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# Exposure to butylparaben induces oxidative stress and apoptosis in the liver of *Polypedates megacephalus* tadpole

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Butylparaben (BuP) is a ubiquitous preservative in cosmetics, foods, and medicine. As an emerging pollutant, it has gained wide attention in recent years. Because amphibians have both aquatic and terrestrial life stages and high skin permeability, they are sensitive to environmental changes. To assess the toxic effects of BuP on amphibians, the present study was conducted on the spot-legged tree frog (*Polypedates megacephalus*) tadpoles. In the present study, LC<sub>50</sub> of BuP for 96 h exposures was 3509 µg/L. Then, the tadpoles were exposed to 0, 350 and 2100 µg/L BuP (hereafter named as CK, B1 and B2) for 12, 24 and 48 h, the oxidative stress and apoptosis in the liver were analyzed. The results showed that the content of MDA and activity of SOD and GSH-PX were increased with an increase in BuP exposure. BuP stress significantly altered the expression of Nrf2-Keap1 signal pathway genes. An increase in the expression of JAK-STAT signal pathway genes was observed with an increase in the exposure to BuP concentration at 24 and 48 h. In contrast, the expression of negative regulators of JAK-STAT signal pathway were significantly increased at 12 h and decreased at 24 and 48 h. A significant decrease in pro-apoptosis genes expression (*Bax* and *Caspase3*) were observed at 12 h but increased at 24 and 48 h, while the expression of anti-apoptosis gene (*Bcl-2*) decreased at 24 and 48 h. These results provide a valuable reference point for assessing the ecological and health risks associated with BuP in the environment.

## KEYWORDS

butylparaben, *Polypedates megacephalus* tadpoles, antioxidant enzymes, apoptosis, JAK-STAT signal pathway

# 1 Introduction

Amphibians stand at the forefront of a global biodiversity crisis (Whitfield et al., 2007). According to the second Global Amphibian Assessment findings, amphibians are the most threatened class of vertebrates (40.7% of species are globally imperiled) (Luedtke et al., 2023). The number of extinct amphibian species was 23 by 1980, and four times more by 2022. Among these factors resulting in the decline of the amphibian population, water pollution is one of the leading causes (Park and Do, 2023). Because of the hypertonicity of the skin, external fertilization and embryonic development, amphibians are sensitive to the pollutants in water, which will affect their physiological activities and growth (Brühl et al., 2011). Some studies have shown that the sharp decline in the population of the northern dusky salamander (*Desmognathus fuscus fuscus*) population in the Acadia National Park (Maine, USA) has been linked to mercury and aluminum pollution, and the decline of the mountain yellow-legged frog (*Rana muscosa*) population was caused by airborne pesticide pollutions (Slaby et al., 2019). Therefore, the relationship between water pollution and amphibian population is of great concern.

Parabens are used in a variety of food, pharmaceuticals and personal care products due to their broad spectrum of antimicrobial activity and their preservative properties. They are esters of p-hydroxybenzoic acid with alkyl substituents including methylparaben (MeP), ethylparaben (EtP), propylparaben (PrP), butylparaben (BuP) and pentylparaben (PeP) (Darbre et al., 2004). Traditionally seen as harmless, non-polluting, and non-irritating, parabens have recently been scrutinized for possible toxicity. Recently, more and more research has focusing on the ubiquitous presence of BuP, their environmental metabolites and high levels of parabens in the human body. Among alkyl parabens, BuP is the most powerful and has the most substantial estrogenic effects (Moreno-Marenco et al., 2020). In the meanwhile, toxicity studies including BuP become more common, because BuP is readily absorbed and maintained in bodily tissues. Ingestion, inhalation, or skin absorption of BuP can cause oxidative stress and harm the endocrine system (Belenguer et al., 2014). It can alter enzyme activity and cellular homeostasis inducing organ toxicity and damaging tissues through oxidative imbalance. This leads to damage to lipids, DNA, and proteins, which in turn triggers the development of various diseases (Aydemir et al., 2019). Some studies on BuP-induced oxidative stress mainly focused on mice liver, human spermatozoa, rat testis and turtle liver (Samarasinghe et al., 2018; Schreiber et al., 2019). There is still a lack of available data on the effects of BuP on the early life stages of amphibians.

Amphibians are very susceptible to environmental changes and contaminants in the aquatic ecosystem because of their porous skin, which is essential for gas exchange and osmotic balance (Linder et al., 2010). Compared to other model organisms like zebrafish, amphibians have not been the subject of many ecotoxicological studies, despite being excellent bioindicator species (Burgos-Aceves et al., 2022; Castro-Castellon et al., 2022; Dong et al., 2022). Furthermore, water quality standards developed based on information on fish toxicity do not protect amphibians

(Sparling et al., 2010). Numerous studies have demonstrated that the physiological indices associated with oxidative stress in aquatic animals may serve as an invaluable foundation for tracking water pollution levels and assessing the biological safety of contaminants. It can be used as a bioindicator species for assessing environmental risk. In this study, the spot-legged tree frog *Polypedates megacephalus* tadpole was chosen to study the toxicity of BuP because it widely distributes throughout subtropical and tropical areas such as China, India, northern Vietnam, and so on (Zug, 2022). The study aimed to ascertain the detrimental effects and threshold concentrations of BuP on *P. megacephalus* tadpoles. Firstly, the LC<sub>50</sub> concentration of BuP in the tadpoles was studied. Secondly, according to the LC<sub>50</sub> concentration, the responses of the antioxidant system and oxidative damage in the livers of the tadpoles after 12, 24 and 48 h of exposure to BuP were analyzed. The present investigation offers factual support for evaluating the possible ecological hazards associated with BuP in amphibians.

