



ORIGINAL ARTICLE

Toxic effects of imidacloprid and sulfoxaflor on *Rana nigromaculata* tadpoles: Growth, antioxidant indices and thyroid hormone-related endocrine system



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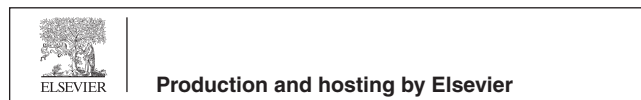
Imidacloprid;
Sulfoxaflor;
Rana nigromaculata tadpoles;
Thyroid disruption;
Gene expression

Abstract Imidacloprid and sulfoxaflor have potential damage to nontarget aquatic organisms. However, limited information has been provided on their underlying toxicity effects on *Rana nigromaculata* tadpoles. Thus, the acute toxicity and chronic effects of imidacloprid and sulfoxaflor on *R. nigromaculata* tadpoles were studied. Acute toxicity indicated that 96 h for LC₅₀ values of imidacloprid and sulfoxaflor were 173.55 and 427.37 mg/L, respectively. In this research, we explored antioxidant enzymes, some biological indexes, hormone levels and expression of relative tadpole genes involved in thyroid hormone-dependent metaplastic development after exposure for 28 days under 1/10 and 1/100 LC₅₀. Results showed an increase in the activity of antioxidant enzymes such as superoxide dismutase (SOD), catalase (CAT) and glutathione S-transferase (GST) in the tadpoles, and a low bioconcentration level with a bioconcentration factor (BCFs) < 1. The impact on the development of tadpoles was induced by the breakdown of hormonal levels engaged in metamorphosis. According to the real-time PCR results, imidacloprid and sulfoxaflor delayed amphibian metamorphosis by modifying mRNA expression, indicating that imidacloprid and sulfoxaflor

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may have an endocrine-disrupting effect on *R. nigromaculata* tadpoles. These findings were indicative of the toxicity of imidacloprid and sulfoxaflor to *R. nigromaculata* tadpoles.

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1. Introduction

Rice is one of the major food crops in China and plays an important role in ensuring national food security. Over the past few decades, the output of rice in China has increased rapidly due to improved varieties, advances in farming techniques and the extensive use of chemical fertilizers and pesticides. However, the period when pesticides are often used in agricultural crops corresponds to the period when amphibian larvae breed and develop (Mann et al., 2009). Owing to pesticide runoff and atmospheric drift, amphibian habitats are likely to be influenced (Lehman and Williams, 2010), and amphibian species encounter great risk due to the high penetration into their skin and their susceptibility to environmental chemicals, especially for tadpoles.

Rana nigromaculata tadpoles is a species of amphibians with wide distribution and abundance in China, where they live in various habitats, including agricultural sites. Thus, damage from toxic ingestion starts at birth. Notably, amphibians are known to prey on crop pests, their movement on agricultural land can reduce the occurrence of rice pests (Teng et al., 2016), and tadpoles are the larvae of amphibians that supply the critical link in the transition between lower and higher trophic levels (Junges et al., 2012). *R. nigromaculata* tadpoles is an excellent model to be used for the assessment of environmental risk due to its naturally low mortality rate with comparatively rapid production times (Zhang et al., 2019). Therefore, it is essential to assess the toxicity of insecticides to *R. nigromaculata* tadpoles.

Neonicotinoids are extensively employed to protect crops and prevent insect pests due to their promising wide spectrum and highly effective pesticidal activity (Frank and Tooker, 2020; Jacob et al., 2019). Imidacloprid [1-(6-chloro-3-pyridylmethyl)-*N*-nitroimidazolidin-2-ylideneamine], a neonicotinoid synthetic insecticide, is a stimulator of postsynaptic nicotinic acetylcholine receptors (nAChRs) (Radwan and Mohamed, 2013). It is the most used seed or soil treatment for some agricultural products, such as rice, cereals, maize and potatoes (Badgajar et al., 2013), and has become a widely used and highly effective insecticide (Rios et al., 2017). Sulfoxaflor [[methyl-oxo-[1-[6-(tri fluoromethyl) pyridin-3-yl] ethyl]-λ6-sulfanylidene] cyanamide], a fourth-generation neonicotinoid, affects the nAChR of insects in a unique way compared with other neonicotinoids, and it is recognized as a significant new tool in insecticide resistance management programs because of its absence of insecticidal cross resistance (Cutler et al., 2013; Jacob et al., 2019).

Researches has shown that with the application of neonicotinoids around the world, neonicotinoids have been detected worldwide in soil and superficial water on account of their elevated levels of solubility in water and extended period of degradation in soil (Yi et al., 2019). Growing evidence indicated that the widespread use of neonicotinoid insecticides may pose potential risks to amphibians (Saka and Tada, 2021; Hrynyk et al., 2018). Our previous study has reported that when exposed to sublethal concentration of sulfoxaflor, sulfoxaflor induced oxidative stress in zebrafish after different times of exposure (Deng et al., 2022), indicating that sulfoxaflor has potential damage to non-target organisms. Several studies have suggested that sulfoxaflor presented a risk to honeybees and sub-lethal concentrations of sulfoxaflor may have caused adverse impacts on ants (Wu et al., 2017; Pan et al., 2017), and sulfoxaflor have acute toxicity and sub-lethal effects to earthworms from oxidative stress and metabolomics (Fang et al., 2018). Meanwhile, researches have reported that imidacloprid exerts sublethal impact on amphibians, for example, imidacloprid inhibited the predator-resistant escape behavior of wood frogs (Lee-Jenkins and Robinson, 2018) and delays the metamorphosis of wood

frogs (Robinson et al., 2017). The metamorphosis is regulated by the amphibian thyroid, and the amphibian thyroid also has the capacity to secrete thyroxine (T3) and triiodothyronine (T4) (Carr and Patiño, 2011), having a critical role in the growth, and metabolism of vertebrates (Heijlen et al., 2013). Studies have demonstrated that insecticides not only cause delays in growth and development, but also change the genes related to TH levels in *Xenopus laevis* tadpoles, leading to the disruption of thyroid endocrine (Li et al., 2016; Liu et al., 2021). However, few studies have been performed to evaluate the impact (especially, sublethal effects) using imidacloprid and sulfoxaflor on *R. nigromaculata* tadpoles.

