Biological effects of the benzotriazole ultraviolet stabilizers UV-234 and UV-320 in early-staged zebrafish (Danio rerio)

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**A B S T R A C T**

Among the benzotriazole ultraviolet stabilizers (BUVSs), UV-234 and UV-320 are frequently detected in aquatic ecosystem. Despite the fact that these chemicals are present in low ng/L levels in surface water, they show high bio-accumulation potential and pose exposure risks to aquatic organisms. However, there are limited toxicological data available in fish. In this study, zebrafish embryos were exposed to 0.01, 0.1 and 1 μM UV-234 or UV-320 for up to 6 days. Developmental toxicity as well as effects on mitochondrial bioenergetics, immune system responses, and locomotor activity in zebrafish were measured. After UV-234 treatment (0.1–1 μM), hatching time of embryos was increased compared to controls. There was also a ~20–40% reduction in non-mitochondrial respiration and oligomycin-dependent mitochondrial respiration in embryos treated with 1 μM UV-234 for 24 and 48 h respectively; conversely basal respiration and non-mitochondrial respiration were increased ~20–30% in embryos treated with 1 μM UV-320 at 48 h. Transcript levels of sod1 mRNA were down-regulated with both UV-234 and UV-320, suggesting an oxidative damage response. Considering that mitochondrial signaling regulates innate immune pathways, we measured the expression of immune related transcripts (il8, irf7, mmp9, il1, irf9, cxcl-C1c, nfkβ1, and ifng). Of these, only il8 and cxcl-C1c mRNA were decreased in response to 0.1 μM UV-320. To associate early molecular events with behavior, locomotor activity was assessed. UV-234 reduced larval activity in a dark photokinesis assay by ~15%, however behavioral responses at environmentally-relevant concentrations of BUVSs were not consistent across experiments nor BUVSs. These data suggest that BUVSs can perturb mitochondrial bioenergetics, embryonic development, and locomotor activity of zebrafish, but these responses appear to be dose-, time- and BUVSs dependent, suggesting these chemicals may have unique modes of action.

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1. Introduction

Benzotriazole ultraviolet stabilizers (BUVSs), a group of (2-hydroxyphenyl) benzotriazole derivatives, have excellent absorption capacity with a full spectrum of UV-light. They are widely used in personal care products (e.g. cosmetics and shampoos) as well as industrial products (e.g. plastics, coatings, adhesives and construction material) to protect against ultraviolet radiation from sunlight (Nakata et al., 2009; Ruan et al., 2012). Due to their high consumption volume, BUVSs have been widely detected in surface waters, wastewaters, sewage sludge, and biota (Kameda et al., 2011; Montesdeoca-Esporda et al., 2012; Nakata et al., 2009; Nakata et al., 2010; Nakata et al., 2012; Ruan et al., 2012; Wick et al., 2016; Zhang et al., 2011; Zhao et al., 2017). Among BUVSs, 2-(benzotriazol-2-yl)-4,6-bis(2-phenylpropan-2-yl)phenol (UV-234) (Figure S1) is one of the most frequently detected BUVSs in
Aquatic pollutants that disturb the balance between production and depletion of reactive oxygen species (ROS) can induce oxidative stress. A recent study has shown that UV-234 and UV-328 can increase ROS production as well as oxidative stress related genes that included glutathione peroxidase (gpx), superoxide dismutase (sod), and catalase (cat) in freshwater green algae *Chlamydomonas reinhardii* (Giraudo et al., 2017). In biological systems, mitochondrial oxidative phosphorylation is a major cellular source of ROS (Lushchak, 2011). Recent studies have shown that mitochondria participate in a broad range of immune pathways, functioning as signaling platforms and contributing to effector responses (West et al., 2011). Mounting evidence suggests that mitochondria facilitate antibacterial immunity by generating reactive oxygen species (ROS) and contribute to innate immune activation following cellular damage and stress (Cloonan and Choi, 2013; Larigule and Faustine, 2013; West et al., 2011). The innate immune system has a key role in the vertebrate immune response, and is exquisitely sensitive to environmental contaminants (Bols et al., 2001). Immunootoxicity can influence the development, growth, and survival of fish (Wester et al., 1994), which can ultimately affect the stability of fish populations. In zebrafish, the active innate immune system is detectable at day 1 of embryogenesis (Tredec et al., 2004). However, functionally mature adaptive immune system is not active until 4–6 weeks post-fertilization (Tredec et al., 2004). This temporal separation makes zebrafish (*Danio rerio*) embryos an appropriate model for studying innate immune in vertebrates (Novoa and Figueras, 2012).

