



## Assessment of multiple hormone activities of a UV-filter (octocrylene) in zebrafish (*Danio rerio*)



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### HIGHLIGHTS

- Octocrylene (OCT) can accumulate in fishes up to sufficiently high levels to cause adverse effects on the endocrine system.
- The accelerated ovary development indicates that OCT has the effect on sex-endocrinology.
- The gene alterations address that OCT has the multiple hormone activities.

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### ABSTRACT

In this study, zebrafish (*Danio rerio*) were exposed to a UV-filter-octocrylene (OCT) with elevated concentrations for 28 d. The total body accumulation of OCT in zebrafish was found to reach 2321.01 (“L” level), 31,234.80 (“M” level), and 70,593.38 ng g<sup>-1</sup> (“H” level) when the average OCT exposure concentration was controlled at 28.61, 505.62, and 1248.70 µg L<sup>-1</sup>, respectively. Gross and histological observations as well as RT-qPCR analysis were conducted to determine the effects of OCT accumulation on zebrafish. After exposure, the gonad-somatic index and percentage of vitellogenic oocytes were found to increase significantly in the ovaries of female zebrafish at the H accumulation level. Significant up-regulation of *esr1* and *cyp19b* were observed in the gonads, as well as *vtg1* in the livers for both female and male zebrafish. At M and H accumulation levels, apparent down-regulation of *ar* was observed in the ovaries and testis of the female and male zebrafish, respectively. Although the extent of the effects on zebrafish differed at different accumulation levels, the induction of *vtg1* and histological changes in the ovaries are indications of estrogenic activity and the inhibition of *esr1* and *ar* showed antiestrogenic and antiandrogenic activity, respectively. Thus, as OCT could easily accumulate in aquatic life such as zebrafish, one of its most of concern hazards would be the disturbance of the histological development and its multiple hormonal activities.

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### 1. Introduction

By virtue of the increased public awareness of the hazards associated with overexposure to ultraviolet (UV) radiation, UV filters are now commonly added to products such as creams, lipsticks,

and in the UV-protection of numerous materials and products (Balmer et al., 2005; Fent et al., 2010). It is estimated that about 10,000 tons of UV filters are produced annually for the global market (Danovaro et al., 2008). There are two main categories of UV filters: 1) Inorganic filters composed of titanium dioxide and/or zinc oxide for scattering and reflecting UV light, and 2) organic compounds such as 2-ethyl-hexyl-4-trimethoxycinnamate (EHMC) or benzophenone-3 (BP-3), which work by absorbing UV-light. In the European Union, 28 UV filters have been registered in total (Schlumpf et al., 2008). All have been identified as potentially dangerous pollutants to the aquatic environment.

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UV filters can easily enter the aquatic system via direct or indirect pathway such as being washed off the skin during bathing and swimming. Because the molecules of most UV filters are very stable, it is difficult for them to be completely removed by conventional wastewater treatment processes. The treated effluents may thus be the primary source of UV filters entering the aquatic environment (Buser et al., 2006; Fent et al., 2010). High concentrations of EHMC (up to 20,070 ng L<sup>-1</sup>) and 4-methylbenzylidene camphor (up to 960 ng L<sup>-1</sup>) have been measured from domestic wastewaters (Kupper et al., 2006), and, if they are not sufficiently removed by treatment, may significantly contaminate the receiving water bodies due to effluent discharge. In surface waters, benzophenone-4 (BP-4) has been measured at levels as high as 849 ng L<sup>-1</sup> (Rodil et al., 2008). EHMC was measured at 26–5610 ng L<sup>-1</sup> in the source water for drinking water supply (Loraine and Pettigrove, 2006).

Of the most commonly used UV filters, octocrylene (OCT) has attracted significant research attention in the environmental protection field because it is more refractory and hydrophobic than other UV filters (Rodil et al., 2009). OCT has been found in several different water bodies across the globe. Seawater collected from Hong Kong, for example, can be contaminated with OCT at up to 103–6812 ng L<sup>-1</sup> (Tsui et al., 2015). It has also been reported that the concentration of OCT in lake water ranges from 2 to 27 ng L<sup>-1</sup> (Poiger et al., 2004). OCT has even been detected in tap water at levels as high as 170 ng L<sup>-1</sup> (Díaz-Cruz et al., 2012). Authors have also monitored the presence of OCT in the effluent of domestic wastewater treatment plant to find that its concentration can reach 0.56–1.8 µg L<sup>-1</sup> (unpublished data).

The Kow of OCT is as high as about 10<sup>7</sup>, which allows it to easily accumulate in aquatic life. In a Swiss lake, OCT was detected from the mussel of brown trout as 2400 ng g<sup>-1</sup> lipid weight (l.w.) (Buser et al., 2006). In a remote Brazilian coastal area, the accumulated OCT in Franciscana dolphins was as high as 89–782 ng g<sup>-1</sup> (l.w.) (Gago-Ferrero et al., 2013). OCT accumulations of 25–11,875 ng g<sup>-1</sup> body weight (b.w.) were also reported in the liver of cod in Oslofjord (Langford et al., 2015). Unfortunately, relatively little information is available regarding the toxic effect of OCT accumulation in aquatic organisms.