Hence, this study aimed to evaluate the risk of imidacloprid and sulfoxaflor on *R. nigromaculata* tadpoles. In order to assess the impact of imidacloprid and sulfoxaflor, thyroid hormone (TH) levels and genes which are involved in the metabolic pathway, including and deiodinases (dio2 and dio3) and thyroid hormone receptors (TR α and TR β), and their expression in relation to TH levels were investigated after 28 days exposure. Besides biological indexes including wet weight, snout-to-vent length (SVL), antioxidant enzyme activity and the accumulation of imidacloprid and sulfoxaflor in tadpole was investigated to understand the comprehensive risk to *Rana nigromaculata* tadpole.

2. Materials and methods

2.1. Chemical reagent

Analytical imidacloprid (CAS: 138261-41, 99.70%) and sulfoxaflor (CAS: 946578-00-3, 98.0%) standards were obtained from the Ministry of Agriculture, Tianjin Agricultural Environment Station (Tianjin, China). Commercial assay kits (ELISA) for T3 and T4 were obtained from Shanghai Yaji Biological Technology Co., Ltd. (Shanghai, China). Commercial assay kits for superoxide dismutase (SOD), catalase (CAT), and glutathione S-transferase (GST) were obtained from Suzhou Comin Biotech (Suzhou, China). 3-Aminobenzoic acid ethyl ester (MS-222) was sourced from Sigma-Aldrich (USA). Other analytical reagents were acquired from Youpu Reagent Co., Ltd. (Tianjin, China).

2.2. Experimental animals and treatment

R. nigromaculata (Gosner 25–26) tadpoles were purchased from the State Key Laboratory of Environmental Chemistry and Ecotoxicology, Research Center for Eco Environmental Sciences (Beijing, China). The tadpoles were housed in a 20 L aquarium containing charcoal filtered-tap water, adapting to laboratory requirements where the temperature remained 24 ± 1 °C and a cycle of 12 h of light/12 h of darkness, and were fed 2 times by commercial diets every day (Totoro Supplies, Hong Kong, China). The mortality rate was under 5% during the domestication period and refreshed water at 24 h intervals. All experiments were carried out under review and permission of the Guizhou University Laboratory Animal Ethics Committee with the ethics authorization number EAE-GZU-2019-E003.

2.3. Experimental design

2.3.1. Bioconcentration-elimination of imidacloprid and sulfoxaflor

In bioconcentration experiments of imidacloprid and sulfoxaflor, two exposure concentrations were set in accordance with 1/100 and 1/10 of the results of 96 h LC₅₀. One hundred tadpoles were picked at random and moved into glasses filled with 20 L of the corresponding concentration of imidacloprid and sulfoxaflor, separately. The exposed solution was changed every 2 days, each treatment group was set with three repetitions, and other breeding requirements as mentioned above. Tadpoles were randomly removed from each glass after 1, 3, 5, 7, 9, 14, 21, and 28 days of the exposure enrichment stage and 2 h, and 0.5, 1, 2, 3, 4, and 5 days of the water elimination stage. The tadpole and water samples were stored in a refrigerator at -20 °C for liquid chromatography with tandem mass spectrometry (LC-MS/MS) analysis. The bioconcentration and elimination test were carried out according to the guideline of OECD 305 (OECD, 2011).

2.3.2. Acute exposure experiments

Tadpoles were selected and placed randomly into a 5-L flask with 4-L of filtered solution. Imidacloprid and sulfoxaflor solvents were made up with concentration 90–300 mg/L, and 244–645 mg/L, respectively. The concentrations leading to the death of 50% of the experimental groups (LC₅₀) was documented. The whole process was conducted on basis of instructions for testing the environmental security of pesticides in accordance with the OECD guideline 203 (fish, acute toxicity test) (OECD, 2019) and GB/T 31270.18–2014 (amphibian acute toxicity test) (AQSIQ and SAC, 2014).

2.3.3. Toxicity experiment

Sixty tadpoles were picked at random and put in each aquarium with 20 L of the test solution of imidacloprid and sulfoxaflor. The exposure concentration for the toxicity test were 1.74, 17.36 mg/L, and 4.27, 42.73 mg/L (1/100 and 1/10 of the LC₅₀ of tadpole, respectively). Dechlorinated tap water was set as a control, and every treatment had three replicates. The exposures were conducted under the laboratory conditions mentioned above. The exposed solution was to be renewed at 3-day intervals. the tadpoles were fed twice a day. After 1, 7, 14, 21 and 28 days of exposure, the wet weight and SVL of tad-

poles were measured, antioxidant enzymes were determined after anesthetization in MS-222 (100 mg/L). After 28 days of exposure, tadpoles were randomly selected and anesthetized using MS-222 (100 mg/L) for TH measurement and gene expression.

2.3.4. Measurement of antioxidant enzyme activities

Ten tadpoles from each aquarium were selected for the analysis (n = 3). After 1, 7, 14, 21 and 28 days of exposure, MS-222 was employed to anesthetize the tadpoles. Tadpoles were homogenized, then centrifuged for 10 min (10,000 g, 4 °C). The supernatant was employed to determine the SOD, CAT and GST activities by using assay kits (Suzhou Comin Biotech).

2.3.5. Levels of TH

After 28 days of exposure, five tadpoles per aquarium were euthanized using MS-222, weighed and placed into a 10 mL centrifuge tube. A proper amount of normal saline was put to the centrifuge tube, homogenized in an ice water bath for 3 min, and then centrifuged for 10 min (4 °C, 3,000 g). Subsequently, T3 and T4 in the supernatant were measured using the ELISA kit.

2.3.6. cDNA synthesis and real-time RT-PCR

All tadpoles were dissected and gathered and TRIzol reagent (Tiangen Biotech, Beijing, China) was employed to extract the total RNA of tadpoles after 28 days of exposure. The concentration of total RNA was determined using a micro spectrophotometer with absorbance at 260 nm (DS-11, 164 DeNovix Inc, USA), and then gel electrophoresis in 1% agarose gel was used to verify the RNA quality. Then RNA was reversely transferred to cDNA in accordance with the manufacturer's instructions with a FastQuant RT kit (Tiangen Biotech, Beijing, China).

A SLightCycler480 II system (Roche, Rotkreuz, Switzerland) was used to confirm and quantify the changes in gene expression. Specific primers are provided in Table 1. Amplification was performed following the protocol of the manufacturer to identify these genes for specific amplification with PCR requirements of 95 °C for 3 min, then 45 cycles of 95 °C for 5 s of annealing at different temperatures (Table 1) and 60 °C for 30 s. The expression of ribosomal protein L8 was normalized using the 2^{-ΔΔCt} method to determine the fold change in gene expression.

Table 1 Real-Time PCR Primers and Conditions.