## 2 Materials and methods

### 2.1 Animals culture and handling

In July, 2021, *P. megacephalus* tadpoles were collected from the puddles in the vegetable garden near Dongzhai Harbor (Haikou, Hainan, China). The tadpoles were acclimated in 5 L glass tanks with dechlorinated water for at least 3 d before the experiment at Hainan Normal University, and the individuals at Gosner stage (GS) 25 (Gosner, 1960) were chosen to study. The egg yolk was fed to them daily, and the water was changed. The water temperature was  $28 \pm 1^\circ\text{C}$  and the photoperiod varied naturally. The pH values of the culture water were maintained at  $7.47 \pm 0.12$ , and the dissolved oxygen concentration was  $8.32 \pm 0.25$  mg/L.

BuP [CAS: 94-26-8, Purity: 99%] was acquired from Macklin (Shanghai, China). For complete dissolution, 75% ethanol was added into the BuP solution, ensuring that the ethanol concentration in the water did not exceed 1  $\mu\text{L/L}$ . The preliminary experiment indicated that the ethanol concentration of ethanol < 1  $\mu\text{L/L}$  was safe for the tadpoles.

Approval for this study was obtained from the ethics committee overseeing experimental animal research at the Hainan Ecological and Environmental Education Center (No: HNECEE-2020-003).

### 2.2 LC<sub>50</sub> of BuP to the tadpoles

Tadpoles were exposed to five groups with BuP concentrations of 0, 400, 800, 1600, 3200 and 6400  $\mu\text{g/L}$ , respectively, and the exact concentrations of BuP were 0, 398.21, 789.54, 1602.36, 3195.47 and 6394.62, respectively, which were analyzed by HPLC (Agilent 1260, USA). The live and dead tadpoles were scored after exposure, and the percent mortality was determined. The experiment was repeated thrice independently, using 20 tadpoles of Stage 25 per treatment group. EPA Probit Analysis Program (Version 1.5) was employed to determine LC<sub>50</sub>.

## 2.3 Acute exposure of BuP to the tadpoles

The 96 h-LC<sub>50</sub> of BuP to the tadpoles, the safe concentration (SC) = 0.1 × LC<sub>50, 96 h</sub> = 350 µg/L (Zang et al., 1993), a higher concentration group used 6-fold SC is 2100 µg/L. So, three groups of 0, 350 and 2100 µg/L BuP exposure concentration (for all the other assays listed in this paper as CK, B1 and B2) were set up. To ensure the experimental integrity, an equal amount of alcohol in the group of 2100 µg/L is added into the other two groups (0 and 350 µg/L). The exact concentrations of BuP in the groups of 350 and 2100 µg/L groups were 348.21 and 2103.01, respectively, which were analyzed by HPLC (Agilent 1260, USA). Water parameters were monitored daily and adjusted as necessary. Thirty-six tadpoles were randomly selected in each group. Liver samples were collected after exposure to BuP for 12, 24 and 48 h.

## 2.4 Biochemical analysis

Following the procedures described by Liang et al. (2020), 0.1 g of liver tissue was homogenized with 1ml ddH<sub>2</sub>O in a high-throughput tissue grinder (CIENZ-48, Ningbo, China). The homogenate underwent centrifugation at 3,000 rpm for 10 min at 4°C, and the resulting supernatant was retained for biochemical analysis. The customized ELISA kits produced by Nanjing Jiancheng Bioengineering Institute (Nanjing, China) were used to measure the activities of superoxide dismutase (SOD), malondialdehyde (MDA) and glutathione reductase (GSH-PX) in the liver.

## 2.5 RNA extraction and qRT-PCR

The liver samples were extracted by TriZol (Invitrogen, Carlsbad, USA) according to the standard manufacturer's protocol. The RNA was dissolved in 0.1% DEPC water, and its integrity and concentration were measured by 1.2% agarose gel electrophoresis and an ultramicro spectrophotometer (Thermo Fisher, USA). The RNA was kept in an ultra-low temperature refrigerator set at -80°C. The cDNA was synthesized according to the instructions of the manufacturer of the Prime Script Reverse Transcriptase (Takara, Tokyo, Japan). A two-step procedure following the SYBR<sup>®</sup>PrimeScript<sup>™</sup> Kit (Takara, Tokyo, Japan) instructions was used to detect the mRNA expression levels of some genes. The primers were designed using the NCBI online primer design tool and the target genes are listed in Table 1. Relative expression was calculated using the 2<sup>-ΔΔCt</sup> method with *β-actin* as the internal control.

## 2.6 TUNEL assay

Sections of the liver were embedded in paraffin wax, and xylene was used to fix the samples in 4% paraformaldehyde fixative, and ethanol was used to dehydrate them. Proteinase K working solution was applied to the tissue followed by a membrane-breaking working

TABLE 1 The specific primers for qRT-PCR.