There has been growing concern regarding the endocrine-disrupting effect of UV filters on aquatic life, for example, the alterations in gonad histology of mature fathead minnows caused by benzophenone-2 (BP-2) (Weisbrod et al., 2007). Many UV filters have been found to be hormonally active and show agonistic and/or antagonistic effects towards nuclear receptors, estrogen responsive genes, and steroidogenesis (Zucchi et al., 2010; Christen et al., 2011; Kim et al., 2014). Accordingly, any study on the potential risk associated with OCT may also need to be concentrated on its possible endocrine disrupting effect when it is accumulated in the bodies of aquatic animals. Zebrafish (*Danio rerio*) makes an excellent vertebrate model for assessing the toxicity of this manner of chemicals *in vivo*, especially when analyzing the action mode (Reimers et al., 2004). Previous studies have also shown that in order to evaluate the ultimate toxicological effect of a target chemical on aquatic life, experimental conditions should be established appropriately for the chemical to be accumulated in the test aquatic animal to appreciably high levels during a certain duration of exposure (Zucchi et al., 2010; Blüthgen et al., 2012). In the case of zebrafish, OECD guidelines specify that 28 d is a reasonable exposure time (OECD, 1996) to control target chemical in the experimental solution to sufficiently elevated concentration.

The objective of this study was to gain knowledge on the endocrine disrupting potential of OCT on aquatic life by using zebrafish for *in vivo* bioassay. We conducted gross and histological observations as well as RT-qPCR analysis to monitor the multiple

hormone activities of OCT after the chemical was accumulated to a certain level.

## 2. Materials and methods

### 2.1. Chemicals

2-Ethylhexyl-2-cyano-3,3-diphenylpropenoate (OCT, CAS No:6197-30-4, purity ≥ 95%) was purchased from TCI (Tokyo, Japan). Dimethylsulfoxide (DMSO), methanol, and dichloromethane of HPLC grade were purchased from Fisher Scientific (Shanghai, China). Stock solutions of OCT (1, 10, 30 g L<sup>-1</sup>) were prepared by dissolution in DMSO and stored in a dark environment at 4 °C between uses. Before exposure, the stock solutions were diluted and DMSO solvent concentration maintained at ≤ 0.01% (v/v).

### 2.2. Maintenance of zebrafish

Juvenile zebrafish (about five months old, mean body length 3.22 cm, mean body weight 0.28 g) was obtained from an animal lab (Hebei, Shijiazhuang) and transferred to an ultra-white fish tank. Before conducting the experiment, female and male zebrafish was separately acclimatized for one month in reconstituted tap water with a total hardness of 125 mg L<sup>-1</sup> as CaCO<sub>3</sub> and an electrical conductivity of 270 µS cm<sup>-1</sup>. The water temperature was kept constant at 27 ± 1 °C with a photoperiod set to 14:10 h light/dark. Zebrafish was fed twice daily with a combination of brine shrimp and flake fish food. Water parameters (nitrate, nitrite, pH) were measured daily and oxygen saturation was maintained at ≥80%.

### 2.3. Exposure of zebrafish to OCT

Adult zebrafish was exposed to OCT solutions including a blank control, solvent control (DMSO, ≤0.01%), and a series of prepared OCT solution (100, 1,000, 3000 µg L<sup>-1</sup>) for 28 d. The concentrations and duration of exposure were determined based on previous studies on other UV filters (Christen et al., 2011) and OECD guidelines (OECD, 1996). A static-renewal procedure was conducted during exposure. Zebrafish of similar size was removed from the culture tank randomly and placed into 5 L glass beakers covered with gauze. Every 48 h, zebrafish was transferred to new exposure solutions and the surplus food and faeces were immediately removed by siphoning. Throughout the entire exposure period, zebrafish was fed as previously described. Their mortality, development, and abnormal behaviors were recorded daily.

### 2.4. Analysis of OCT in exposure water and zebrafish

To determine the actual OCT concentrations during exposure, aliquots of 200 mL exposure water from each treatment group were taken at the beginning (0 h), after 24 h, and prior to water renewal (48 h). The adsorption and accumulation of OCT in the zebrafish may not have been consistent during the whole exposure, resulting in variations in OCT level in the water; thus, in order to accurately obtain actual OCT concentrations, samples were taken on days 1–3, 10–12, and 19–21, respectively.

About 200 mL aliquots of each nominal concentration were taken for analysis at 0 h, 24 h, and 48 h. The water samples were stored at –20 °C in brown glass bottles until analysis. Watersample extraction and chemical analysis were performed as follows. Before solid-phase-extraction, the pH of exposure samples was adjusted to 3 with hydrochloric acid. Agilent C18 (500 mg and 3 cc) cartridges were conditioned with 10 mL dichloromethane, 10 mL methanol, and 10 mL Milli-Q water, then water samples were passed through

at a rate of 5 mL/min. The cartridges were washed with 10 mL Milli-Q water and air dried for about 30 min, then the retained analyte was eluted with 10 mL dichloromethane. The extraction was evaporated to dryness under a gentle stream of N<sub>2</sub> at 30 °C, reconstituted in 1 mL methanol, and stored in the dark at 4 °C until analysis. Chemical analysis of OCT concentration was carried out by high performance liquid chromatography (HPLC) coupled to UV-detection at 290 nm. The chromatographic mobile phase composed of 85% (v/v) methanol in Milli-Q water was subjected to a flow rate of 0.9 mL/min. The chromatographic separation was performed with a Hypersil BDS-C18 column (4.6 mm × 250 mm, 5 μm particle size) at a temperature of 30 °C. Extract samples 20 μL in volume were injected in duplicate and finished in 12 min with an auto-sampler.

Zebrafish was extracted according to previously published methods (Weisbrod et al., 2007) with some modifications. Individual whole zebrafish was homogenized in a tissue grinder (Sangon, EQ 6002) together with 3 mL HPLC water (the same quantity as fish weight). Liquid-liquid extraction was then performed with a mixture of ethyl acetate, *n*-heptane, and HPLC water (1:1:1, v/v/v). The volume of each eluent added was proportional to the initial amount of HPLC water. The mixture was rigorously shaken for 10 min; then the supernatant was collected after centrifugation at 5000 g for 10 min at 4 °C. The extraction was repeated twice. The supernatants were combined and filtered through 0.45 μm Acrodisc® syringe filter, then the extraction was dried with a gentle stream of nitrogen at 30 °C and the analytes were reconstituted in 200 μL methanol. Finally, the residues were centrifuged for 10 min at room temperature and the final supernatant was subjected to HPLC analysis. The detection of OCT in exposed zebrafish was similar to that of water samples. At the end of the exposure period, the gonads, livers, and brains were excised, snap-frozen in liquid nitrogen, and stored at –80 °C until qRT-PCR analysis.