To evaluate the sub-lethal effects of BUVSs in early-staged zebrafish, we firstly assessed the developmental toxicity of UV-234 and UV-320. Bioenergetic perturbations induced by exogenous chemicals during embryogenesis are expected to affect the developmental process, thus indicators to assess mitochondrial function including oxidative respiration (i.e. mitochondrial bioenergetics) and sod1 and sod2 expression levels were evaluated. Furthermore, several transcripts related to the innate immune system were measured to better address the interaction between mitochondria and the immune system. To provide relevance to higher level endpoints, we also investigated the effects of UV-234 and UV-320 on locomotor behaviors in zebrafish larvae, as early evidence of toxicity could lead to detrimental ecologically relevant responses (i.e. adverse outcome pathway).

### 2. Materials and methods

#### 2.1. Chemicals

2-(benzotriazol-2-yl)-4,6-bis[2-phenylpropan-2-yl]phenol (UV-234, CAS no. 70321-86-7, batch number: MKBW2452V) was purchased from Sigma-Aldrich Co. LLC (USA). 2-(benzotriazol-2-yl)-4,6-ditert-butylphenol (UV-320, CAS no. 38467-71-7, purity >98%) was purchased from Ark Pharm, Inc. (USA). Stock solutions of 0.1, 1 and 10 mM UV-234 or UV-320 were prepared by dilution in dimethyl sulfoxide (DMSO, CAS no. 67-68-5, purity >99.9%, Sigma). The exposure solutions were prepared by adding 10 μL of each stock solution into 100 mL of embryo rearing medium (ERM; 13.69 mM NaCl, 0.54 mM KCl, 0.025 mM Na₂HPO₄, 0.044 mM KH₂PO₄, 1.3 mM CaCl₂, 1 mM MgSO₄, 4.17 mM NaHCO₃, pH 7.2) yielding the final concentrations of 0.01, 0.1 and 1 μM UV-234 (equal 4.48, 44.8, and 447.6 μg/L) or UV-320 (equal 3.23, 32.3, and 323.4 μg/L). A vehicle treatment of DMSO served as control. The ratio of vehicle to ERM (1:10,000) was used in this study. Exposure concentrations used in this study were higher than those occurring in the environment (in the low ng/L).
2.2. Experimental design

Experimental design for exposures and endpoints are similar to that reported previously (Liang et al., 2017b). Please refer to the Supporting Information for more specific details. Briefly, adult zebrafish (2F, Wild type AB) were randomly selected from a breeding stock and placed in two tanks to generate the fertilized embryos. The ratio was three males to one female per tank. The fertilized eggs (~300–400) were determined to be at the same stage (shield stage, ~6 h post fertilization, hpf) and these embryos were used in subsequent experiments. Embryos were staged as outlined previously (Kimmel et al., 1995). Ten embryos were equally distributed into one glass beaker containing 10 mL of each exposure solution. Five replicate beakers were performed for each condition (control, 0.01, 0.1, and 1 μM UV-234 or UV-320). Embryos were maintained in an incubator at 26.0 ± 1 °C with a photoperiod of 14:10 h light: dark. For the duration of the experiment, a static waterborne exposure was conducted over 96 h with 90% water renewal with the chemical each day.

2.3. Embryo development

Developmental experiments were performed according to Liang et al. (2017b). Briefly, embryos were transferred to wells of a 96-well culture plate and were treated with the solvent control or one of the three concentrations of UV-234 or UV-320. UV-234 and UV-320 experimental groups were measured separately on two 96-well plates. Each treatment included n = 24 embryos. The 96-well plate was incubated at 26 ± 1 °C in the EVOS FL Cell Imaging System (Thermo Scientific, USA) and images of each well were automatically acquired every hour for 96 h. Developmental parameters that included hatching time, mortality, and malformation were manually scored using the method of Reimers et al. (2004).

