CYP1A/Regucalcin Gene Expression and Edema Formation in Zebrafish Embryos Exposed to 2,3,7,8-Tetrachlorodibenzo-p-dioxin

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Abstract In this study, zebrafish eggs were exposed to a relatively low concentration (50 pg/mL) of 2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD) for 72 h and then transferred to vehicle/TCDD-free water for the remainder of the experiments. Mortality, heart rates, edema severity, CYP1A, and regucalcin gene expressions were investigated to study TCDD-induced toxicity in zebrafish during the early life stage. Results indicated that the 50 pg/mL TCDD caused severe and visible developmental toxicity. Further research of the long term and low concentration of TCDD exposure is required.

Keywords 2,3,7,8-Tetrachlorodibenzo-p-dioxin · Zebrafish · Edema · Gene expression

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As an unwanted byproduct, 2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD) is frequently detected in environmental samples. In 1990, whole squawfish contained detectable TCDD at all selected sample sites in Willamette River, ranging from 0.5 to 1.5 pg/g (Curtis et al. 1993). Besides, TCDD bioaccumulation by fish in Great Lakes is wide-spread, which indicates that fish consumption is the major pathway of exposure to human beings (Cook et al. 2003). Concerns over low concentration and long-term exposure of TCDD to fish may help to understand and commence the existing environmental situations.

TCDD caused several developmental toxic responses to teleost fish through the AHR/ARNT signaling pathway (Schmidt and Bradfield 1996), particularly the yolk sac and pericardial edema as one of the most vital responses. Likewise, the occurrence of edema in laboratory-reared larvae of wild lake trout from Lake Ontario has been linked to exposure to TCDD and structurally related compounds (Cook et al. 2003). TCDD exposure to lake trout eggs produces a dose-related induction of CYP1A in vascular endothelial cells of embryos and sac fry, which correlates with yolk sac edema, hemodynamic dysfunction, and subsequent sac fry mortality (Guiney et al. 1997). On the other hand, exposure to TCDD substantially increases the intracellular free Ca²⁺ level in several cell types (Puga et al. 1997; Canga et al. 1993), and regucalcin (RC), a Ca²⁺-binding protein, plays a role in the maintenance of intracellular Ca²⁺ homeostasis by activating Ca²⁺ pump enzymes, was also studied in rat model to evaluate the risk posed by TCDD (Takahashi and Yamaguchi 1997; Pastorelli et al. 2006). In this study, mortality, heart rates, edema severity, CYP1A, and RC gene expressions were studied to investigate low dose of TCDD-induced toxicity in zebrafish during the early life stage.

Materials and Methods

TCDD (99% purity) dissolved in 100% DMSO was purchased from Cambridge Isotopes (Cambridge Isotope Laboratories, Andover, MA), and DMSO (AMRESCO) was used as vehicle control in all experiments.

AB-line zebrafish were raised and maintained according to the procedures described by Westerfield (1995). Embryos and juvenile fish were maintained at 26 ± 1.5 °C in a 14 h light and 10 h dark cycle. Newly fertilized eggs, approximately beginning at 2 h poster-fertilization (hpf), were exposed to vehicle (0.5% DMSO) or to 50 pg/mL TCDD for 72 h in a 6-well plate with gentle rocking. During the 72 h exposure time, 40% solution of each well was replaced with fresh confected reagent. Following the 72 h static exposure, the embryos were rinsed with water and maintained in aerating water for further observation. Each group had two parallels and experiments were repeated at least two times under the same condition. As an index of toxic responses, heart rate was determined in a 30 s interval using 40× microscope (Olympus, Japan). Embryos were randomly chosen in vehicle or TCDD beginning from 36 hpf, when the embryos already had heartbeats, and this was repeated every other day since then.

Edema incidence was scored as the percent of live embryos showing any discernable signs of edema at the indicated time (Henry et al. 1997). If needed, juvenile fish were anesthetized with tricaine (MS222; Sigma). The area was calculated as previously described (Prasch et al. 2004). The area of yolk sac and pericardium were determined using the area measurement function of BI 2000 Image (TME Tech Ltd, Chengdu, China). All photos were taken at $40\times$ with the fish in a lateral orientation at the same resolution.