## 2.5. Gross and histological observations

To assess the effect of OCT on the development of zebrafish after exposure, the condition factor ( $CF = (\text{weight (g)}/\text{length (cm)} \times 100)$ ) was measured as well as the gonad-somatic index ( $GSI = \text{gonad weight (g)}/\text{body weight (g)}$ ) according to methods used by previous researchers (Weisbrod et al., 2007; Christen et al., 2011). At the end of exposure, the ovaries of five female zebrafish were excised and encased in O.C.T. (Sakura, USA). Sections (10 μm) were cut with a CM1850 Freezing Microtome (Leica Biosystems, Germany) and stained with haematoxylin and eosin (H&E) and pictures of three randomly selected fields of vision were taken with an optical microscope (Nikon 90i) to observe the histology of the

ovaries of the female zebrafish. Three fields from each region of the ovary were randomly selected to observe the distribution of primary oocytes, previtellogenic oocytes, and vitellogenic oocytes in the ovaries. The frequencies of the above cells were determined according to Pereira et al. (2013).

## 2.6. RT-qPCR analysis

Total RNA was extracted from the brains, livers, and gonads ( $n = 5$ ) and pooled using a Takara MiniBest Universal RNA Extraction kit (Takara, Japan) according to the manufacturer's instructions. To guarantee the concentration and quality of RNA, RNA was analyzed with a NanoDrop 1000 spectrophotometer and electrophoresis. Only RNA samples of purity between 1.8 and 2.0 for both ratios of 260/280 nm and 260/230 nm and RNA intensity ratio for 28S/18S of about 2:1 were used for RT-qPCR.

Total RNA templates were reverse transcribed to cDNA and stored at –20 °C for subsequent RT-qPCR analysis. The three-step real-time PCR profile was completed according to previously published methods (Christen et al., 2011). In this study, certain target genes involved in hormonal activity (Table 1) were selected for analysis: estrogen responsive gene (*vtg1*), nuclear receptor (*esr1*, *ar*), and steroid metabolism (*cyp19b*, *rpb2a*). The 18S ribosomal RNA (18S *rRNA*) was selected as a reference gene for normalization, because its expression profile did not vary either in different conditions or different tissues. The relative linear quantity of target molecules relative to the calibrator was calculated by  $2^{-\Delta\Delta ct}$  (Livak and Schmittgen, 2001). Transcriptional alterations of different genes are expressed as fold change ( $\log_2$ ).

## 2.7. Data analysis and statistics

Data were illustrated graphically as shown below with GraphPad® Prism5 (GraphPad Software, San Diego, CA, USA). Significant differences in transcript levels were analyzed by one-way analysis of variance (ANOVA). Differences between treatments were analyzed by Newman-Keuls test to compare treatment means with controls. Results are given as mean ± standard deviation (SD). Differences were considered statistically significant at  $p \leq 0.05$ .

## 3. Results and discussion

### 3.1. Chemical analysis

Samples were analyzed by HPLC at 0 h, 24 h, and 48 h to determine the actual concentration of OCT in the prepared solution (Fig. 1). All actual concentrations were lower than the

**Table 1**

Primer sequences for quantitative real-time PCR analysis and sources: *vtg1* (vitellogenin 1), *esr1* (estrogen receptor), *ar* (androgen receptor), *cyp19b* (P450aromB), *rpb2a* (retinol-binding protein type II), 18S *rRNA* (18S ribosomal RNA).

Target gene	GenBank number	Sense primer (5'-3')	Antisense primer (5'-3')	Product size (bp)
<i>vtg1</i> <sup>a</sup>	AY034146	AGCTGCTGAGAGGCTTGTTA	GTCCAGGATTTCCCTCAGT	94
<i>esr1</i> <sup>b</sup>	NM_152959	TGAGCAACAAAGGAATGGAG	GTGGGTGTAGATGGAGGTTT	163
<i>ar</i> <sup>c</sup>	NM_001083123	CACTACGGAGCCCTCACTTGC	GCCCTGAAGTGTCCGACCTC	237
<i>cyp19b</i> <sup>d</sup>	AF183908	CGACAGGCCATCAATAACA	CGTCCACAGACAGCTCATC	94
<i>rpb2a</i> <sup>e</sup>	AF363957	GGAGATGCTCAGCAATGACA	TCTGCACAATGACCTTCGTC	110
18S <i>rRNA</i> <sup>f</sup>	Y855349.1	AAACGGCTACCACATCCAAG	TTACAGGGCTCGAAAGAGA	116

Date sources:

<sup>a</sup> (Hoffmann et al., 2006).

<sup>b</sup> (Martyniuk et al., 2007).

<sup>c</sup> (Hossain et al., 2008).

<sup>d</sup> (Arukwe et al., 2008).

<sup>e</sup> (Zucchi et al., 2011).

<sup>f</sup> (Wintz et al., 2006).