Genes	Primer sequence (5'-3')	Product length (bp)
<i>β-actin</i>	F: CTGGCTGTCAACATGGTCCC R: TACTGTTGGCTACCACGGCT	86
<i>Nrf2</i>	F: TCTCGCATGACACAGATCCC R: TGCTCTTTGCTCGTCTCTGG	152
<i>Keap1</i>	F: GGGAGAGAAGTGCCTCATCC R: TGGATGAGAAAGTCGCAGCA	90
<i>TOR</i>	F: GGAGCACGTTAAGACTGCCT R: ACAGACAACATGGGCTCCAG	165
<i>S6K1</i>	F: CCGAATGGAAACAAGTGGCG R: GTATGGCGCAGCATTGGATT	97
<i>4EBP1</i>	F: ATCCCACCAGGACCATCCA R: ACGGTCCAACAGAACTTGC	137
<i>JAK1</i>	F: AGCTGTGCAGATATGACCCG R: TATGAGTCCCTCCGGTTCCA	87
<i>JAK2</i>	F: GCGGCACACCAATGTGAATC R: CTCTCACGTTTGCCTCGGA	118
<i>JAK3</i>	F: CTGGAAGCTGGAAGTGGTCA R: TCCTTGTACCTTCACGGGT	120
<i>STAT3</i>	F: GGCCCATGCTGGAAGAGAGA R: GCATTGGCATAACATGGCTGT	90
<i>SOCS1</i>	F: CAATGCCTGGAAGCCTCAAAC R: CTGACTGAGAGCGGAAGGTT	133
<i>SOCS2</i>	F: TAGAGGTGGGCTCTGACCAA R: CTGTCCGTCCTCTATGGTGC	131
<i>SOCS3</i>	F: GTACGGTGACAGGAAGCCAG R: AGTGAAGAAGTGGCGTTGT	98
<i>PIAS1</i>	F: ACAAACACGGTCGCAAACAC R: CCGCCGATACAACCTCTTGA	104
<i>PIAS2</i>	F: GTCCAGAACAACATGCAGCG R: AGGCACAACCTTAGCTGGAC	146
<i>PIAS3</i>	F: TTCGCTGTGAGAAGCCCTC R: TGGCTGAGGGTGTCTTGTTC	201
<i>Bcl-2</i>	F: GGCCGTACAACCTGAGGACAA R: CTGACTCGTCTGTCTTT	190
<i>Bax</i>	F: CCAGCACACGTCAGCACTTA R: AACTGCCCTGGAGAAACCTG	111
<i>Caspase3</i>	F: GCAGGGGAACAGAGCTTGAC R: TGGATGAACCAGGAGCCATT	163

solution. The TUNEL reaction solution was applied according to the Tunel Cell Apoptosis Detection Kit (Servicebio, Wuhan, China) after DAPI re-staining and sealed with a sealer to prevent fluorescence quenching. The images were observed and captured using a Nikon light microscope (Nikon, Eclipse C1 and DS-U3, Japan).

## 2.7 Data analysis

Experiment results are reported as mean ± S.E., and were analyzed using SPSS 24.0 statistical software. One-way ANOVA was used to evaluate the data after ensuring that the data were

homogeneous regarding variances and normality. If one of the non-conforming and was used nonparametric tests. In statistical tests, differences were considered significant when  $P < 0.05$ .

### 3 Results

#### 3.1 LC<sub>50</sub> value of BuP for *P. megacephalus* tadpoles

The cumulative mortality and LC<sub>50</sub> of BuP on the tadpoles are shown in Table 2. After 96 h of exposure, the mortality rate of tadpoles in the control group was 0, the cumulative mortality rate of tadpoles in the low-concentration group (BuP 400 µg/L) was 10%, and the mortality rate of tadpoles in the high-concentration group (BuP 6400 µg/L) reached 80%, with the LC<sub>50</sub> of 6748, 5183, 4182 and 3509 µg/L, respectively.

#### 3.2 Effect of BuP on MDA content and antioxidant enzyme activity in the liver

The levels of hepatic MDA were shown to be substantially elevated after exposure to BuP at 12 h ( $p < 0.05$ ). Notably, the B2 group exhibited the highest MDA content after 48 h of exposure (Figure 1A). The SOD activity was shown to be considerably elevated after exposure to BuP at 12, 24 and 48 h. The maximum level of SOD activity was seen after 24 h of stress in the B2 group (Figure 1B). The GSH-PX activity exhibited a considerable increase in the B2 group after 48 h of stress, demonstrating a notable difference when compared to the control group ( $p < 0.05$ ). However, no significant changes in GSH-PX activity were seen after 12 and 24 h of exposure (Figure 1C).

#### 3.3 Effect of BuP on the mRNA expression of antioxidant genes in the liver

The mRNA expression levels of *Nrf2* exhibited a statistically significant increase ( $p < 0.05$ ) compared to the control group after 12

h exposure to BuP in both the B1 and B2 groups. The B1 group was 1.21-fold and the B2 group was 1.50-fold of the control group. However, after 24 h of exposure, the B1 group was significantly higher 1.32-fold than the control group. Subsequently, after 48 h, no significant differences were observed in the treated groups (Figure 2A). The mRNA expression levels of *Keap1* in the B1 and B2 groups exhibited substantial reductions compared to the control group after 12, 24 and 48 h of exposure (Figure 2B). The levels of *TOR* mRNA expression exhibited a significant elevation in all exposure groups after 12, 24 and 48 h of treatment ( $p < 0.05$ ). Notably, the B2 group had the highest expression level at the 12 h time point of exposure was 4.32-fold of the control group (Figure 2C). The mRNA expression level of *S6K1* exhibited a substantial rise in the B2 group after 24 h exposure was 1.24-fold of the control group. However, in the B1 group, there was a significant increase in *S6K1* mRNA expression after 48 h of exposure, which was 1.30-fold of the control group, but in the B2 group, a reduction in expression was seen (Figure 2D). The mRNA expression level of *4E-BP1* exhibited a significant decrease in the B2 group after 12 h of exposure was 0.37-fold of the control group. However, these levels showed recovery after 24 h of exposure. Subsequently, both the B1 and B2 groups demonstrated a significant increase in *4E-BP1* mRNA expression levels after 48 h of exposure (1.22-, 1.33-fold of the control group, respectively), with a notable distinction when compared to the control group ( $p < 0.05$ ; Figure 2E).