2.4. Analysis of metabolic rates

Extracellular flux analyzer assay was optimized according to Souders et al. (2018). Please refer to the Supporting Information for more specific details. Two separate experiments were conducted for oxygen consumption rate (OCR) for both chemicals. After exposure for either 24 or 48 h, one embryo (n = 5 total) from separate biological replicates from each treatment was selected for mitochondrial respiration assessment using the XF24 Extracellular Flux Analyzer (Agilent Seahorse XF Technology, USA). Basal respiration [defined as (final basal OCR measurement)], mitochondrial respiration [defined as (final basal OCR measurement – final NaN3 OCR measurement)], and non-mitochondrial respiration [defined as (final NaN3 OCR measurement)] were calculated as per Seahorse XF Cell Mito Stress Test Kit User Guide (User Guide Kit 103015-100, Agilent). Here, basal respiration is baseline OCR data for each individual collected over a number of pre-defined cycles. Oligomycin is then used to assess ATP dependent respiration in the embryos (oligomycin-induced ATP production). Following NaN3, oxidative phosphorylation was completely blocked and non-mitochondrial respiration (e.g. oxygenase activity) was determined. The relevance of these endpoints for mitochondrial function and bioenergetics are provided in Souders et al. (2018); in short, constitute endpoints that suggest mitochondrial dysfunction following treatment with specific-acting mitochondrial toxicants.

2.5. Gene expression analysis

After 6 d exposure, larvae in each group were collected and 5 larval fish were pooled into one centrifuge tube from each beaker (n = 5/condition). Extraction of RNA and cDNA synthesis were performed according to Liang et al. (2017b). Primer sets for target genes were collected from literature (Table S1). The genes were selected to map the modulation of several important inflammatory mediators like interferon, gamma (Ifng), tumor necrosis factor a (tnfa), chemokine (C-X-C motif) ligand 8α (il8), chemokine (C-X-C motif) ligand 18β (cxcl-C1c), and matrix metallopeptidase 9 (mmp9) upon BUVSSs exposure (Jin et al., 2008). To better understand to upstream regulatory mechanism, we included regulators of the innate immunity toll-like receptors (tir5α and tir5b) and nuclear factor of kappa light polyepptide gene enhancer in B-cells 1 (nkb1), both trigger and regulate mentioned immune mediators and are often highly expressed in fish immune cells upon stimulation (Hayashi et al., 2001). Transcript levels of superoxide dismutase (sod1 and sod2) were also measured as an indicator of oxidative stress. Three reference genes (beta actin, β-actin; ribosomal subunit 18, rps18; and ribosomal protein L13a, rpl13a) were combined for normalization. Real-time PCR was performed according to Liang et al. (2017b). More specific details are provided in the Supporting Information.

2.6. Dark photokinesis behavioral assay

After exposure for 5d in beakers, larval zebrafish were individually transferred into a 96-well culture plate and placed in a Noldus DanioVision Observation Chamber (Noldus Information Technology, Netherlands). For each treatment, 24 fish were assessed for behavior in mid-afternoon. After being acclimated at 26 °C the instrument for 2h, the activities of fish were recorded by an infrared analog camera within the Danio Vision observation chamber, and the resulting video was collected and analyzed using the EthoVision software version 12.0 (Noldus Information Technology, Leesburg, VA). Larvae were tracked following a standard 50 min “white light routine” of alternating 10 min periods of light and dark beginning with a dark period (i.e. dark photokinesis). Analysis profiles were generated in EthoVision for distance moved and activity (i.e. percent pixel change across wells over time with an activity threshold of 16 to remove background noise) and an excel file was exported for further analysis.

2.7. Statistical analysis

Statistical analysis of variance was performed using SPSS (version 17.0). The normality of data was assessed using the Kolmogorov-Smirnov test and logarithmic transformation was performed prior to ANOVA to ensure data conformed to assumptions. The Levene’s test of homogeneity of variance was used to test for adherence to one-way analysis of variance (ANOVA) assumptions. Hatching time was analyzed using one-way ANOVA followed by Dunnnett’s post-hoc test was used to compare each treatment with the control. For hatching rates and survival rates, the Log-rank (Mantel-Cox) test was used to compare the differences between treatments and the control. Mitochondrial respiration was analyzed by two-way ANOVA and comparison of means between each treatment and the control was carried out using Dunnnett’s test as post hoc test. All quantitative data above are expressed as mean ± standard error (SEM). The qPCR results were analyzed by Kruskal-Wallis followed by Dunn’s multiple comparison tests. For behavior analysis, exported data were first pooled into 5 sections according to each of the dark/light periods measured. Treatments included DMSO control, 0.01, 0.1 and 1 μM BUVSSs. Data were analyzed using one-way ANOVA followed by Holm-Sidak’s multiple comparison tests. A probability of p < 0.05 was considered to be statistically significant. All figures were prepared using Graph-Pad Prism version 6.0 (GraphPad Software Inc., La Jolla, CA, USA).
3. Results