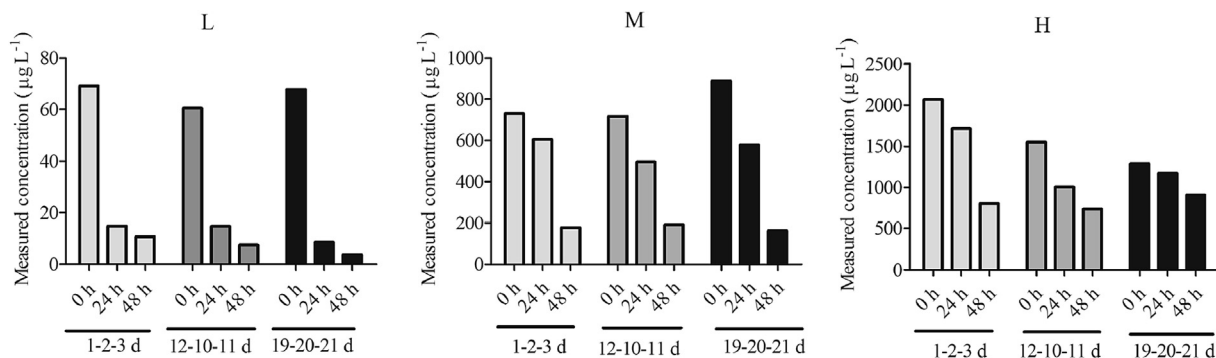


Fig. 1. Measured concentrations of OCT at prepared concentrations of 100, 1,000, and 3000  $\mu\text{g L}^{-1}$ .

corresponding prepared levels. The actual concentration of OCT markedly decreased with the extension of exposure time during each water renewal. The mean actual exposure concentration within 48 h was 28.61, 505.62, and 1248.70  $\mu\text{g L}^{-1}$  at the prepared level of 100, 1,000, and 3000  $\mu\text{g L}^{-1}$ , respectively. Based on the results of our preliminary experiment on the physicochemical properties of OCT, there was no significant change in OCT solubility in water over 72 h (unpublished data), which ruled out the degradation of OCT during exposure. Additionally, the data obtained from the exposure solution without zebrafish showed no further significant decrease in OCT. To this effect, we hypothesized that the decrease in OCT in the exposure water was caused by several factors including uptake into zebrafish or adsorption to surplus food or faeces rather than degradation.

OCT was also detected in all exposed zebrafish except the control. To be precise, the total body burden (TBB) of OCT as shown in Fig. 2 reached up to 2321.01 (L), 31,234.70 (M), and 70,593.38  $\text{ng g}^{-1}$  (b.w.) (H). By comparing the TBB of OCT in zebrafish in the present study to that observed in previous studies, we confirmed that the residual concentrations of OCT in zebrafish were within the scope of the concentrations in the fishes living in surface water over long periods of time. For example, the OCT content can reach up to 2400  $\text{ng g}^{-1}$  (l.w.) in brown trout (Buser et al., 2006) and 1,1875  $\text{ng g}^{-1}$  (b.w.) in cod (liver) (Langford et al., 2015).

The bioaccumulation factors (BCF) of OCT in zebrafish at the accumulation levels of “L”, “M”, and “H” were 81, 62 and 56, respectively (Fig. 2) – a similar but lower range compared to the TBB. In general, without any adverse effect of the OCT exposure, the

enrichment of OCT in zebrafish can be expected to equilibrate eventually, and the BCF to become constant. In this study, OCT displayed a similar bioaccumulation profile with another quickly metabolized UV filter, BP-3 (19–94) (Blüthgen et al., 2012). Gomez et al. (2012) pointed out that the uptake of OCT is rapid, but is followed by elimination within 24 h. OCT is also known to be excreted to some extent (Gago-Ferrero et al., 2015). Thus, the metabolism and/or excretion of OCT *in vivo* can explain the low BCF despite the high lipophilicity of OCT.

### 3.2. Gross observation

During exposure, no abnormal behavior or mortality were observed among the exposed zebrafish. Furthermore, no significant differences in CF occurred in the OCT-exposed zebrafish compared to the control (Fig. 3A and B). The GSI of males did not change significantly (Fig. 3C). However, the H accumulation level showed a significant increase in GSI in the ovaries of females (Fig. 3D).

GSI is a highly comprehensive physiological index that reflects the level of sex hormones and reveals the impact of exogenous estrogen to fishes (Ankley et al., 2001). Diuron metabolites lead to significant estrogenic activity associated with ovary development with the increase of GSI in *O. Niloticus* females (Boscolo Pereira et al., 2016), for example. The results of the present study indicate that OCT has estrogenic activity effects once accumulated in female zebrafish to a certain degree.

### 3.3. Ovary histology

Based on the GSI changes in females, we further investigated the histological development of ovaries in the exposure solution. Ovarian sections of zebrafish in the blank (water) showed a normal frequency distribution at ovarian lamellae, including primary oocytes (PV, with a large nucleus, centrally positioned with numerous nucleoli) previtellogenic oocytes (V, with a large number of cortical alveoli vesicles) and vitellogenic oocytes (Oo, where cortical alveoli vesicles were no longer observed, oocytes were at their maximum size and were filled with protein yolk granules) (Fig. 4A). The ovaries of female zebrafish with H level OCT accumulation mainly revealed heavy vitellogenic oocytes characterized by deposition in the cytoplasm exogenous yolk (Fig. 4C and D). Moreover, quantitative analysis of the oogenesis (Fig. 5) showed a decrease in the percentage of primary oocytes (13%,  $p \leq 0.001$ ), but a significant increase in vitellogenic oocytes (20%,  $p \leq 0.001$ ) of ovaries with the H accumulation level.