Using Trizol reagent (Invitrogen, Carlsbad, CA), total RNA was isolated from five zebrafish embryos selected randomly from each replicate of the control and 50 pg/mL TCDD group during three times of embryo development: 52, 96, and 168 hpf. Then, each total RNA (2 µg) was reversetranscribed to single strand cDNA using M-MLV Reverse Transcriptase reagents (Promega UK Ltd.) according to the supplier's instructions. To detect zfCYP1A (GenBank Accession no. BC094977) and zfRC mRNA (GenBank Accession no. BC075882) expression at the three above stages, we used related primers (5'-GGAACTTCT TCA TCGTCGTG-3' and 5'-CATATCGTAGTATCCGTGGCT-3' for ZfCYP1A, 5'-CTGGAGGATTGTCT AACCG-3' and 5'-CCAAAACAGCATGAAGTGA-3' for ZfRC) designed by Primer Premier 5.0 (Premier, Palo Alto, USA) to amplify a 188 base-pair (bp) fragment of the zfCYP1A gene, and a 202 bp fragment of the zfRC gene. A denaturing step (94°C, 5 min) was followed by both of 35 cycles in the following conditions: denaturation (94°C for 30 s), annealing (56°C for 30 s), and extension (72°C for 1.5 min) in TC-312 (TECHNE). PCR products were size-fractionated in 1.5% agarose gels, stained with ethidium bromide, and measured densitometrically using JD-801 (JEDA Tech Ltd, Jiangsu, China) image software. β -Actin (GenBank Accession no. BC045846; 5'-AGCTCATAGCTCTTCTCCAG-3' and 5'-CTGGTATCGTGATG GACTCT-3' for primers) was used as an internal standard, with the amplifying cycles of 30.

Statistical analysis was done using the SPSS package (version 10.0; SPSS Company, Chicago, IL, USA). Mortality and edema rate were statistically compared between TCDD and control-DMSO using Chi-square test. For other time course experiment a t-test was employed to compare TCDD to DMSO at each time point independently, after ensuring homogeneity of variances between groups. For all experiments p < 0.05 was used to determine significance.

Results and Discussion

To perform the experiment successfully, normally developing eggs were chosen, so the fertilization rate was regarded as 100%. As in Fig. 1, mortality of the control and TCDD exposed group had fine discrimination at the first 180 hpf (p < 0.05). Embryos of the control went to a steadily developing stage after removing from their chorion and swimming for food, while the TCDD-treated embryos exhibited different types of developmental toxicity. Hence, the mortality of TCDD-treated embryos elevated and

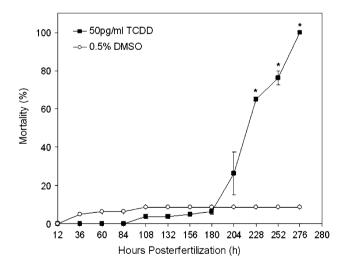


Fig. 1 Comparison of mortality between 0.5% DMSO-treated embryos and 50 pg/mL TCDD-treated embryos. After 72 h static exposure, embryos were transferred to aerating water for further observation. At the first 204 hpf, there was no significant difference between the control and the TCDD exposed group, where after TCDD-treated embryos decreased abundantly because of TCDD developmental toxicity. Each symbol represents the mean \pm SEM. *Significance compared to the equivalent stage embryos of control (p < 0.05)



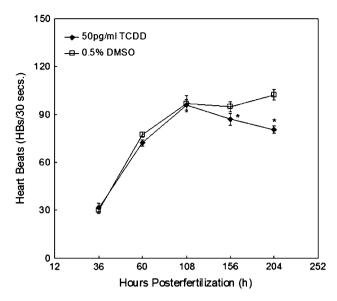


Fig. 2 Comparison of heart beats during a 30 s interval between 0.5% DMSO-treated embryos and 50 pg/mL TCDD-treated embryos (n = 6). From 156 hpf, there were significant differences between the control and the TCDD-exposed embryos. After 204 hpf heart beats decreased notably by TCDD, and the heart failed to palpitate till death (data was not expressed in this figure). Asterisks show significant difference from the equivalent stage embryos of control (p < 0.05)

finally reached 100% at 276 hpf. The heart in zebrafish is the primary target apparatus for TCDD exposure, which is not only affected in morphology but also in its function (Antkiewicz et al. 2005), and in some extent the rate of heart can reflect the affection of blood flow. Results revealed that TCDD prominently weakened the heart rate of treated embryos compared to the control since 156 h (Fig. 2).