Gene	Genbank accession	Primer sequences (5' to 3')	Size (bp)	Annealing (°C)
<i>RPL13A</i>	MG844184	Forward: TACCGCAACAACTCAAATACC Revers: CAAAGACCTTCAGTCGCTCC	175	60
<i>dio2</i>	MH892455	Forward: TGCCTACAAACAGGTGAAGCT Revers: CCAAAGTTGACAACAAGAGGG	150	60
<i>dio3</i>	MH892456	Forward: GAGTCCCTGAAGGCGGTCT Revers: AGTCCAGGATCCGGCACA	127	60
<i>TRα</i>	KC139354.1	Forward: GGGGTGGTCTCAGATGCTATAT Revers: TTCAAAGGCGAGGAGGTAAG	174	60
<i>TRβ</i>	KC139355.1	Forward: AAGTGAGACCTTAACGCTGAATG Revers: TGAAGACAGTGATACGCCAA	115	60

2.4. Extraction and analysis of imidacloprid and sulfoxaflor

2.4.1. Sample pretreatment

The water sample was filtered using a nylon filter with 0.22 μm and moved to a vial for the determination of imidacloprid and sulfoxaflor without pretreatment.

For the tadpole samples, 0.20 g of tadpoles were weighed, added to 2 mL of methanol in a plastic centrifuge tube and ground with an electric tissue homogenizer. The tadpole samples were vortexed at 2500 rpm/min for 5 min. Purification agents were added to the samples exposed to imidacloprid (0.2 g MgSO_4) and sulfoxaflor (0.2 g MgSO_4 and 0.1 g NaCl), then vortexed for 3 min, separately. The samples were collected by centrifugation for 5 min (8000 rpm/min, 4 $^\circ\text{C}$), then 1.5 mL of top-serum was removed and pipetted into round-bottomed flask, and evaporated by rotation (40 $^\circ\text{C}$ water bath) until vacuum dried. The evaporated residue was dissolved by the addition of acetonitrile (1.5 mL), supernatant was placed in plastic centrifuge tube, 100 mg of PSA was added, and the sample was swirled for 1 min and the centrifugation was carried out for 3 min. The solution was passed with a 0.22 μm nylon filter and moved to a vial with a lined tube, then analyzed by LC-MS/MS.

2.4.2. Quantification analysis

Imidacloprid and sulfoxaflor in *R. nigromaculata* tadpoles and water were analyzed by liquid chromatography-mass spectrometry LC-MS/MS. The detailed analytical methods for imidacloprid and sulfoxaflor are given in [Supporting Information Test S1](#).

2.5. Statistical analyses

IBM SPSS 26.0 was employed for statistics analysis. For SVL, weight, TH levels, gene expression, and antioxidant enzyme activity, differences in statistics with respect to the exposure and control groups were analyzed by one-way ANOVA, with various letters denoting each statistical difference with respect

to the exposure and control groups ($p < 0.05$). The probit equation was performed to calculate the LC_{50} and associated 95% confidence intervals. The results presented the mean standard deviation (SD). The bioconcentration factor (BCF) of imidacloprid and sulfoxaflor in tadpoles was calculated using the following equation:

$$\text{BCF} = C_t / C_w$$

where C_t (mg/kg) denoted the concentration of imidacloprid and sulfoxaflor in tadpoles, C_w (mg/L) denoted concentration of imidacloprid and sulfoxaflor in the water.

3. Results and discussion

3.1. Validation of the LC-MS/MS method

Recovery experiments of imidacloprid and sulfoxaflor were carried out. The limit of quantification (LOQ) for every target compound is computed from the signal-to-noise ratio (S/N) around the target peak, defined as 10 times the S/N ([Guo et al., 2022](#)). The LOQ of imidacloprid and sulfoxaflor of the method for tadpole was 0.05 mg/kg and 0.3 mg/kg, respectively. The method provided average recoveries of imidacloprid and sulfoxaflor from 76.18% and 96.82%, with relative standard deviations (RSD) from 1.75% to 15.41%, from 82.51% to 112.51%, with RSD from 1.06% to 5.79% in the tadpole, respectively ([Table S3 and S5](#)). The calibration curve with $R^2 > 0.993$, was shown in [Tables S2 and S4](#). In conclusion, the approach is well suited for imidacloprid and sulfoxaflor determination.

3.2. Bioconcentration and elimination of imidacloprid and sulfoxaflor

The concentration of imidacloprid and sulfoxaflor in the water was maintained at a constant level as the solution was renewed every two days ([Fig. S1](#)). In our study, it was shown that imidacloprid can accumulate rapidly during the

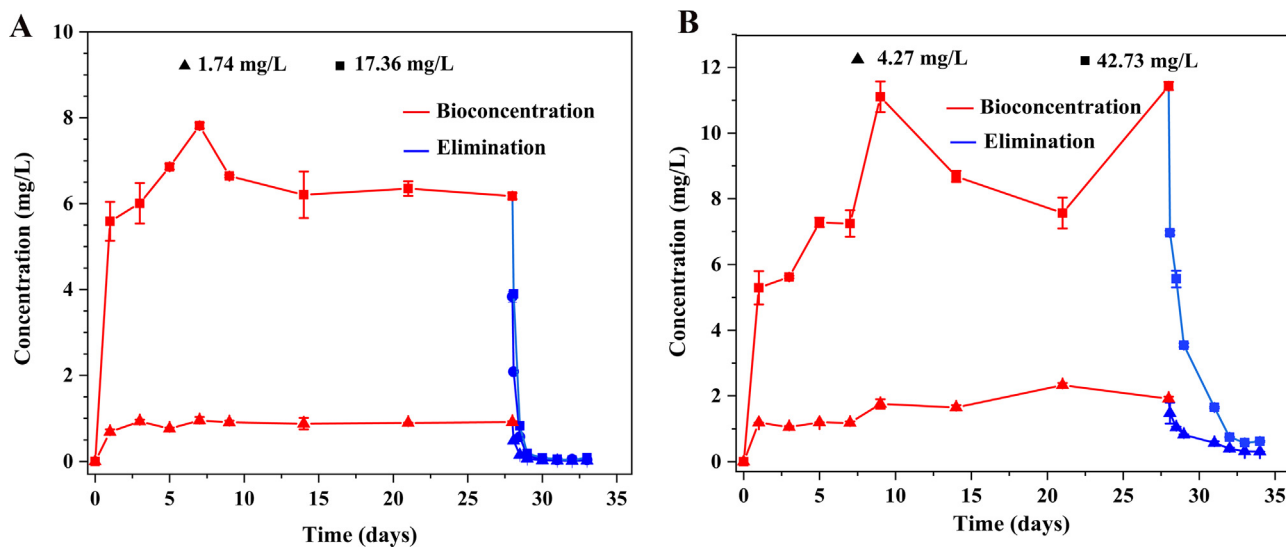


Fig. 1 Concentrations of imidacloprid and sulfoxaflor in tadpoles (A) 1.74 and 17.36 mg/L, (B) 4.27 and 42.73 mg/L. (Error bars represent the mean \pm SD).