Although gonad development is highly complex, external conditions can provide very useful information; sex reversal, for example, may occur if the gonad is exposed to endocrine disruptors

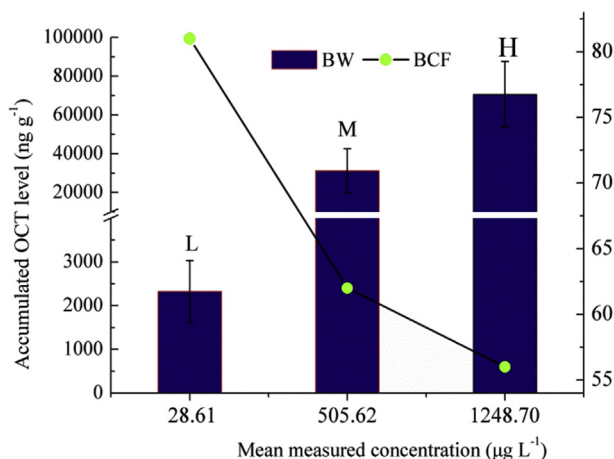
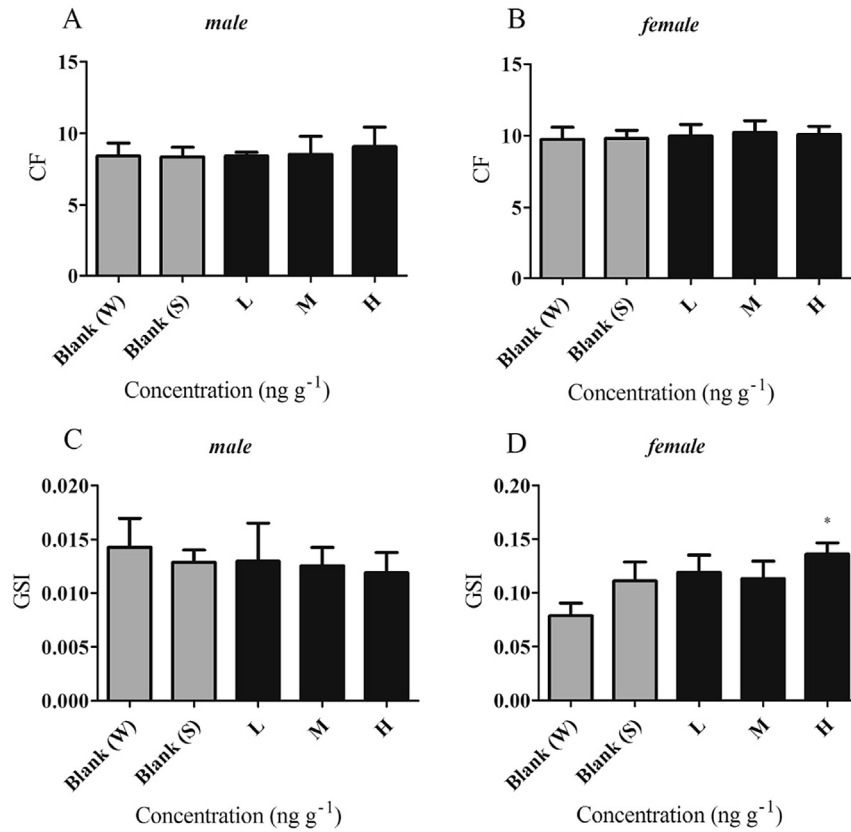


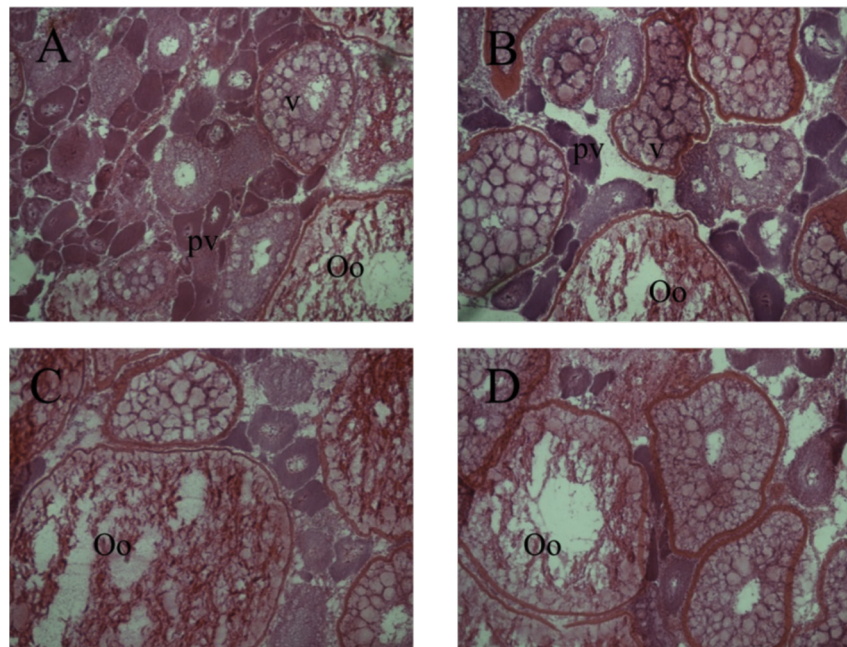
Fig. 2. Accumulated OCT levels and BCF of OCT in zebrafish exposed to actual OCT levels of 28.61, 505.62, and 1248.70  $\mu\text{g L}^{-1}$ .