3.1. Developmental toxicity of UV-234 and UV-320 in zebrafish embryos

Mortality and hatching rate were not different between treatments and the control during 96 h exposure to UV-234 or UV-320 (Table S2 and Figure S2). However, there was an increase in hatching time for the UV-234 exposure groups (i.e. took longer to hatch). The average hatching time of zebrafish embryos in the control group was 74.5 h, while that in treatments was 78.95–79.68 h, ranging from 5.97% to 6.95% (df = 84, F = 7.7, p = 0.0001) (Fig. 1A). No differences were detected among doses, suggesting that this may be associated with a threshold effect as opposed to a dose dependent response at these concentrations. There were no differences observed for embryos exposed to UV-320 (df = 84, F = 1.47, p = 0.23) (Fig. 1B).

3.2. UV-234 and UV-320 exposure altered mitochondrial bioenergetics

To address any effects of UV-234 and UV-320 on mitochondrial bioenergetics, oxygen consumption rate (OCR) was measured in intact zebrafish embryos after 24 and 48-h exposure, respectively (Fig. 2A and B). After 24-h UV-234 exposure, non-mitochondrial respiration was significantly decreased following 1 μM UV-234 treatment (df = 32, F = 3.52, p = 0.02). Conversely, there were no changes in basal respiration (df = 32, F = 3.45, p = 0.17) nor mitochondrial respiration (df = 32, F = 2.68, p = 0.66) in any treatment compared to control (Fig. 3C–E, upper graph). For the 48-h UV-234 exposed embryos, mitochondrial respiration was significantly inhibited in 1 μM UV-234 treatment (df = 32; F = 2.68, p = 0.025). There were no differences detected in fish for basal respiration (df = 32, F = 3.45, p = 0.06) nor non-mitochondrial respiration for any treatment compared to the control (df = 32, F = 3.52, p = 0.16) (Fig. 2C–E, upper graph).

For the 24-h UV-320 treatments, there were no changes in basal respiration (df = 32, F = 2.02, p = 0.9), mitochondrial respiration (df = 32, F = 1.56, p = 0.69), nor non-mitochondrial respiration (df = 32, F = 1.91, p = 0.68) (Fig. 2C–E, lower graph). However, for the 48-h exposure to UV-320, basal respiration (df = 32, F = 2.02, p = 0.024) and non-mitochondrial respiration (df = 32, F = 1.91, p = 0.028) were significantly increased in embryos at 1 μM UV-320 group; no difference was detected in fish for mitochondrial respiration among groups (df = 32, F = 1.56, p = 0.059) (Fig. 2C–E, lower graph). Taken together, there is some evidence that these compounds affect respiration and this effect appears to be dependent upon the dose and the time point examined.

3.3. UV-234 and UV-320 exposure affected the expression of genes related to oxidative stress

To test the hypothesis that BUVs induce mitochondrial damage response, the mRNA levels of sod1 and sod2 were measured in larvae following exposure to UV-234 and UV-320 (Fig. 3). After 6-d exposure of UV-234, a significant decrease in sod1 mRNA levels were detected in fish from the 0.01 μM treatment (−1.87-fold, p = 0.045) (Fig. 3A). Conversely, levels of sod2 mRNA were up-regulated following 1 μM UV-324 exposure (15.1-fold, p = 0.01) (Fig. 3B). After 6-d exposure to UV-320 treatment, levels of sod1 were significantly decreased in fish from the 0.1 μM UV-320 treatment (−6.4-fold, p = 0.017) (Fig. 3C). Conversely, levels of sod2 were increased in fish from the 0.1 μM UV-320 treatment (7.7-fold, p = 0.002) (Fig. 3D). Fold change on a log2 scale depicts changes in gene expression, and relative expression was compared to the control group to reveal differences among the three treatments.