In this research, 50 pg/mL, the relative lower exposure concentration of TCDD, not only weakened the heart rate but also caused serious edema to zebrafish (Fig. 3A). The edema rate was nearly 100% (Fig. 3B). Even though the yolk sac area of control decreased as embryo developed because of yolk absorption, the TCDD-treated yolk sac area elevated visibly (Fig. 3C). Correspondingly, from 144 hpf, TCDD significantly increased the pericardial area of treated embryos (p < 0.05, Fig. 3D). After 192 hpf the yolk sac edema caused by TCDD was quite severe, so the pericardial edema of TCDD-treated embryos was relatively inconspicuous compared with the control.

As in Fig. 4A, in the mass, CYP1A mRNA expressions of the control were equivalent and kept a lower level through the whole three stages. Otherwise, at 52 hpf no

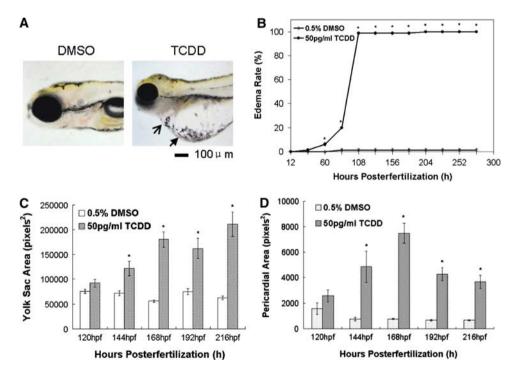


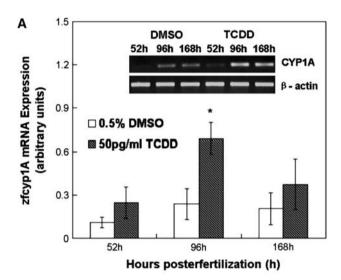
Fig. 3 Comparison of edema rate and edema area between 0.5% DMSO-treated embryos and 50 pg/mL TCDD-treated embryos at indicated time. (**A**) Photos illustrate a lateral view of edema severity. Void arrows indicate the pericardial cavity, and solid arrows represent the yolk sac edema compared to the control. (**B**) Edema rate. (**C**) The comparison of yolk sac area. From 144 hpf, edema severity of TCDD-treated group was rather significant. Yolk sac area of the control

decreased due to nutrition. (**D**) The comparison of pericardial area. Also from 144 hpf, there was significant difference between the control and the TCDD-treated embryos. Each symbol represents the mean \pm SEM of five to eight embryos. Asterisks show significant difference between the same stages of control and TCDD-treated embryos (p < 0.05)



difference was observed between the TCDD-exposed embryos and the vehicle in CYP1A mRNA expression, while it was obviously induced by TCDD at 96 hpf, but at 168 hpf its expression level fell to the normal. On the other hand, there was no significant difference in RC mRNA expression at each selected time compared to the control (Fig. 4B).

Fish are particularly sensitive to chemical exposure, especially during early development (Walker and Peterson 1994). Zebrafish, as a prominent model vertebrate, was successfully applied to the toxicity investigation of TCDD.