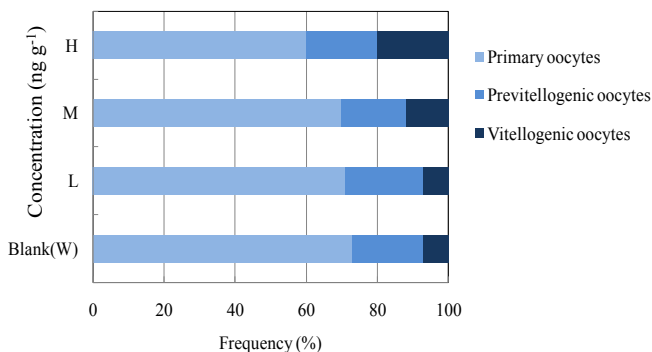
**Fig. 3.** CF and GSI indices of male and female zebrafish exposed to blank (w), blank (s), and zebrafish with OCT accumulation levels of L, M, and H after 28 d exposure. Values are mean  $\pm$  SD. Asterisks (\*) indicate statistical significance (\* $p \leq 0.05$ ).

during critical development periods (Scholz and Gutzeit, 2000; McAllister and Kime, 2003). Nagahama and Yamashita (2008) demonstrated that E2 regulates ovarian development through the

control of vitellogenin synthesis in the liver during the oocyte growth period. Vitellogenic oocytes are also known to significantly increase in size due to the presence of vitellogenin (Lubzens et al.,



**Fig. 4.** Histological section of ovaries of female zebrafish exposed in blank (water) (A), and zebrafish with OCT accumulation level of L (B), M (C), and H (D). PV: previtellogenic oocytes; V: vitellogenic oocytes; Oo: oocytes.



**Fig. 5.** Relative percentage of different stages of oogenesis in ovaries of female zebrafish exposed to blank (water) (A), and zebrafish with L, M, and H OCT accumulation level. Asterisks (\*) indicate statistical significance (\* $p \leq 0.05$ ).

2010). In this study, the increased GSI and percentage of final vitellogenic oocytes in the females with H level accumulation of OCT was likely directly caused by the induction of vitellogenin. This hypothesis was further validated by our subsequent analysis of *vtg1* mRNA expression in the liver. The effect of OCT on the development of ovaries was identical to that of EHMC, whereby a significant increase in vitellogenic oocytes was observed at  $394 \mu\text{g L}^{-1}$  (Christen et al., 2011). Similarly, the UV filter BP-2 has been shown to cause increased follicular atresia and even adverse effects on reproduction (Weisbrod et al., 2007). Taken together, these results indicate that high accumulation of OCT can interfere with the endocrine system by accelerating the ovaries and development of zebrafish.

#### 3.4. Effects of OCT on hormonal activity and steroidogenesis

Our histological observations indicated that high accumulation of OCT may have a significant effect on the endocrine system. In effort to better understand the effects of OCT accumulation on aquatic life, the differential expression of the target genes in different tissues was analyzed. The transcriptions of several genes involved in hormonal pathways were observed in the brains, gonads, and livers (Fig. 6).

The expression pattern of *esr1* in the gonad showed an overall up-regulated tendency. The modulation of this sex-steroid receptor was in line with previous studies on UV filters, where an induction of estrogen receptors (ERs) has been consistently observed (specifically for EHMC) (Inui et al., 2003). There is ample empirical evidence for the fact that the biological impact of exogenous estrogens extend beyond the gonads to other bodily systems, including the brain (Engler-Chiurazzi et al., 2016). An overall down-regulation of *esr1* was observed in the brain in our zebrafish samples.

Interference with ERs is a well-identified key event in the initiation of adverse outcomes (Sonavane et al., 2016). The down-regulation of *esr1* that we observed is related to the anti-estrogenic activity of OCT in the brain, in accordance with results obtained *in vitro* in a previously published recombinant yeast assay (Kunz and Fent, 2006). As a ligand dependant transcription factor, ER transduces hormone signals into a large variety of physiological responses in various organs such as reproductive organs (Ascenzi et al., 2006). Thus, different modulations of OCT on the expression of *esr1* in different organs highlight the importance of analyzing transcription profiles in multiple tissues.

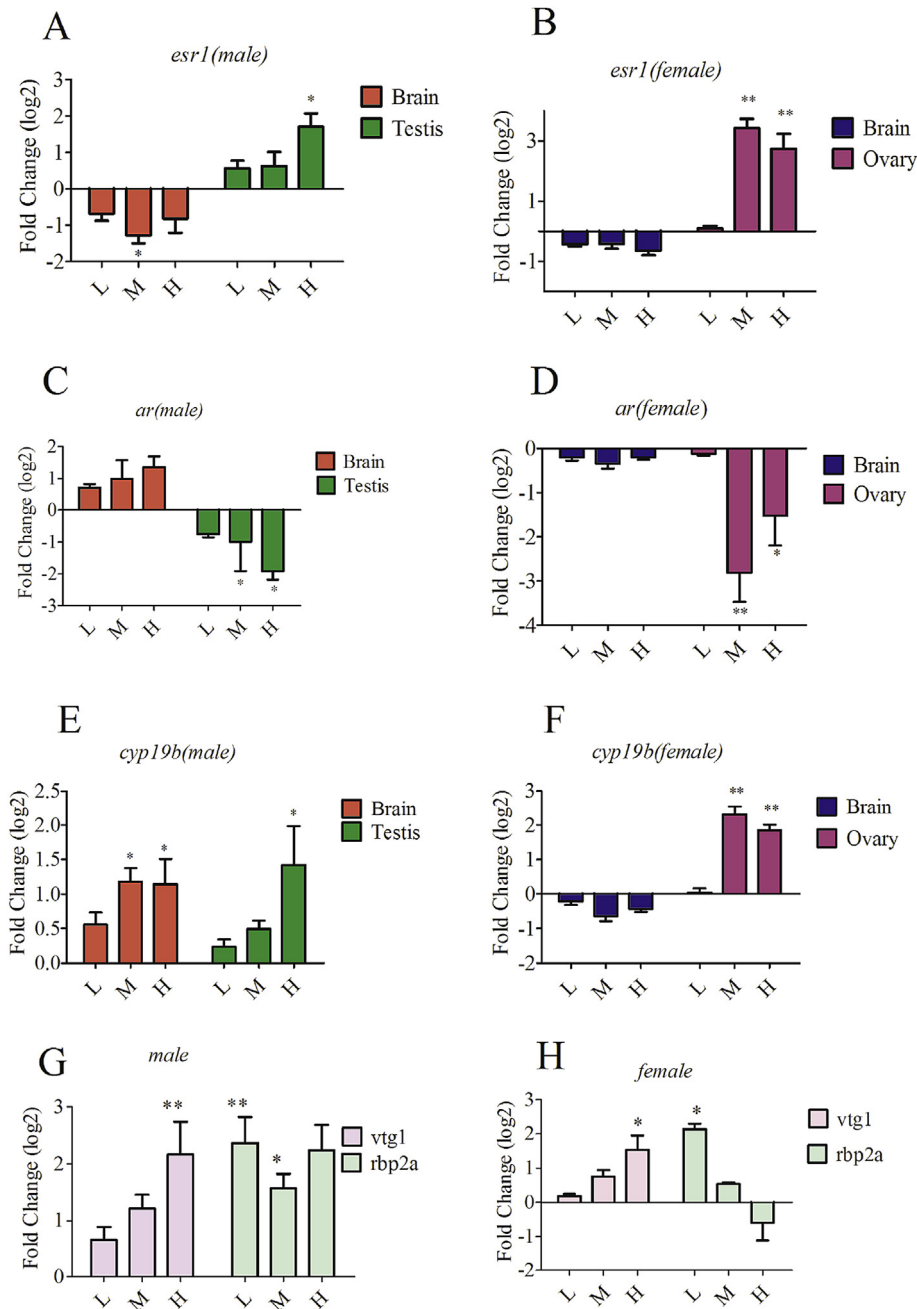
Exogenous estrogen can interfere with the endocrine system via multiple mechanisms of action. A significant inhibition of the expression of *ar* was observed in the gonads of males with M and H OCT accumulation levels and in female zebrafish with the H