biological uptake period and the concentrations of 1.74 and 17.36 mg/L in tadpoles reached the highest level at 7th and 6th days, then kept at a relative steady state until the 28th day (Fig. 1A). The elimination process was not fast, and the concentration of imidacloprid in tadpole at 1.74 and 17.36 mg/L reached the lowest values of 0.015 and 0.08 mg/kg respectively in 5 days after the tadpoles were moved to clean water (Fig. 1A). With the enrichment of sulfoxaflor with the concentration of 4.27 and 42.73 mg/L in tadpoles, the concentrations increased to the level of maxi-

mum at 21th and 9th days. The elimination process shown that the concentration of sulfoxaflor in tadpoles at 4.27 and 42.73 mg/L reached to the lowest value of 0.31 and 0.61 mg/kg, respectively, in 5 days after the tadpoles were moved to clean water (Fig. 1B). As a result, the research showed that imidacloprid can accumulate rapidly in tadpoles shortly after exposure to sublethal concentrations compared with sulfoxaflor. We also inferred that imidacloprid and sulfoxaflor may continue to decline to undetectable levels in tadpoles over a longer period of time.

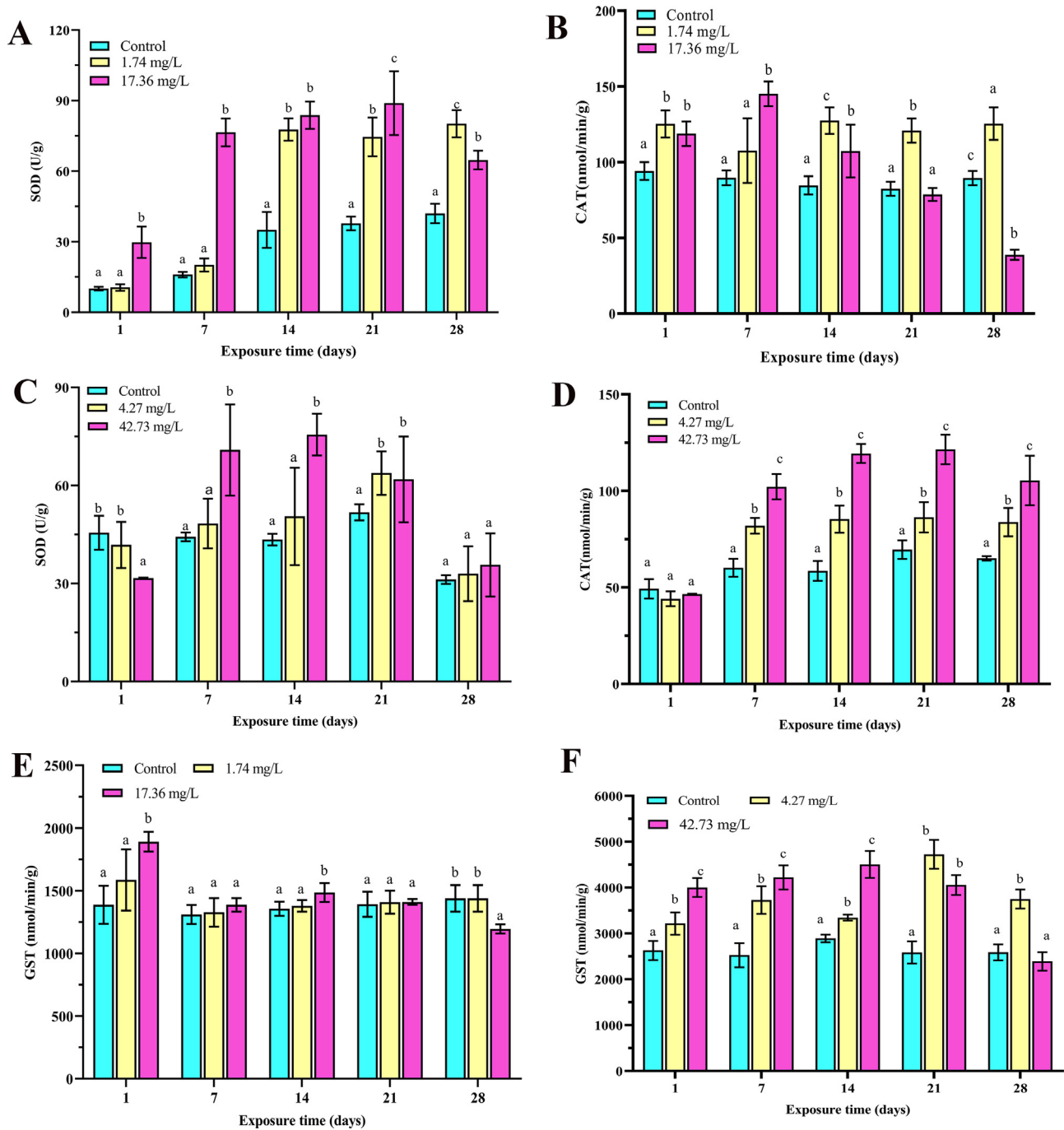


Fig. 2 Activity of SOD, CAT and GST in *R. nigromaculata* tadpoles with exposure to imidacloprid and sulfoxaflor after 1, 7, 14, 21 and 28 days (n = 6). Significant differences between treatments are indicated by different letters, $p < 0.05$. Error bars represent SDs.

The BCFs of imidacloprid at 1.74 and 17.36 mg/L were 0.559 and 0.39, the BCFs of sulfoxaflor at 4.27 and 42.73 mg/L were 0.359 and 0.191, respectively, these values indicated that imidacloprid and sulfoxaflor have potential accumulation in tadpoles.

3.3. Acute toxicity

For the assessment of the effect of imidacloprid and sulfoxaflor on acute toxicity in *R. nigromaculata* tadpoles. 96 h-LC₅₀ of imidacloprid and sulfoxaflor was documented. The 96 h-LC₅₀ of imidacloprid and sulfoxaflor was 173.55 and 427.37 mg/L, respectively (Table S1), indicating that imidacloprid have greater toxicity for *R. nigromaculata* tadpoles than sulfoxaflor. On the basis of their LC₅₀, imidacloprid and sulfoxaflor could be categorized as low-toxicity pesticides to *R. nigromaculata* tadpoles. Although imidacloprid and sulfoxaflor had low toxicity, our previous study showed that the subacute sulfoxaflor posed impact to organisms (Deng et al., 2022). Therefore, the subacute influence of sulfoxaflor and imidacloprid on *R. nigromaculata* tadpoles should be assessed.