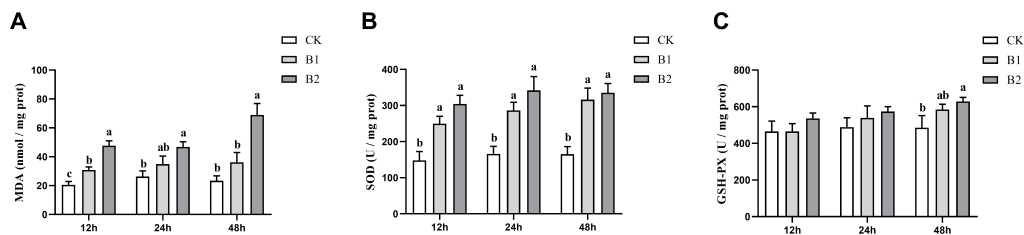
#### 3.4 Effect of BuP on the mRNA expression of JAK-STAT pathway response

The hepatic *JAK1* expression levels in the B1 and B2 treatment groups after 12 h exposure showed no significant difference compared to the control group. However, after 24 h of exposure, the expression level of the B1 group showed a significant increase, whereas the B2 group exhibited a slightly lower expression level than the B1 group. After 48 h of exposure, the hepatic *JAK1* expression levels in the B2 group showed a statistically significant increase (1.33-fold) compared to the control group ( $p < 0.05$ ). The hepatic *JAK2* expression levels exhibited a significant increase in the B1 and B2 groups (1.14-fold, 1.13-fold, respectively) compared to the control group during 48 h of stress ( $p < 0.05$ ). The hepatic *JAK3* expression levels exhibited a statistically significant increase at 24 and 48 h of exposure to BuP in the B1 and B2 groups when compared to the control group ( $p < 0.05$ ). Additionally, the expression levels of *STAT3* demonstrated a significant increase in the B2 group after 24 h and 48 h of exposure to BuP.

The mRNA expression levels of hepatic *SOCS1* and *SOCS2* in the B1 and B2 treatment groups exhibited a significant increase after 12 h, with the B1 group showing the highest expression was 1.33-fold of the control group. Conversely, the mRNA expression levels of hepatic *SOCS1* and *SOCS2* in the B1 and B2 treatment groups were significantly lower than those in the control group after 24 h of exposure ( $p < 0.05$ ). However, these levels rebounded after 48 h, with no significant differences observed in comparison with the control group. The hepatic mRNA expression levels of *SOCS3* and *PIAS1* were significantly elevated after 12 h of exposure in the B1 group

TABLE 2 Deaths and LC<sub>50</sub> value of BuP toxicity to tadpoles of *Polypedates megacephalus* (N = 20).

Concentration (µg/L)	Deaths (N)			
	24 h	48 h	72 h	96 h
0	0	0	0	0
400	0	0	1	2
800	0	1	2	4
1600	1	3	4	6
3200	4	7	9	11
6400	8	12	15	16
LC <sub>50</sub>	6748	5183	4182	3509



**FIGURE 1**  
Effect of BuP on the MDA levels and antioxidant enzyme activity in the liver of *P. megacephalus* tadpoles. (A) MDA content; (B) SOD activity; (C) GSH-PX activity. Different lowercase letters represent significant difference between groups at the same sampling time ( $p < 0.05$ ).

(1.41- and 1.45-fold of the control group, respectively). Additionally, *PIAS1* expression in the B2 group also showed a significant increase. However, after 24 and 48 h, both *SOCS3* and *PIAS1* expression levels in the B1 and B2 groups were significantly reduced. The hepatic mRNA expression levels of *PIAS2* and *PIAS3* were shown to be increased after 12 h of stress in the B1 group, reaching their highest levels (1.45- and 1.25-fold of the control group, respectively), and then reduced after 24 and 48 h of stress (Figure 3).

