive biomarkers to identify the precocious effects of environmental 409 concentrations of BPA on the ovaries after two weeks of BPA exposure. In accord with 410 previous studies demonstrating the utility of the aromatase as a biomarker in rats, we 411 412 show the relevance of determining the changes in transcriptional profiles of the Cyp19a gene in the zebrafish ovary as an early biomarker of BPA exposure. Additionally, we 413 observed that variations in the VTG protein are also indicative of endocrine alterations. 414 415 Both Cyp19a and VTG are more sensitive biomarkers than histological evaluations, as their abundance changes at lower doses than those required to produce morphological 416 alterations. Data suggest that low BPA doses exert a direct negative action on the Cyp19a 417 418 expression, but higher doses are needed to observe BPA acting as a positive inductor of VTG synthesis. 419

420

421 Acknowledgments

The authors wish to thank the Spanish Junta de Andalucía (P09-AGR-514) for theirfinancial support for this study.

The authors wish to thank the Institut für Pathologie Stiftung TierärztlicheHochschule Hannover (Germany), especially Dr. Wolfgang Baumgärtner and the technician operating the electron microscope, for their encouragement during the ultrastructural images evaluation.

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588 **Figure captions**

Figure 1: BPA-exposure-induced histological changes in zebrafish ovaries. Images 589 590 obtained under light microscope (left) are marked as cA (control), A, C, E, G (groups treated with increasing doses of BPA); for these images, bars represent 50 µm. 591 Ultrastructural observations (right) are marked as cB (control), B, D, F, H (exposure 592 593 groups); in these images, bars represent 10 µm. Control group: Sections shown all types of ovarian follicles (FO) (cA) interstitial tissue and ovg follicles (Fovg) (cB) with their 594 normal appearance. Group 1 $\mu g/L$: A) Vacuolizations (V) are visible between the oocyte 595 and the follicular cells (Fovg). B) Detachment of an ovg follicle (Fovg) from the follicular 596 cells of the oocyte. Group 10 µg/L: C) Vacuolizations (V) of the pellucid zone (ZP) can 597 be seen in the Fc, particularly follicular atresia (FA), with the disintegration of all its 598 components. D) Vacuolizations (V) were observed in the pellucid zone (ZP) of the Fc. 599 Group 100 µg/L: E) Strong vacuolizations (V) and separations of the pellucid zone (ZP) 600 601 were observed in the Fv, with evident follicular atresia (FA). F) Disorganization and vacuolization (circle) of the dense and light pellucid zone (ZP) of the Fv. Group 1000 602 $\mu g/L$: G) Follicular atresia (FA) was observed in the Fm, along with the production of 603 604 cellular debris (RC). H) Follicular atresia of Fm, with the disintegration of cellular components and production of cellular debris (RC). 605

Figure 2. Follicular atresia caused by BPA exposure. The percentage (numbers inside
the bars) of atretic follicles in the different follicular populations in each study group is
represented. Fm: mature follicles; Fv: vitellogenic follicles; Fc: cortical alveolar follicles;
Fovg: primordial follicles.

610 Figure 3. VTG levels in the whole-body extracts of zebrafish exposed to increasing

611 concentrations of BPA. The vitellogenin (VTG) concentrations ($\mu g/g$) were determined

612 in whole-body extracts of zebrafish (n = 9 per experimental group) exposed to increasing 613 amounts of BPA using a specific commercial enzyme-linked immunosorbent assay. The 614 data are presented as the means \pm SD of 5 independent determinations. All treatment 615 groups were significantly different from the control at the *P* < 0.001 level; significant 616 differences at the *P* ≤ 0.05 level were also observed for the comparisons between each 617 two consecutive treatment doses.

Figure 4. Changes in the number of *Cyp19a* transcripts in the zebrafish ovary after 618 different concentrations BPA exposure. The relative quantification of the levels of the 619 *Cyp19a* transcripts was calculated by using the $2^{-\Delta\Delta Ct}$ method and the *GusB*, *Nono* or 620 621 *Hprt1* transcripts as internal references. Numbers over the bars indicate the fold change 622 in relation to the control counts. The significant differences between the control and experimental samples were determined using Student's *t*-test and were expressed by ** 623 to denote P < 0.01; and ***, P < 0.001. The statistics program SigmaStat 5.1 (GraphPad 624 Prism) was used throughout the study. 625

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