3.4. Oxidative stress effects

The first line of cellular response to antioxidants is catalyzing the disproportionation of O₂^{•-} to H₂O₂ and O₂, and to prevent the production of superoxide anion radicals by removing reactive oxygen species (ROS) via SOD (Sies, 1997). CAT subsequently degrades H₂O₂ into oxygen and water. SOD and CAT perform a key role in the antioxidant defense system, with their task being to defend from oxidative stress (Menon and Rozman, 2007). In our research, the SOD activity was significantly increased in exposure of imidacloprid compared with those control group (Fig. 2A, $p < 0.05$), compared with the SOD activity under the treatment at 1.74 mg/L, the activity was increased in 17.36 mg/L (1, 7, 14, and 21 days), exhibited a time-dependent increase but decreased at 28 days (Fig. 2A, $p < 0.05$). The CAT activity in 17.36 mg/L was significant increased compared with control (1, 7, 14, 21 and 28 days) (Fig. 2B, $p < 0.05$), and compared with the CAT activity under the treatment at 1.74 mg/L, the activity was decreased in 17.36 mg/L (14, 21 and 28 days) (Fig. 2B, $p < 0.05$). The SOD activity was significantly increased compared with the control in the exposure of sulfoxaflor (7, 14, and 21 days,

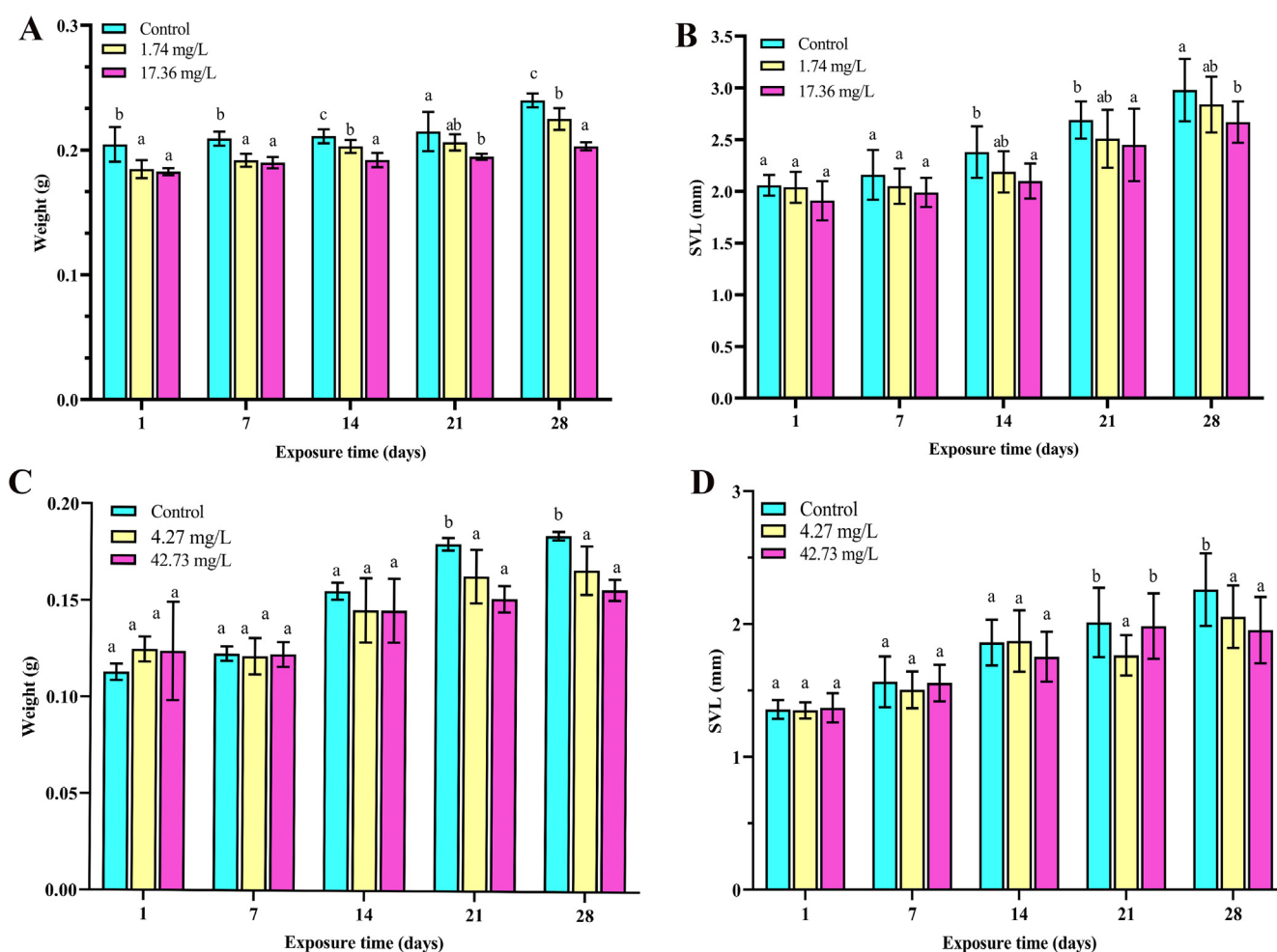


Fig. 3 Developmental endpoints of *R. nigromaculata* tadpoles exposed to different concentrations of imidacloprid and sulfoxaflor for 1, 7, 14, 21 and 28 days ($n = 14$). Mean weight (A C), SVL (B D). Values represent mean \pm SD. Significant differences between treatments are indicated by different letters, $p < 0.05$. Error bars represent SDs.

$p < 0.05$) (Fig. 2C). The CAT activity was all significantly increased in comparison with that under other exposure of sulfoxaflor (Fig. 2D, $p < 0.05$), and exhibited a time-dependent increase (7, 14, 21, and 28 days).