accumulation level, again in accordance with previous *in vitro* data showing that OCT is submaximal to full antiandrogenic activity in recombinant yeast carrying *hAR* (Kunz and Fent, 2006). By comparison, the estrogenic *EE2* produced a down-regulation of the *ar* transcript (Filby et al., 2007). In short, antiandrogenic activity of OCT cannot be ruled out *in vivo*. Christen et al. (2011) pointed out that potential changes in transcriptions of hormone receptors may be partially responsible for the histological changes in the testes and ovaries of exposed fishes. Thus, we would assert that the regulation of *esr1* and *ar* in the ovaries partially promotes ovarian development.

The accumulation of OCT slightly affected the expression of genes involved in steroidogenesis. There was a significant up-regulation of *cyp19b* (cytochrome p450 aromatase) in the brain of males with M and H OCT accumulation levels. *Cyp19* is considered a sensitive, convenient signal of organic xenobiotics in the aquatic environment (Hinfray et al., 2006), and estrogens are known to enhance the transcription of *cyp19b* in the brains of adult zebrafish (Mouriec et al., 2009; Callard et al., 2001). Certainly, up-regulation of *cyp19b* in the brain of male zebrafish can be interpreted as an estrogenic response. In addition, *cyp19b* is an important hormone involved in controlling the reproductive process in teleosts (Kazeto et al., 2001; Kuhl et al., 2005). The ovaries of female zebrafish with the M and H OCT accumulation levels showed a significant increase in *cyp19b*, but the gene expression only increased in the testis of male zebrafish at the H accumulation level (Fig. 6E and F). The OCT accumulation-mediated induction of *cyp19b* mRNA is likely to promote ovarian maturation, because this enzyme catalyzes the final, rate-limiting step in the conversion of testosterone into estradiol in the gonads (Simpson et al., 1994). We found that exposure to OCT inevitably altered the transcription of steroidogenesis (*cyp19b*). Further research is needed to confirm whether OCT affects plasma sex steroid levels and/or fertility and reproduction in adult fish.

Exposure of zebrafish to OCT resulted in a significant up-regulation of *vtg1* in the liver at the H OCT accumulation level. *Vtg* mRNA is always induced in zebrafish by exposure to (xeno) estrogen, so the induction of vitellogenin in the liver is a sensitive and reliable indicator of exposure to exogenous estrogen (Sumpter and Jobling, 1995; Liu et al., 2010). In contrast to our findings, Blüthgen et al. (2014) observed no significant alteration of *vtg1* in the liver after 8 d exposure. The exposure time, concentrations, and development stage may impact the regulation of gene transcription. Especially considering the immediate decrease in OCT after water-renewal, it seems likely that transient OCT exposure affects the transcriptional level. The fact that 30 d exposure of fish to *EE2* results in an up-regulation of *vtg* (Bogers et al., 2006) implies that OCT has estrogenic activity in the liver. As discussed above, endogenous *E2* modulates ovarian development by regulating vitellogenin synthesis. Therefore, it is possible that changes in *vtg1* expression in the liver accelerated ovarian development.

*Rbp2a* retinoids play a significant role in various physiological processes such as cell growth, differentiation, and reproduction (Chen et al., 2012). In this study, the induction of *rbp2a* was observed in the livers of males and females even at the L OCT accumulation level (Fig. 6G and H). Many studies have reported alterations of retinoids in various species due to environmental contaminants – McKearin et al. (1986), for example, unequivocally demonstrated that vertebrate *rbp* mRNA levels are regulated by steroidogenesis. Levy et al. (2004) reported that *rbp* is up-regulated by *E2* and *EE2* in cultured *X. laevis* hepatocytes, which provides evidence that retinoids are a biomarker for detecting the specific actions of pure EDC. The changes in *rbp2a* that we observed therefore suggested an environmental endocrine-disruption effect of OCT on zebrafish. Moreover, when there is inhibited absorption



**Fig. 6.** Relative gene expression of *esr1*, *ar*, and *cyp19b* in the brain and gonad, and *vtg1*, *rbp2a* in the livers after OCT accumulation in zebrafish to L, M, and H level, respectively. Relative transcript abundance was quantified by qRT-PCR. Fold changes ( $\log_2$ ) were determined via  $2^{-\Delta\Delta Ct}$  method; results are given as mean  $\pm$  SD. Asterisks (\*) indicate statistical significance (\* $p \leq 0.05$ , \*\* $p \leq 0.01$ ).

of intestinal retinoids, the depleted stores of liver retinoids are transported to the eyes and ovaries (Chen et al., 2012). This means that the transcriptional result of *rbp2a* was likely to cause an overall promotion of development based on the increased vitellogenic oocytes in the ovaries and the up-regulation of *cyp19b* in the brain. Essentially, the expression of *rbp2a* in the liver is likely a negative feedback response to OCT accumulation in zebrafish.