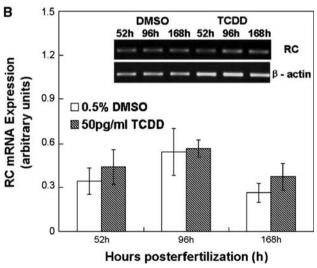


Fig. 4 Comparison of gene expressions between 0.5% DMSO-treated embryos and 50 pg/mL TCDD-treated embryos at indicated time. (**A**) *Zf*CYP1A. The expression in the control was tiny, while the expression caused by TCDD was elevated significantly at 96 hpf and decreased to lower levels at 168 hpf. (**B**) *zf*RC. There was no significant difference in RC mRNA expression at each selected time compared to the control. The insets were typical electrophoresis results. Values are mean \pm SEM of n = 4. Asterisk indicates significant difference from equivalent stage embryos of control (p < 0.05)

Most studies concerning TCDD waterborne exposure to zebrafish have been done at short-term and high doses exposure of dioxin (Antkiewicz et al. 2005; Prasch et al. 2004). Compared to those, we used a lower concentration of TCDD (50 pg/mL) exposed to zebrafish statically for 72 h at its early stage, during which embryos could not swim. As described above, TCDD exposure caused mortality, severe edema and other toxicity endpoints to zebrafish embryos.

As is known now, edema is caused by two mechanisms: water barrier or/and decreased water export (Hill et al. 2004). Many factors are involved in the formation of edema, and some gene expressions are correlative. CYP1A is the best understood target gene (Nebert et al. 1990). TCDD causes a potent and sustained elevation of CYP1A mRNA expression (Andreasen et al. 2002) that precedes development toxicity of the different endpoints (Henry et al. 1997). However, people found TCDD caused developmental toxicity through a CYP1A-independent mechanism in zebrafish because the zfcyp1a-MO effectively prevented CYP1A up-regulation, but it did not prevent the signs of developmental toxicity, including pericardial edema, slowed blood flow, craniofacial malformation, and defects in erythropoiesis (Carney et al. 2004). In this research, CYP1A mRNA expression was investigated in three different periods: 52, 96, and 168 hpf. These three stages represented in, after and later of TCDD exposure, respectively. Results showed that CYP1A level in TCDD-treated group fell after a rise ultimately. The variety of CYP1A transcription paralleled with the TCDD exposure time, which reflected the bioaccumulation and metabolism of TCDD in vivo. The former results revealed that if exposure to TCDD is delayed until after 96 hpf, edema is not observed, which suggest TCDD interferes with cardiovascular and erythropoietic developmental processes that are normally completed by 96 hpf (Belair et al. 2001). In this relatively low concentration of TCDD exposure, zebrafish embryos had no toxic symptoms at 52 hpf, while the CYP1A expression was also not affected. Later, with developmental toxicity occurring, TCDD began to produce its effect in vivo and induced CYP1A expression. Moreover, toxicity endpoints do not occur shortly after exposure, but takes place from 72 hpf or later. This association is likely to explain why CYP1A level induced by TCDD has distinctness between 52 and 96 hpf. Then, at 168 hpf CYP1A mRNA expression in TCDD-treated embryos fell instead of maintaining its high level, which may also be related to TCDD depuration.

Also, the kidney is the most important organ for the homeostasis of water and electrolytes (Cao et al. 2003). Kidney dysfunction can lead to a loss of plasma proteins, such as albumin, in the urine. The hypo-osmotic effect in blood drives extracellular fluid out of the vasculature,



leading to edema (Cao et al. 2003). RC has been demonstrated to have an inhibitory effect on various enzyme activation by Ca²⁺ and calmodulin in liver and kidney cells (Kurota et al. 1997; Yamaguchi and Mori 1990). In zebrafish embryo, TCDD exposure increased the intracellular calcium level, triggering Ca²⁺-mediated endonuclease activity leading to apoptosis (Dong et al. 2002). Our results indicated there was no significant difference in RC mRNA expression at each selected time compared to the control. Similarly, TCDD did not alter the abundance of RC transcripts or proteins in rat model (Pastorelli et al. 2006). So taken together, there may be some other Ca²⁺-regulated protein rather than RC which would be involved in TCDD-induced edema.

Through this experiment, it can be concluded that TCDD exposure with a relative lower concentration can cause severe toxicity endpoints to zebrafish embryos, especially the yolk sac and pericardial edema. Thus, further research at environmentally lower doses of TCDD exposure needs to be carried out.

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