The increase in SOD and CAT activity in the exposure of imidacloprid and sulfoxaflor may be consequence of the increase in ROS, which facilitates the activation of the enzymes (Pinya et al., 2016). Meanwhile, we discovered that higher doses of imidacloprid reduced the SOD activity or CAT activity, which might be attributed to an excessive accumulation of ROS and the suppression of enzymatic activity, resulting in reduced defense response (Rodríguez-Serrano et al., 2006). In return, the enzymes of SOD and CAT were unable to combat the high levels of oxidative stress due to the increased concentrations of imidacloprid, and over oxidative stress might conversely render SOD or CAT inactive activity (Bagnyukova et al., 2006). Researches have reported inhibition of CAT activity in tadpoles with different insecticide exposures (Rutkoski et al., 2021; Sun et al., 2014), which was similar with our results.

GST is a second-stage detoxification enzyme known to catalyze the binding of glutathione (GSH) to exogenous substances and their metabolites (Wu et al., 2007). As for imidacloprid, the concentrations of GST were significantly increased when exposed to 17.36 mg/L (1 and 14 days, $p < 0.05$) (Fig. 2E). The concentrations of GST were significantly increased in tadpoles exposed to sulfoxaflor in all treatment groups when compared with that in the control (1, 7, 14, 21 and 28 days, $p < 0.05$) (Fig. 2F). The increase in GST activ-

ity may be due to the induction of detoxification by imidacloprid and sulfoxaflor through binding intracellular GSH and harmful metabolites, and imidacloprid and sulfoxaflor induced excessive ROS and themselves may also contribute to the activation of GST. However, when exposed to 28 days, the concentrations of GST exposed to imidacloprid were significantly decreased in 17.36 mg/L ($p < 0.05$) (Fig. 2E). For sulfoxaflor exposure, the GST levels were significantly decreased to 42.73 mg/L at 21 and 28 days when compared with that in 4.27 mg/L ($p < 0.05$) (Fig. 2F). The study explained that this reduction in activity of the enzyme appears to be associated with an overconsumption of GSH which serves as a precursor and variation in GST formation induced by various intermediary metabolites (Ge et al., 2015). In this research, the GST activity of tadpoles was affected by imidacloprid and sulfoxaflor, indicating that the GST system is probably the pathway of detoxification from imidacloprid and sulfoxaflor in tadpoles.

3.5. Growth index and TH

3.5.1. Growth indexes

The inhibition of growth is a biomarker of toxicity development (Xu and Huang, 2017). In this research, the effects on biological indexes (weight, SVL) were evaluated when tadpoles were exposed to imidacloprid and sulfoxaflor. For the imidacloprid-treated group, the weight of imidacloprid exposure was significantly decrease ($p < 0.05$) in comparison to that for control (Fig. 3A) and SVL was significantly decreased

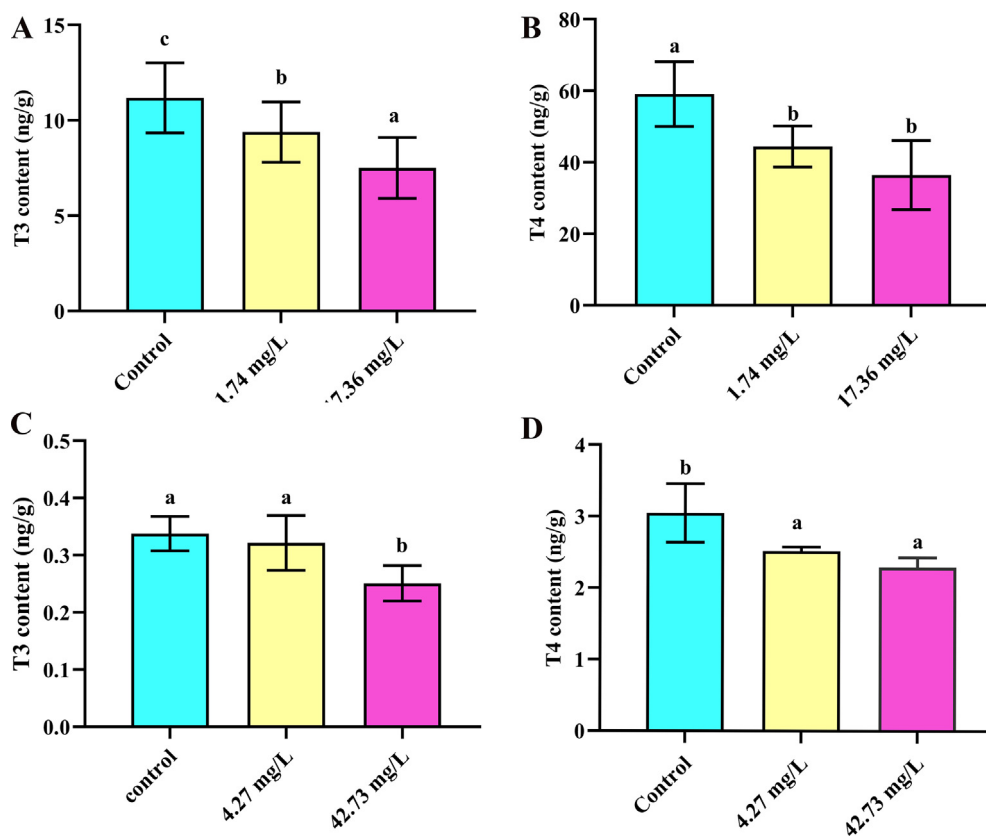


Fig. 4 T3 and T4 levels in *R. nigromaculata* tadpoles exposed to imidacloprid and sulfoxaflor for 28 days ($n = 6$). Significant differences between treatments are indicated by different letters, $p < 0.05$. Error bars represent SDs.

(14, 21, and 28 days, $p < 0.05$) (Fig. 3B). Weight was significantly decrease in all treatments of sulfoxaflor at 21 and 28 days compared with that in the control (Fig. 3C), SVL at 14 days was not significantly difference from that of the control, decreasing at 21 days only by 4.27 mg/L exposure ($p < 0.05$), and SVL was significantly decreased at 28 days in all treatments of sulfoxaflor ($p < 0.05$) (Fig. 3D). Some studies have reported that development and metamorphosis in *Xenopus laevis* were altered by exposing to insecticides (Liu et al., 2021), and the weight of *R. nigromaculata* tadpoles was significantly decreased after 28 days of exposure to triadimefon (Zhang et al., 2018), which was in accordance with our findings. In our study, reduction in SVL and weight exposed to sulfoxaflor and imidacloprid in all treatments after 28 days of exposure may be ascribed to disturbances in metabolism of the thyroid.