3.4. UV-320, but not UV-234, altered the expression of genes related to the immune system

We examined transcript levels for genes involved in toll-like receptor (TLR) signaling pathways that included toll-like receptors (tlr5a and tlr5b), immune response regulator nfkβ1, pro-inflammatory cytokines (ifng, tnfa), chemokines (il8, cxcl-C1c) and matrix metalloprotease mmp9 (Fig. 4 and Figure S3). All transcripts tested were not altered in abundance following UV-234 exposure (Figure S3). In the UV-320 exposure experiments, tlr5a, tlr5b, nfkβ1, ifng, tnfa and mmp9 were not different in relative abundance in the zebrafish larvae (Fig. 4A–E and H). However, the level of il8 was decreased in fish exposed to 0.1 μM UV-320 group compared to the control group (7.9-fold, p = 0.02) (Fig. 4F). Similarly, levels of chemokine cxcl-C1c were significantly decreased in fish exposed to 0.1 μM UV-320 group compared to the control (2.8-fold, p = 0.03) (Fig. 4G).

3.5. Locomotor behavior of zebrafish larvae was affected by both UV-234 and UV-320

Behavioral responses were subtle for both UV stabilizers and were dependent upon the dose and chemical. In the first 10 min, decreased activity of larval zebrafish pre-adapted to darkness was detected in larvae exposed to UV-234 (df = 36, F = 34, p = 0.0001) (Figure S4A, B). In the following light-dark cycle, as expected with

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Fig. 1. Hatching time of zebrafish embryos following ~96 hpf exposure to different concentrations of (A) UV-234 or (B) UV-320. Data were analyzed using one-way ANOVA followed by Dunnett’s post-hoc test. All quantitative data above are expressed as mean ± standard error (SEM). Asterisk denote significant differences (*p < 0.05, **p < 0.01, ***p < 0.001) between treatments and controls.
dark photokinesis, locomotor activity increased with loss of illumination, but there was no difference in the activity of larvae in UV-234 groups compared to the control (Figure S4A, B). In contrast, a decrease in movement (distance moved) was detected in larvae following treatment to 1 \( \mu M \) UV-234 (\( df = 36, F = 4.1, p = 0.028 \)) in 20–30 min following light extinction (Fig. 5A and B). During subsequent light (30–40 min), the total distance moved in larvae was different than controls (\( df = 36, F = 33, p = 0.003 \)). Distance moved was increased in response to 0.01 \( \mu M \) while distance moved was lower in larvae exposed to 0.1 \( \mu M \) UV-234 compared to control (Fig. 5A and B). Taken together, there was some evidence that UV-234 altered behavior; however there was no discernible pattern in response.

For UV-320 treatments, increased activity of larval zebrafish pre-adapted to darkness was detected in larvae exposed to 1 \( \mu M \) (\( df = 36, F = 37, p < 0.0001 \)) compared to the control (assessed in first 10 min) (Figure S4C, D). Increased distance moved for larvae was also observed with 1 \( \mu M \) UV-320 exposure (\( df = 36, F = 12, p < 0.0001 \)) in the first 10 min (dark) (Fig. 5C and D). In response to illumination, larvae increased their distance moved in the 0.1 \( \mu M \) UV-320 treatment (\( df = 36, F = 6.3, p = 0.004 \)) compared to control (10–20 min) (Fig. 5C and D). However, following a light-dark cycle, fish exhibited a decrease in distance moved in the 1 \( \mu M \) UV-320 treatment (\( df = 36, F = 4.3, p = 0.016 \)) in 30–40 min (light) (Fig. 5C and D). Taken together, changes in activity was small and dependent upon the dose.

4. Discussion

BUVSs are emerging pollutants of concern which have been widely detected in aquatic environments (Nakata et al., 2009; Song et al., 2014). Depending on their structure, BUVSs and their
metabolites can exert structure-dependent effects in both mammals (Nagayoshi et al., 2014) and aquatic organisms (Fent et al., 2014; Liang et al., 2017a; Zhuang et al., 2017). Despite several reports on their endocrine disrupting effects in mammals and fish (Fent et al., 2014; Nagayoshi et al., 2014; Zhuang et al., 2017), studies investigating their potential for developmental toxicity and immunotoxicity have not yet been conducted. In this study, zebrafish were exposed to two commonly detected BUVSs, UV-234 and UV-320, for up to 6 days and developmental toxicity, mitochondrial bioenergetics, behavior, and molecular endpoints related to the oxidative stress and the immune system were measured.

