



Effect of acute ammonia exposure on the glutathione redox system in FFRC strain common carp (*Cyprinus carpio* L.)

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Abstract

Ammonia is one of the most common aquatic pollutants. To analyze the effect of ammonia exposure on the glutathione redox system, we investigated the levels of hydrogen peroxide (H₂O₂) and glutathione, and transcription and activities of glutathione-related enzymes in liver and gills of FFRC strain common carp (*Cyprinus carpio* L.) exposed to 0, 10, 20, and 30 mg/L of ammonia. The results showed that H₂O₂ content reached a maximum level at 48 h of exposure in the liver of fish. In gills, H₂O₂ increased rapidly at 6 h and reached to maximum levels at 24 h of exposure, indicating that gills experienced oxidative stress earlier than the liver of fish exposed to ammonia. Reduced glutathione (GSH) content and reduced glutathione/oxidized glutathione (GSH/GSSG) ratio increased significantly within 24 h of exposure. Meanwhile, the transcription and activities of glutathione S-transferase (GST) and glutathione reductase (GR) increased significantly in the liver, and glutathione peroxidase (GSH-Px) and GST increased in the gills of fish exposed to ammonia. Malondialdehyde (MDA) content kept at a low level after exposure to low concentration of ammonia, but increased significantly after exposure to 30 mg/L ammonia for 48 h along with a decrease in GSH content and GSH/GSSG ratio. These data showed that the glutathione redox system played an important role in protection against ammonia-induced oxidative stress in the liver and gills of FFRC strain common carp, though the defense capacity was not able to completely prevent oxidative damage occurring after exposure to higher concentration of ammonia. This research systematically studied the response of the glutathione redox system to ammonia stress and would provide novel information for a better understanding of the adaptive mechanisms of fish to environmental stress.

Keywords Ammonia · FFRC strain common carp · Glutathione redox system · Hydrogen peroxide (H₂O₂) · Oxidative stress

Introduction

In aquaculture systems, ammonia is one of the major environmental pollutants of concern. It has been reported that the concentration of ammonia increased directly with culture period, and might reach as high as 46 mg/L in intensive aquaculture systems (Cheng et al. 2015). High concentration of ammonia can cause growth reduction, histological damage, immune suppression, and high mortality (Qi et al. 2017; Liang et al. 2018). Exogenous ammonia enters into fish mainly through gills, and is then transported through the blood to

the liver and other organs. One of the known mechanisms of ammonia toxicity is the induction of oxidative stress, through increasing the concentration of reactive oxygen species (ROS) such as hydrogen peroxide (H₂O₂), hydroxyl radical (HO·), and superoxide radical (O₂^{·-}) (Murthy et al. 2001; Rama and Manjabhat 2014). Overproduction of ROS in cells can result in the oxidation of proteins, DNA, and lipids, eventually leading to cell death (Martinez-Alvarez et al. 2005; Hegazi et al. 2010). To cope with oxidative stress and keep cellular redox state in balance, aquatic organisms evolved both non-enzymatic and enzymatic antioxidant defense system to convert ROS to harmless metabolites (Sinha et al. 2015; Ramirez-Duarte et al. 2016). It has been reported that ammonia exposure can alter activities of antioxidant enzymes to prevent oxidative damage in the gills, muscle, liver, and brain of fish (Qi et al. 2017; Hegazi et al. 2010; Lorenzo et al. 2017). The induction of antioxidant defense is thought to be a protective reaction against ammonia stress in fish, but the exact defense mechanism is not clear.