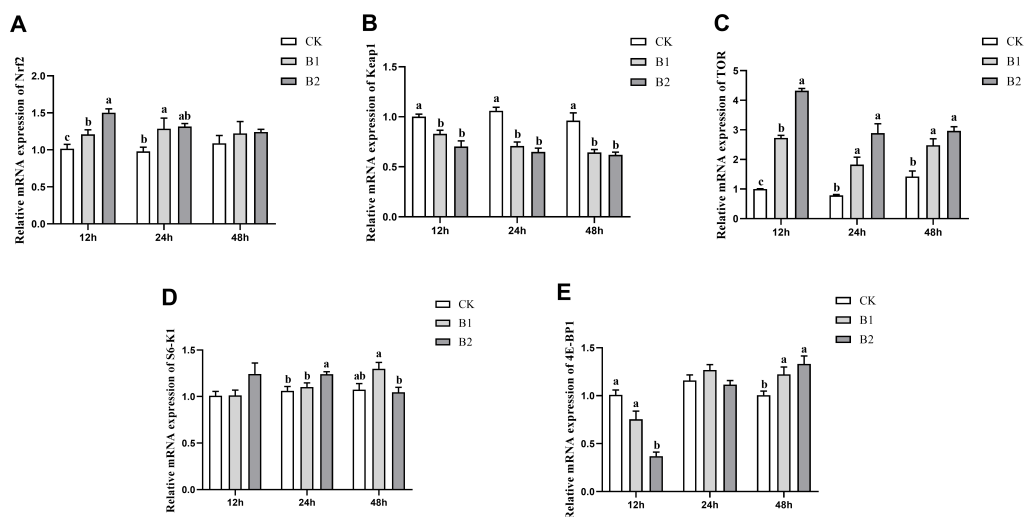
### 3.5 Effect of BuP on cell apoptosis in the liver

The expression levels of the anti-apoptotic gene *Bcl-2* showed a substantial drop after 24 and 48 h of exposure in both the B1 and B2 treatment groups. The B2 group exhibited the lowest expression level (0.72-fold of the control group) after 48 h of stress, as seen in Figure 4A. The levels of expression for pro-apoptotic genes *Bax* and *Caspase3* showed a substantial drop after 12 h of stress in the B1 and B2 treatment groups. Conversely, there was a rise in expression levels after 48 h of stress, as seen in Figures 4B, C.

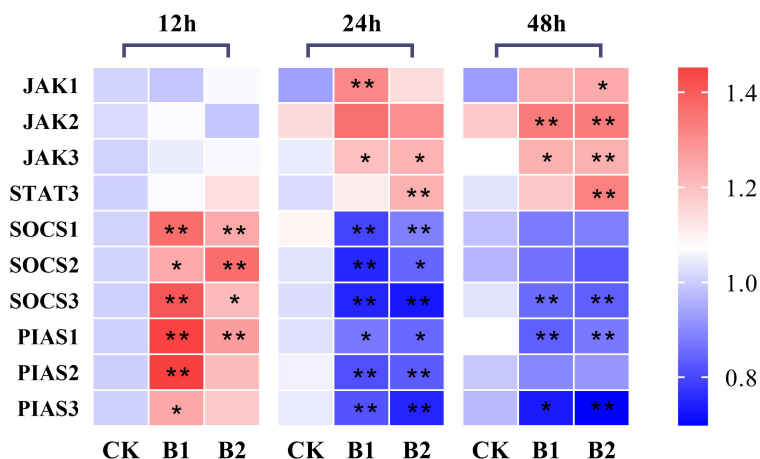
The TUNEL assay indicated a low presence of apoptotic cells in the liver of the control group. Following 24 and 48 h exposure to BuP, there was an observed rise in the number of positive cells. However, it was noted that the number of apoptotic cells in the B2 group at 24 h was somewhat lower compared to the B1 group at 24 h during the BuP exposure period. The B2 group at 48 h has the highest number of positive cells, as seen in Figure 5.

## 4 Discussion

Some studies on the toxicity of BuP using fish as animal models indicated that the 96 h-LC<sub>50</sub> BuP was 2.9 mg/L in the early life stages of medaka (*Oryzias latipes*), 4.2 mg/L in fathead minnow (*Pimephales promelas*), 1.359 mg/L and 2.34 mg/L in embryos and early-life stages of zebrafish (*Danio rerio*), respectively (Yamamoto et al., 2007; Dobbins et al., 2009; Merola et al., 2020; Li et al., 2023a). The present investigation showed that 96 h-LC<sub>50</sub> BuP in *P. megacephalus* tadpoles was 3.5 mg/L, which is lower than those reported in fathead minnow (*Pimephales promelas*), higher than that in the early life stages of medaka and zebrafish. Compared with



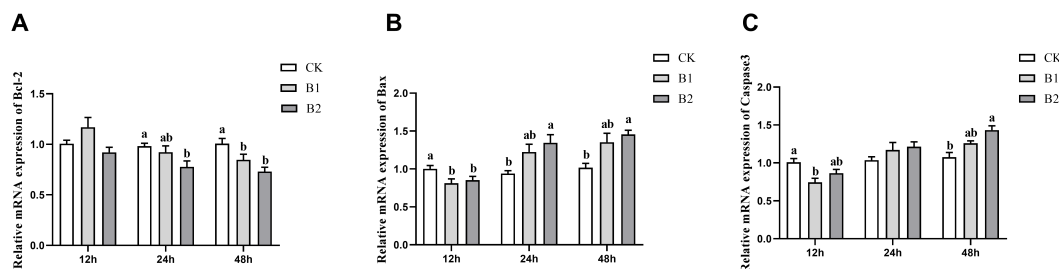
**FIGURE 2**  
Effect of BuP on the mRNA expression of antioxidant genes in the liver of *P. megacephalus* tadpoles. (A) relative mRNA expression of *Nrf2*; (B) relative mRNA expression of *Keap1*; (C) relative mRNA expression of *TOR*; (D) relative mRNA expression of *S6-K1*; (E) relative mRNA expression of *4E-BP1*. Different lowercase letters represent significant difference between groups at the same sampling time ( $p < 0.05$ ).