3.5.2. Levels of thyroid hormone

The influence of imidacloprid and sulfoxaflor on T3 and T4 concentrations was evaluated. T3 concentrations were observed to decrease significantly during exposure to all treatments of imidacloprid, and T4 concentrations were also significantly decreased in all treatments compared with control ($p < 0.05$) (Fig. 4A–B). When exposed to 42.73 mg/L of sulfoxaflor, the levels of T3 and T4 were decreased significantly compared with the control ($p < 0.05$) (Fig. 4C–D). These results clearly showed an endocrine-disrupting effect of imida-

cloprid and sulfoxaflor on the thyroid system of tadpoles. In our study, significant suppression of metamorphosis development (SVL and weight) in parallel with a decrease in T3 and T4 levels in 28 days was noticed, demonstrating that suppressive effect of imidacloprid and sulfoxaflor on metamorphosis might be attributed to reduction of thyroid hormones levels. A previous study showed that imidacloprid altered TH levels of lizards (Wang et al., 2020). Above all, we concluded that the T3 and T4 synthesis in *R. nigromaculata* tadpoles was inhibited by imidacloprid and sulfoxaflor.

3.5.3. Relative expression of TH-dependent genes

The mRNA levels of *dio2* and *dio3* (encoded as iodothyronine deiodinase (DIO2 and 3)). Type 2 deiodinase plays an essential part in transforming T4 into the more inactive T3, while T4 and T3 are inactivated to the bio-inactive inverted T3 and T2 by type 3 deiodinase, separately (Lorenz et al., 2018). As for imidacloprid, the levels of *dio2* gene expression increased at 1.74 mg/L, and significantly downregulated at 17.36 mg/L ($p < 0.05$) (Fig. 5A). The *dio3* gene expression was significantly upregulated at in all treatments ($p < 0.05$) (Fig. 5B). The results revealed that imidacloprid reduced the *dio2* gene expression, which decreased the conversion from T4 to T3 leading to a decrease in T3 levels. In contrast with the results for *dio2*, the expression of *dio3* gene was increased, which inactivated T4 and T3. The downregulation of *dio2* expression and the upregulation of *dio3* expression by 17.36 mg/L of imi-

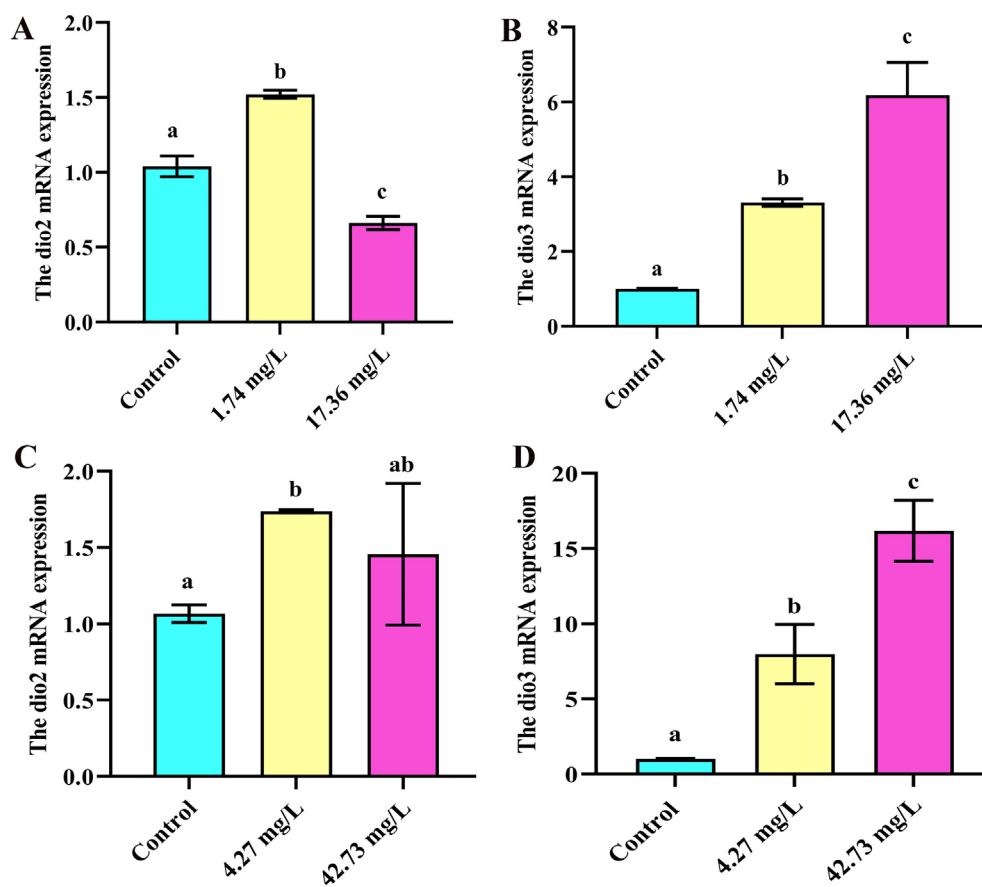


Fig. 5 Expression of *dio2* and *dio3* exposed to imidacloprid and sulfoxaflor after 28 days ($n = 3$). Significant differences between treatments are indicated by different letters, $p < 0.05$. Error bars represent SDs.

clodolprid were in accordance with the inhibition of the T3 and T4 we detected. When exposed to 42.73 mg/L of sulfoxaflor, compared with the control, the *dio2* gene expression was not significantly different ($p < 0.05$, Fig. 5C). Therefore, there was no significant effect of sulfoxaflor on *dio2* gene expression at 42.73 mg/L. However, compared with the control, during exposure to 4.27 mg/L, an upregulation was observed ($p < 0.05$, Fig. 5C). After 28 days exposure, the expression of the *dio3* gene was remarkably upregulated in all treatments ($p < 0.05$) (Fig. 5D). The *dio2* expression had no significant changes, and the upregulation of *dio3* was in accordance with the levels of T3 and T4 we detected. On the contrary, a previous study observed a downregulation of *dio3* in *R. nigromaculata* tadpoles after 21 days of triazolone exposure, which was likely related to the observed recovery of T3 concentrations (Li et al., 2016).

In our study, the *dio3* expression inactivates T4 and T3 and the elevated *dio3* expression levels contribute to reduction in TH levels in the target tissue. TH is indispensable to normal growth. For instance, tail muscles die as a result of TH induction; limb muscles grow and differentiate in response to the hormone (Brown and Cai, 2007). As a result, reducing the TH levels in the target tissue might result in suppression of both growth and metamorphosis. On basis of the results of the expression of *dio2* and *dio3*, we inferred that imidacloprid and sulfoxaflor disrupt HPT regulation on the genetic level.