### 3.5. Comparison of hormonal activity of OCT accumulation in vivo and in vitro assay

Multiple hormonal activities of OCT were also observed in an *in vitro* assay by Kunz and Fent. (2006). Comparison of the

hormonal activity of OCT evaluated by *in vivo* and *in vitro* assay is shown in Table 2. Both types of assay reveal the antiandrogenicity, androgenicity, and antiestrogenicity of OCT, but the estrogenic activity of OCT was not observed in the *in vitro* assay. A recent study found that OCT provokes a significant expression of ecdysone receptors in *Chironomus riparius* embryos, which also indicates the estrogenic activity of OCT *in vivo* (Ozáez et al., 2016).

Of course, the basic principle of *in vitro* assay is a simplification of the mechanism *in vivo*. To a certain extent, *in vivo* assay can be incarnated by *in vitro* assay but with simpler operation and high sensitivity. *In vitro* assay does not allow the researcher to metabolize chemicals as easily, however, so some of the more complex biological effects of chemicals on organisms do not occur.

**Table 2**  
Comparison of hormonal activities of OCT between *in vitro* and *in vivo* assays of zebrafish with H-level OCT accumulation.

Type of assay	Estrogenic activity	Antiandrogenic activity	Androgenic activity	Antiandrogenic activity
<i>In vitro</i> <sup>a</sup> (EC <sub>50</sub> )	×	2.57E-3 <sup>b</sup> (mol L <sup>-1</sup> )	6.27E-4 <sup>b</sup> (mol L <sup>-1</sup> )	2.45E-5 <sup>b</sup> (mol L <sup>-1</sup> )
<i>In vivo</i> <sup>b</sup> (“H” level)	<i>esr1</i> (+): gonad <i>cyp19b</i> (+): brain	<i>esr1</i> (-): brain	<i>ar</i> (+): brain	<i>ar</i> (-): gonad

Note:

<sup>a</sup> (Kunz and Fent, 2006).

<sup>b</sup> This study; “+” indicates up-regulation of target genes, “-” indicates the target gene was down-regulated, and “×” that it was not detected.

Metabolization may lead to some loss of the parent compounds, but the production of metabolites with potential activity should also be considered. For example, the UV filter BP-1 as a metabolite of BP-3 shows stronger estrogenic potency than BP-3, which mainly possesses antiestrogenic and antiandrogenic activities (Kunz and Fent, 2006). The metabolites of OCT formed *in vivo* may have higher hormonal activities than its corresponding parent compound, so it is not surprising that the estrogenic activity of OCT was detected *in vivo* but not *in vitro*. Moreover, EDCs affect estrogenic activity through a number of mechanisms including direct interactions (estrogen receptors) or indirect interactions (synthesis of sex steroid hormones, gonad development, and reproduction) (Sun et al., 2014). In some cases, *in vivo* assay has shown a comparable to better sensitivity to (xeno) estrogens than *in vitro* assay (Sonavane et al., 2016). It is advisable to apply *in vitro* assay to detect a specific toxicity, and essentially to create a useful reference for further *in vivo* assay. Overall, as discussed above, the results of both our *in vivo* assay and previous *in vitro* assays provide detailed information for the ecological risk assessment of OCT.

#### 4. Conclusions

By the exposure of female and male zebrafish to solutions of elevated OCT concentrations, we found that OCT molecules can indeed transfer rapidly from the solution to zebrafish. As a result, OCT was ultimately accumulated to 2321.01 (L), 31,234.80 (M), and 70,593.38 ng g<sup>-1</sup> (H) after 28 d when the average concentrations of the solutions were controlled at 28.61, 505.62, and 1248.70 μg L<sup>-1</sup>, respectively.

The multiple hormone activities of OCT and their effects on zebrafish were assessed via gross observations, histological observations, and RT-qPCR analysis. Even at the L OCT accumulation level, apparent up-regulation of *rbp2a* was observed in the livers of female and male zebrafish. The higher the OCT accumulation level, the more apparent the adverse effects – for example, female zebrafish with high levels of OCT accumulation showed significant increase in *GSI* and percentage of vitellogenic oocytes in their ovaries. High OCT accumulation also resulted in multiple hormonal effects in different tissues such as the up-regulation of *esr1* in the gonads and the induction of *vtg1* in the liver, indicating that OCT possesses estrogenic activity. The down-regulation of *esr1* in the brains of female zebrafish and inhibition of *ar* in the testes of male zebrafish also demonstrated the antiestrogenicity and antiandrogenicity of OCT. The accumulation of OCT to elevated levels also altered the expression of genes involved in steroidogenesis, *i.e.*, caused the up-regulation in the transcription of *cyp19b* in the brains of female zebrafish. We also compared *in vivo* and *in vitro* assay results regarding antiestrogenic, androgenic, and antiandrogenic activities. We concluded that the hormonal activities revealed by *in vivo* assay in this study can deepen our understanding of the toxic effects of OCT in the aquatic environment. Overuse of OCT should be avoided, or at least cautiously considered, from the viewpoint of ecological risk control in the future.

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