**FIGURE 3**  
The heatmap of the mRNA expression of JAK-STAT pathway genes under BuP stress at different sampling times in the liver of *P. megacephalus* tadpoles. The symbol of \* means significant difference when  $p < 0.05$ , \*\* highly significant difference when  $p < 0.01$ .

other parabens, MeP was most toxic while BuP was least toxic to *Dugesia jaonica*, for example, 96 h-LC<sub>50</sub> of MeP, EtP, PrP and BuP exposures were found to be 77, 31, 12.3, and 7.3 mg/L, respectively (Li, 2012). However, parabens concentrations in surface water ranged from 3.31 to 55.2 ng/L in the Yellow River (China) and from 15.0 to 164 ng/L in the Huai River (China) (Feng et al., 2019), which was much lower than 96 h-LC<sub>50</sub> BuP to the tadpoles. However, higher BuP concentrations can be detected in the tissues of animals or human beings, such as 70 ng/g in fish (Ramaswamy et al., 2011), and 137 µg/L in women’s urine (Smith et al., 2013), which indicated its bioaccumulation and biomagnifications. As endocrine-disrupting chemicals, parabens would exert severe effects on the body even at sublethal exposure. The study on adult zebrafish (*Danio rerio*) showed the hepatotoxicity of MeP pollutants even at environmentally realistic concentrations (Hu et al., 2022). Exposure to 1 mg/L of BuP had a detrimental impact on the growth of fathead minnow (*Pimephales promelas*) (Dobbins et al., 2009). So, it is necessary to discuss the effect of higher BuP exposure on aquatic animals although the ambient concentration of BuP is not so high.

Numerous studies have shown that the body’s antioxidant enzyme system can adapt to environmental stress by altering the activity of antioxidant enzymes during oxidative stress when aquatic animals are stressed by pollution (Silva et al., 2018; Yang et al., 2018; Schreiber et al., 2019). The promotion of antioxidant systems that consist of enzymatic antioxidants such as superoxide dismutase (SOD), catalase (CAT), and glutathione peroxidases (GSH-PXs) as well as the non-enzymatic antioxidants which collectively reduce oxidative state. Primarily, SOD functions as a catalyst for the dismutation of superoxide anions (O<sub>2</sub><sup>-</sup>) into O<sub>2</sub> and H<sub>2</sub>O<sub>2</sub>, acting as the main protective mechanism against free radicals. CAT and GSH-PX play complementary roles in the removal of H<sub>2</sub>O<sub>2</sub>, and GSH acts as a primary electron donor in the reduction of H<sub>2</sub>O<sub>2</sub> and organic peroxides, thereby scavenging O<sub>2</sub><sup>-</sup> and other reactive oxygen species and safeguarding the organisms from oxidative stress (Hermes-Lima, 2004; Pacoñ and Pacoñ, 2021). BuP has potential cytotoxic and genotoxic impacts in a clinical application and it enhances the production of mitochondrial ROS and triggers oxidative DNA damage. The concentration of BuP in our study was positively correlated with



**FIGURE 4**  
Effect of BuP on the mRNA expression of apoptosis genes in the liver of *P. megacephalus* tadpoles. (A) relative mRNA expression of *Bcl-2*; (B) relative mRNA expression of *Bax*; (C) relative mRNA expression of *Caspase3*. Different lowercase letters represent significant difference between groups at the same sampling time ( $p < 0.05$ ).