For all vertebrates, it is known that two thyroid hormone receptors (TR), called TR α and TR β , affect the development of organisms primarily by combining with the TR (Darras et al., 2011). Abnormalities in TR α and TR β might contribute to a cascade in which THs fail to associate with TRs and activate post-receptor reactions (Yu et al., 2013). In the present study, imidacloprid and sulfoxaflor exposure had a negative effect on TR α expression. When exposed to imidacloprid, gene expression levels of TR α were significantly decreased to 17.36 mg/L compared with the control ($p < 0.05$) (Fig. 6A), and gene expression levels of TR α showed an increase to 4.27 mg/L and 42.73 mg/L in sulfoxaflor treatments. ($p < 0.05$) (Fig. 6C). The TR α mRNA levels have a significant change in zebrafish (Wang et al., 2020), which was similar with our results. Imidacloprid and sulfoxaflor exposure also had a negative effect on TR β expression. In particular, the level of expressions of TR β was decreased in all treatments of imidacloprid ($p < 0.05$) (Fig. 6B). Likewise, there was significantly downregulated TR β expression detected in lizards, which was same as our results (Wang et al., 2020). As for sulfoxaflor exposure, the expression of TR β was dramatically downregulated in all treatments compared with control group ($p < 0.05$), and groups have no significant difference from each other (Fig. 6D). The reduced TR β in tadpoles was associated with the delayed metamorphic response (Navarro-Martín et al., 2014). Accordingly, we concluded that sulfox-

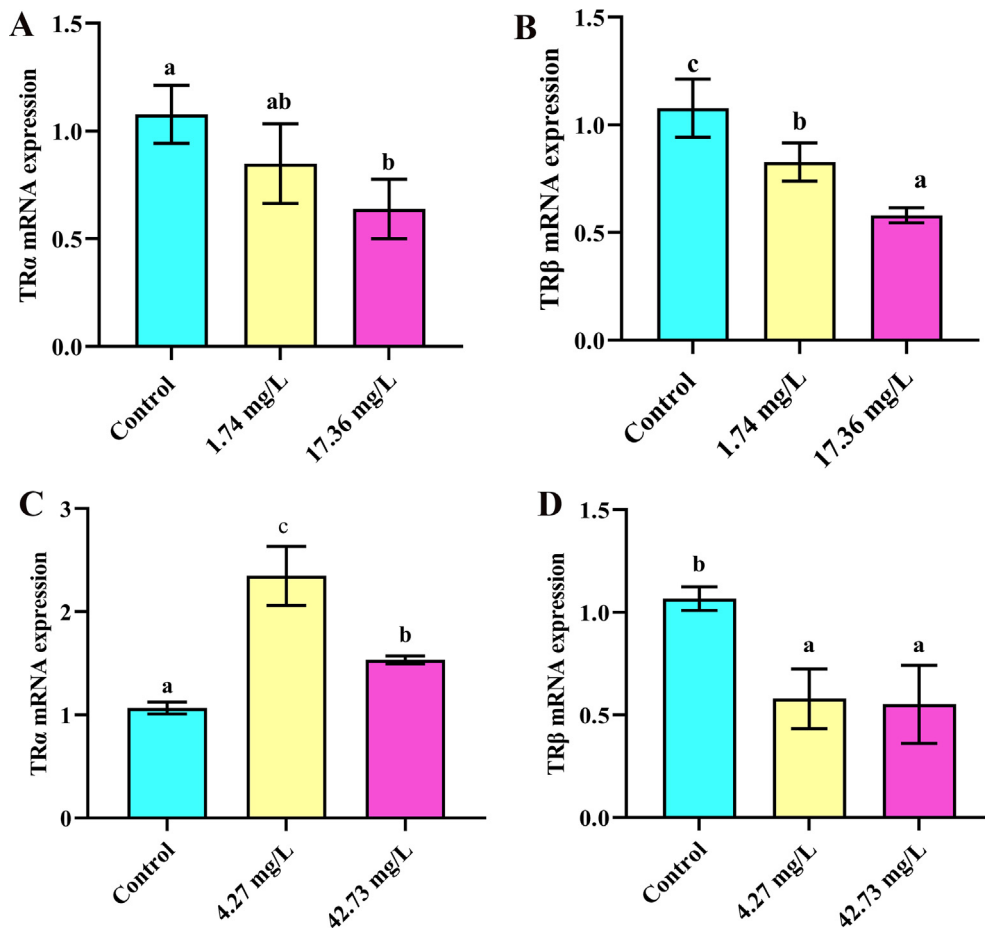


Fig. 6 TR α and TR β expression exposed to imidacloprid and sulfoxaflor after 28 days ($n = 3$). Significant differences between treatments are indicated by different letters, $p < 0.05$. Error bars represent SDs.

aflor and imidacloprid may delay metamorphosis through controlling the expression of TR β and have an influence on the endocrine system.

4. Conclusion

In the present study, the aim of the study was to investigate the impact of exposure to imidacloprid and sulfoxaflor on *R. nigromaculata* tadpoles. Our results indicated that imidacloprid and sulfoxaflor have shown low-toxicity to *R. nigromaculata* tadpoles based on the acute toxicity. Oxidative stress was caused when exposed to imidacloprid and sulfoxaflor, respectively, and imidacloprid and sulfoxaflor inhibited the growth of *R. nigromaculata* tadpoles due to decreased TH levels. Although exposure to imidacloprid and sulfoxaflor showed same trend of the dio3 and TR β gene expression, there was no significant effect of sulfoxaflor on dio2 gene expression exposed to high level compared with imidacloprid, the gene expression levels of TR α showed an increase in sulfoxaflor but decreased in imidacloprid treatments. On basis of the results of the expression of dio2 and dio3, we inferred that imidacloprid and sulfoxaflor disrupt HPT regulation on the genetic level, and imidacloprid and sulfoxaflor may delay metamorphosis through controlling the expression of TR β and have an influence on the endocrine system. In sum, imidacloprid and sulfoxaflor probably have negative impacts on the development of *R. nigromaculata* tadpoles through disruption of the thyroid hormone pathway. This research may contribute to more stringent management of the levels of insecticides in the aquatic environment, providing support for maintaining a healthy ecosystem.

CRedit authorship contribution statement

Xia Zhou: Writing – original draft, Visualization, Data curation. **Yao Deng:** Formal analysis, Investigation. **Ran Wang:** Methodology. **Fang Wang:** Investigation. **Honghao Cui:** Investigation. **Deyu Hu:** Funding acquisition, Supervision, Project administration. **Ping Lu:** Funding acquisition, Supervision, Project administration.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary material

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.arabjc.2023.104723>.

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