Fig. 3. The mRNA levels of *sod1* (A & C) and *sod2* (B & D) in zebrafish larvae following treatment with UV-234 or UV-320 for 6 days. Upper graphs are data for UV-234 exposure and the lower graphs are data for UV-320. All quantitative data above are expressed as mean ± SEM. Asterisk denote significant differences (*p < 0.05) between treatments and controls.

Fig. 4. The mRNA levels of (A) *tlr5a*, (B) *tlr5b*, (C) *nkb*, (D) *il6*, (E) *tnfa*, (F) *il8*, (G) *cxcl-1c*, and (H) *mmp9*, transcripts associated with innate immunity, were measured after exposure to UV-320 for 6 days. All quantitative data above are expressed as mean ± SEM. Asterisk denote significant differences (*p < 0.05) between treatments and controls.
4.1. Hatching time of zebrafish embryos is affected differently by BUVSs

In this study, hatching time was significantly increased (i.e. delayed hatching) with exposure to UV-234, but this was not the case for UV-320 treatments. As pointed out in our previous study, time to hatch may be a sensitive indicator for adverse effects in fish embryos exposed to chemicals (Liang et al., 2017b). The mechanism for the delayed hatching may involve a change in the elasticity of the chorion or in chorionic hatching enzyme function (Winnicki et al., 1970), although this must be rigorously tested with the chemical under study. Another hypothesis related to the delay in hatching may involve oxidative respiration in developing fish; exposure to hypoxic environments delayed hatching of sheepshead minnow (Cyprinodon variegatus) embryos (Hendon et al., 2008). In the present study, oxygen consumption rate (OCR) was decreased by UV-234 while there was an increase in OCR with UV-320. Previous studies have suggested that chemical-induced alterations in mitochondrial bioenergetics in zebrafish embryos may contribute to developmental toxicity (e.g. curved tails or abnormal hatching time) (Wang et al., 2018; Zhang et al., 2017). Noteworthy is that hatching delay is reported to be associated with the interference in the thyroid hormone (TH) system (Timme-Laragy et al., 2006). In our previous study, UV-234 was shown to alter the expression of the thyroid hormone pathway in zebrafish embryos (Liang et al., 2017a). Thus, we hypothesize that mitochondrial dysfunction and disruptions in TH system induced by UV-234 may both contribute to the hatching delay. However, further studies are required to elucidate the exact role of each in a biologically complex process.

4.2. BUVSs induce oxidative damage in developing zebrafish at the transcript level

Mitochondrial dysfunction can lead to increased reactive oxygen species (ROS) and apoptosis. Mitochondrial oxidative phosphorylation is a major cellular source of ROS and levels of ROS are not only determined by production, but also by the rate of ROS degradation or inactivation (Mailloux and Harper, 2012). ROS can be converted into hydrogen peroxide (H₂O₂) in the mitochondrial matrix by superoxide dismutase 2 (SOD2). H₂O₂ can then freely cross mitochondrial membranes or can be further detoxified by additional mitochondrial antioxidant enzymes, such as glutathione peroxidase (GPX). In the cytosol, superoxide is converted by SOD1 into H₂O₂, which is further detoxified by the peroxisomal enzyme catalase (Li et al., 2013). Redox imbalance can be therefore generated when a cell accumulates an excessive amount of ROS which overwhelms antioxidant defenses.