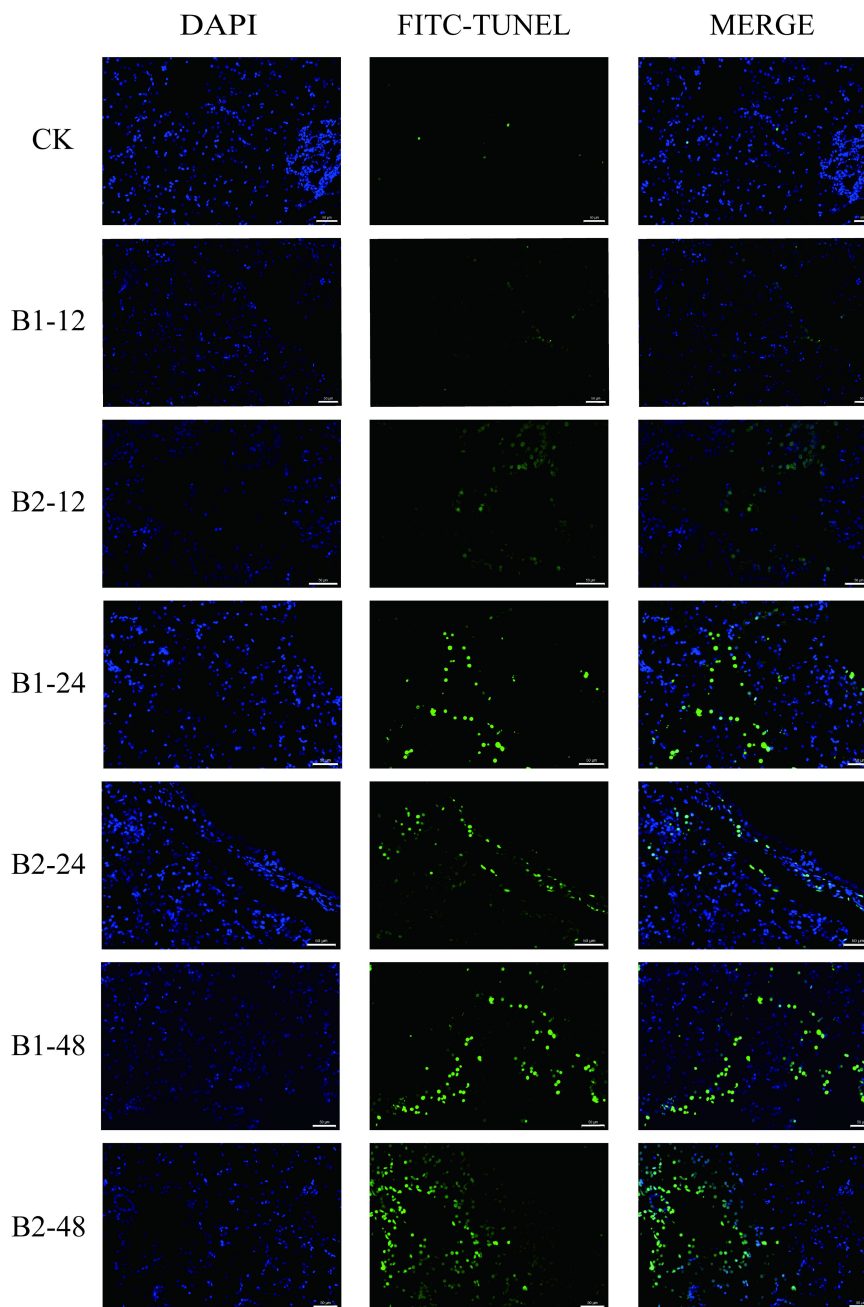


FIGURE 5

Apoptosis of liver cells in *P. megacephalus* tadpoles. CK, control group; B1-12, B1-24 and B1-48: B1 group sampling at 12, 24 and 48h; B2-12, B2-24 and B2-48: B2 group sampling at 12, 24 and 48h. Note: The first column showed the nucleus (blue); the second column showed the positive cell of apoptosis (green); the third column showed the merge; bar = 50  $\mu$ m.

the activities of SOD and GSH-PX, indicating activation of the antioxidant defense mechanism. The activities of SOD exhibited a statistically significant increase compared to the control group after 12, 24 and 48 h exposure to BuP in both the B1 and B2 groups ( $p < 0.05$ ). The activities of GSH-PX were significantly increased in the B2 group than the control group after 48 h of stress ( $p < 0.05$ ). This increased activity of antioxidant enzymes in tadpoles exposed to BuP demonstrated that there was an increase in ROS in the liver, and the tadpoles showed signs of lipid peroxidation in the liver, indicating that antioxidant defenses and neutralized ROS were

weakened in prolonged and highly concentrated environments. We infer that antioxidant defenses are upregulated in the initial phases of exposure as a result of oxidative stress, including superoxide dismutase (SOD) and glutathione peroxidase (GSH-PX). As a result of the BuP exposure, cells attempt to neutralize reactive oxygen species (ROS). Nevertheless, prolonged exposure can overwhelm these antioxidant systems. Due to persistent ROS generation, key antioxidants like GSH-PX may be depleted and antioxidant enzymes may become less active, which leads to a weakening of defenses. A compromised antioxidant defense system

can result in oxidative damage over time. A similar phenomenon was observed in Nile tilapia (*Oreochromis niloticus*), with liver SOD and GSH-PX activities increased after 6 d and 12 d BuP exposure respectively. Furthermore, having strong antioxidant defenses at all times might quickly prevent, or prior research has shown that when ROS are not promptly removed (Pacoñ et al., 2021), they may induce lipid peroxidation and generate a substantial quantity of peroxides, such as MDA. In our study, the MDA contents were significantly increased in the B2 group compared to the control group.

The Keap1-Nrf2 system plays a vital role in the transcriptional activation of an array of antioxidant and detoxification genes (Wang et al., 2018). In response to oxidative stress, Keap1 is a sensor for the stress signals that lose their activity to interact with transcription factor Nrf2. Subsequently, Nrf2 translocates into the nucleus, which initiates the activation of genes encoding antioxidant enzymes (Taguchi et al., 2011). The mRNA levels of Nrf2 in the large yellow croaker (*Pseudosciaena crocea*) were increased sharply when exposed to 8 mg/L Zn and 32, 64 µg/L Hg (Zheng et al., 2016). The activity of Nrf2 is primarily controlled by Keap1 (Itoh et al., 2010). In this study, the relative expression of *Keap1* was inversely linked with the concentration of BuP in all groups, but the mRNA expression level of *Nrf2* rose and peaked in the B2 group. The mRNA expression level of *Nrf2* was significantly increased in the three different concentrations of BuP groups after 12 h of exposure, while only the B1 group was significantly increased than the control group after 24 h of exposure ( $p < 0.05$ ). The relative expression of *Keap1* in the B1 and B2 groups exhibited a significant decrease compared to the control group after 12, 24 and 48 h of exposure ( $p < 0.05$ ). Therefore, we speculated that increasing *Nrf2* mRNA expression related to a decrease of *Keap1* could elevate the antioxidant ability to BuP stress. On the other hand, by activating S6K1 and suppressing 4E-BP1, the TOR signaling pathway may promote the production of anti-inflammatory cytokines (Weichhart et al., 2011). In a sensitive manner, cytokines, growth factors, amino acids, or insulin activate mTOR and dramatically increase the phosphorylation status of 4E-BP1 and S6K1 (Hay and Sonenberg, 2004). A loss of mTOR function causes a cell cycle arrest as well as a severe reduction in protein synthesis. Our findings demonstrated that exposure to BuP significantly increased TOR transcription in the liver of *P. megacephalus* tadpoles. After exposure for 12 h, the transcriptional levels of *4E-BP1* dramatically dropped in the B2 group, and after 48 h, it significantly increased in both the B1 and B2 groups. A substantial increase in *S6K1* transcriptional levels was seen in the B1 group, whereas a reduction in activity was observed in the B2 group. Gonzalez-Rodriguez et al. examined the impact of S6K1 on the vulnerability of hepatocytes to apoptosis triggered by death receptor activation within a hepatic cellular model (González-Rodríguez et al., 2009). The results of our study are consistent with the findings of Pardo et al. which indicate that the expression of *S6K1* in hepatocytes altered following 16 h treatment with palmitic acid, as compared to the control group (Pardo et al., 2015).