In this study, oxidative and non-mitochondrial respiration of zebrafish embryos were suppressed by 1 μM UV-234 exposure. Noteworthy was that mRNA level of sod2 was increased at 1 μM UV-234 in larval fish (6 dpf). The upregulation of sod2 suggests that the antioxidant defense system was activated due to redox imbalance (Knecht et al., 2013) and it is hypothesized that this could be due to the impaired bioenergetics observed with BUVSs. In mammal and fish cells, 10% of cellular oxygen uptake is due to non-mitochondrial (e.g. oxidase), rather than mitochondrial use of oxygen for respiration (Raftery et al., 1999). The increase in non-mitochondrial oxygen consumption has been considered to be a protective mechanism by some to minimize oxidative stress (Raftery et al., 2017).
Similar to the pattern observed with UV-234, the mRNA levels of \textit{sod1} were down-regulated while that of \textit{sod2} were upregulated in fish exposed to 0.1 \textmu M UV-320. This may be associated with the upregulation of basal and non-mitochondrial respiration at 1 \textmu M UV-320, indicative of redox imbalance. The elevated \textit{sod2} and decreased \textit{sod1} mRNA levels suggest that there is a molecular response to increased mitochondrial ROS production as expected. Previous studies have shown that there can be increased levels of \textit{sod2} mRNA in response to oxidative stress caused by environmental stress or chemicals in zebrafish (Cai et al., 2012; Tiedke et al., 2013). Moreover, Zou et al. (2017) reported that isoniazide induces oxidative stress in developing zebrafish by increasing ROS, and that this was associated with a decrease in \textit{sod1} mRNA levels. Our data indicate that UV-320 may affect mitochondrial bioenergetics and induce oxidative stress in developing zebrafish.

4.3. The effects of BUVSSs on locomotor activity in zebrafish

Chemicals that impair mitochondrial function may also underlie abnormal development in fish embryos (Bestman et al., 2015). In early life stages of fish, behavioral responses can act as sensitive indicators of toxicant exposure (Liang et al., 2017b). Here, activity of zebrafish larvae was assessed using a standard light-dark challenge. The activity of zebrafish larvae is characterized by higher activity in the dark compared to light at early stages of development. This hyperactive state has been termed the visual motor response (VMR) (Mueller and Neuhaus, 2012). Hypoactivity following UV-234 exposure in the larvae was observed in these experiments and implies that the chemical caused a decrease in visual sensitivity. To be noted, during the second light-dark cycle (40–50 min), the behavior of UV-234 exposed larvae became indistinguishable from that of control animals. Thus, at these environmentally relevant concentrations, it is somewhat unclear how BUVSSs affect behavior and the response may be clutch/dose specific. In a previous study of zebrafish larvae exposed to cocaine, loss of a difference in locomotor activity was also observed after longer exposure regimes; in that study, the possible reason was proposed to be the elimination of accumulated cocaine in the brain of fish (Kirla et al., 2016). The effect of UV-234 was subtle, and the loss of a response at the end of the assay may be attributed to the biotransformation or habituation in the assay. Nevertheless, these results suggest that UV-234 may influence neuronal pathways that mediate behavioral responses to light in zebrafish larvae.

In contrast to UV-234, UV-320 did not significantly alter locomotor activity of zebrafish larvae in the dark cycle, but it altered larvae travel distance in light periods. Moreover, in continuous darkness (first 10 min of the assay), increased activity of larvae was observed in the highest dose. Naturally, zebrafish prefer dark areas over light areas. Larval fish also display a startle response when exposed to a sudden change in light intensity (Colwill and Creton, 2011). It has been reported that stress responsive behavior is highly sensitive to neuroactive chemicals (e.g. diazepam and carbamazepine) (Brandao et al., 2013). However, to date, few studies have demonstrated that BUVSSs induce neurotoxicity in fish at the physiological or molecular level. Liang et al. (2016) demonstrated that > 0.42 \textmu M benzotriazole can induce alterations of brain proteomic profiles related to neurotoxicity in female Chinese rare minnows (Gobiocypris rarus). However, currently strong evidence is lacking for physiological or behavioral responses to support neuroactivity of benzotriazole and its derivatives (e.g. BUVSSs). Additional studies investigating other behaviors such as anxiety or feeding behavior may reveal evidence for neurological or energy-related limitations.

4.4. The immune system: a potential target for BUVSSs toxicity?

Apart from producing ATP, controlling calcium homeostasis, and regulating apoptosis, mitochondria also act as a sensor for stressors and can initiate inflammatory signals in the cell by activating the innate immune system (Lartigue and Faustin, 2013; West et al., 2011). Studies show that dysfunction in mitochondria is related to inflammatory diseases and disorders (reviewed in Cloonan and Choi, 2013). Altered oxygen consumption in the embryos as well as changes in the mRNA levels of \textit{sod1} and \textit{sod2} in response to BUVSSs in the present study led us to the hypothesis that the innate immune system may also be affected by BUVSSs toxicity.

To explore the potential for immunotoxicity of BUVSSs, we measured transcripts of several key inflammatory markers (\textit{tnfa} and \textit{ifng}) and signals that attract other immune cell types to the site of inflammation — chemokines IL-8, CXC-C1c and mmp9. All of these genes are upregulated by toll-like receptor (TLR) dependent NF-\textbf{κ}B signaling pathway, that function as enhancers and regulators of genes involved in innate immune response (Hayashi et al., 2001). Despite the similarity in the structure of UV-234 and UV-320, only UV-320 showed any evidence for an immunomodulatory response. All concentration of UV-234 tested did not interfere with the expression of transcripts investigated, but fish in the 0.1 \textmu M UV-320 group showed evidence for downregulation of chemokines (CXC-C1c and il8). Both of these transcripts are regulated by nuclear factor NF-\textbf{κ}B. However, we did not observe any changes in the transcriptomic level of \textit{tnfk1} in our exposures. In contrast to chemokines, transcriptomic levels of pro-inflammatory cytokines (\textit{tnfa} and \textit{ifng}) and upstream regulators (\textit{tnfk1} and tlr5qa/b) were not changed, indicating that the signaling pathway TLR- NF-\textbf{κ}B -proinflammatory cytokines may not be modified by UV-320.

Both downregulated cytokines Cxcl-c1c (also known as Cxcl18b) and IL-8 (Cxc18) are chemokines molecules, that were shown to be a reliable inflammatory markers. These CXC subgroup of chemokines are the key controllers of neutrophil chemotaxis in mammals and in teleost fish including zebrafish (de Oliveira et al., 2013; Deng et al., 2013). They are produced by innate immune cell (e.g. macrophages), but also by non-infected cells that immediate surround the infected/inflamed tissue (Torraca et al., 2017). They control several biological function including homestasis, cell recruitment and immune function (Russo et al., 2014). Dysregulation of these chemokines have the direct effect on the ability of the host to respond the microbial infection and to eliminate the pathogens (Torraca et al., 2014). In addition, Cxcl signaling axis significantly contribute to disease-associated processes, including tissue injury, fibrosis, angiogenesis and tumorigenesis (Russo et al., 2014; Torraca et al., 2014). Inhibition of cxc-c1c and il8 by UV-320 suggest that chemokines may be potential molecular targets for BUVSSs.

4.5. Conclusions

In conclusion, both UV-234 and UV-320 can affect mitochondrial bioenergetics and the expression of transcripts related to oxidative stress (i.e. \textit{sod1} and \textit{sod2}) in zebrafish embryos. However, our data support the notion that these UV stabilizers may also act differently in terms of their mechanisms. We hypothesize that UV-234 can affect mitochondrial bioenergetics and cause hatching delay in zebrafish embryos. However, UV-320 is proposed to alter immune response via transcriptional mechanisms following disrupting mitochondrial bioenergetics in zebrafish. The underlying mechanisms may contribute to the differential activity behaviors of larval zebrafish in response to changes in illumination. Further research in neuro-toxicological effects of BUVSSs after longer-term exposure is needed to provide additional evidence for the behavioral response observed here. Additional molecular targets should
also be investigated as potential biomarkers for the exposure to BUVSs in fish.

A final point to make is that we understand little regarding the toxicity and biological effects of biotransformation products of BUVSs. Zhuang et al. (2017) observed that UV-328 had an enhanced anti-androgenic effect following CYP3A4-mediated biotransformation, and showed that hydroxylation is the dominant process involved in the biotransformation. Recently, Damalas et al. (2018) assessed the toxicity of 3 benzotriazoles that included 1-H-benzotriazole (BT), 4-Methyl-1-H-benzotriazole (4-MeBT), and 5-methyl-1-H-benzotriazole (5-MeBT) as well as their biotransformation products in zebrafish (Danio rerio) larvae; it was demonstrated that parent compounds had a higher toxicity potential compared with their biotransformation products, and 4-MeBT showed the highest toxicity potential. However, no data on transformation products of UV-234 and UV-320 exist to date and this remains a significant knowledge gap moving forward in assessing environmental risks associated with BUVSs.

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Appendix A. Supplementary data

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References