The present investigation revealed that exposure to BuP produced oxidative stress resulting in an up-regulation of *JAK1*,

*JAK2*, *JAK3*, and *STAT3*, hence initiating the JAK-STAT3 signaling pathway. Our findings are consistent with Li et al. (2023b) findings that, compared to the controls, grass carp liver cells subjected to TBBA and Pb showed substantial oxidative stress as well as increased expression levels of the hepatic JAK2-STAT3 signaling pathways. On the other hand, preventing oxidative stress from activating the JAK-STAT3 signaling pathway may be a more efficient strategy to shield the body's tissues and cells from harmful external stimuli. SOCS, PIAS, and protein tyrosine phosphatases (PTPs) are regulators that adversely regulate the JAK-STAT signaling pathway in addition to its constituent parts (Wormald and Hilton, 2004). The initial elevation and subsequent downregulation of the genes *SOCS1*, *SOCS2*, *SOCS3*, *PIAS1*, *PIAS2*, and *PIAS3* were seen in this research after 24 h of exposure to BuP. It is suggested that these findings, namely the initially elevated SOCS genes and PIAS proteins, may represent a kind of adaptive cellular reaction to the JAK-STAT signaling system, which is necessary to control or modify the BuP-induced JAK-STAT pathway activation. For example, Adeyi et al. showed that rats exposed to 2-methoxyethanol first showed an elevation of SOCS genes and PIAS proteins, and then later on, a downregulation was seen (Adeyi et al., 2023).

Oxidative stress may trigger programmed cell death, such as apoptosis, a protective process that controls the number of cells in the body and eliminates undesirable and perhaps hazardous ones (Estaquier et al., 2012). Pro- and anti-apoptotic Bcl protein family members and Caspase inhibitor proteins regulate apoptosis (Logan and Storey, 2016). Bax is a pro-apoptotic protein that translocates from the cytoplasm to the mitochondria upon cellular injury or stress, promoting an increase in permeability of the outer mitochondrial membrane, leading to the release of cytochrome c, which in turn activates the downstream apoptotic cascade response. Like Bax, the B-cell lymphocytic leukemia proto-oncogene (*Bcl-2*) family is an anti-death protein that prevents cell apoptosis (Yin et al., 2023). *Bcl-2* family proteins move from the cytoplasm to the mitochondrial membrane when the cells receive an apoptotic signal. Caspases may readily identify stress-induced apoptosis. The intrinsic pathway is initiated by the release of cytochrome c from the mitochondria, which then activates Caspase3 and Caspase9. Numerous proteins, such as fodrin and nuclear laminae, are cleaved by Caspase3 and Caspase9, which ultimately results in cell death (Liang et al., 2016). The results of this experiment show that increasing the concentration of BuP can cause a gradual decrease in the mRNA expression level of *Bcl-2* in the liver, while *Bax* and *Caspase3* levels first decrease and then increase, suggesting that BuP can cause apoptosis in *P. megacephalus* tadpoles' hepatocytes through the accumulation of toxicity over time. The mRNA expression level of *Bcl-2* was significantly decreased in the B2 group after 24 and 48 h of exposure to BuP ( $p < 0.05$ ). The *Bax* and *Caspase3* levels were significantly decreased in the B1 group after 12 h of stress and then significant increased in the B2 group after 48 h of stress ( $p < 0.05$ ). The TUNEL assay revealed a significant increase in the quantity of apoptotic cells inside the liver. The findings of this study suggest that exposure to BuP stress may potentially trigger apoptosis in *P. megacephalus* tadpoles.



## 5 Conclusions

Following exposure to BuP, the amount of MDA in the liver increased as exposure duration and concentration increased. The antioxidant defense response to BuP stress included the activation of the antioxidant enzymes (SOD and GSH-PX), and the Nrf2-Keap1 signaling pathway contributes to tadpoles' antioxidant defense response. The JAK-STAT pathway inhibited toxic stress after 24 h, but with prolonged exposure, BuP stress induced apoptosis in tadpole hepatic cells.

## Data availability statement

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding authors.

## Ethics statement

All animal protocols described herein were performed according to the guiding principles of the Hainan Provincial Ecological Environment Education Center Experimental Animal Ethics Committee (No: HNECEE-2020-003).

## Author contributions

XA: Data curation, Software, Writing – original draft, Writing – review & editing. MI: Formal analysis, Writing – review & editing.

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## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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