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The glutathione redox system, including glutathione, glutathione peroxidase (GSH-Px), glutathione S-transferases (GST), and glutathione reductase (GR), plays important roles in maintaining cellular redox homeostasis and protecting cells from oxidative damage during environmental stress (Lesser 2011; Lin et al. 2018). Glutathione, including reduced glutathione (GSH) and oxidized glutathione (GSSG), is an effective non-enzymatic antioxidant that can modulate the redox status of protein thiols, and directly scavenge singlet oxygen and hydroxyl radicals (Li et al. 2003; Al-Ghais 2013; Xia and Wu 2018). GSH-Px detoxifies the cytosolic H_2O_2 and hydroperoxides using GSH as a cofactor (Arthur 2001; Srikanth et al. 2013). GST functions to detoxify xenobiotics or lipid peroxidation end products by conjugation GSH (Comakli et al. 2011). GR is a NADPH-dependent oxidoreductase which can catalyze the conversion of GSSG to GSH (Lin et al. 2018). The dynamic balance between GSH and GSSG is important for ROS detoxification and cellular survival, and GSH/GSSG ratio has been widely used as an indicator of cellular redox status (Sies 1999). Previous studies have reported that glutathione and other glutathione-related enzymes were involved in the detoxification of harmful ROS in ammonia-exposed fish (Sinha et al. 2014). However, there are no detailed studies on the molecular patterns of these antioxidative enzymes in fish under ammonia stress.

FFRC strain common carp, a new strain of *Cyprinus carpio* with the advantages of fast growth, good body shape, strong adaptability, and stable genetic traits, has been cultured widely in many provinces of China (Dong 2011). However, the stress response of FFRC strain common carp to environmental pollutants has not been studied. In this study, a comprehensive analysis of the glutathione redox system, including glutathione, GSH-Px, GST, and GR, was conducted in the gills and liver to understand the antioxidant response of FFRC strain common carp to ammonia pollution. These results will offer a better insight into the mechanism of fish adaption to ammonia.

Materials and methods

Animals

FFRC strain common carps (50 ± 0.16 g body mass, 1 month old) were obtained from a fish farm in Qinxian (Shanxi, China), and acclimated for 2 weeks in cycling-filtered plastic tanks containing continuously circulating dechlorinated tap water at 25 °C (pH 7.5; dissolved oxygen 6.0–7.0 mg/L) with natural light and photoperiod. The fish were fed twice a day with commercial pellets, and were fasted for 24 h prior to experimentation.

Experimental design and sample collection

Ammonium chloride (NH_4Cl) was used as a source of the total ammonia-nitrogen (T-AN). Based on the value of 96 h median lethal concentration (96 h LC50, T-AN: 35.6 mg/L) established by our group, test concentrations of ammonia were 0 (control), 10 (low), 20 (middle), and 30 (high) mg/L. The exposure was conducted for 48 h with three replicates of four treatments (total of twelve tanks). Twenty fish were selected randomly and placed in a 150L plastic tank containing 100 L of test solution. Ammonia concentration was measured by nesslerization (Hegazi et al. 2010) every 6 h and adjusted by adding the calculated amount of the NH_4Cl solution. Fish were not fed during the whole experimental period. Each tank was cleaned to remove feces and approximately 50% of the water was changed daily with dechlorinated tap water containing the respective amount of ammonia.

After exposure to ammonia for 6, 24, and 48 h, three fish from each tank were randomly sampled and dissected after anesthesia with tricaine methanesulfonate MS-222 (0.2%). The gills and liver were quickly excised within 2 min on ice, and frozen in liquid nitrogen for further analysis.

In this study, all fish were administered in strict accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals (China).

Quantification of oxidative stress markers

The frozen samples were homogenized with ice-cold normal saline at a ratio of 10% (w:v). The homogenates were centrifuged at $12,000 \times g$ for 15 min at 4 °C, and the supernatants were used for determination of biochemical parameters.

H_2O_2 content was assessed using a commercial assay kit (Nanjing Jiancheng Bioengineering Institute, Nanjing, China). H_2O_2 binds with molybdenic acid to form a complex, which was measured at 405 nm, and the content of H_2O_2 was then calculated.

MDA content was determined using a commercial assay kit (Nanjing Jiancheng Bioengineering Institute, Nanjing, China) based on the thiobarbituric acid (TBA) reactivity. Briefly, the supernatants were collected and reacted with an equal volume of 0.67% (w/v) thiobarbituric acid (TBA) in a boiling water bath for 30 min. After cooling, the mixture was centrifuged at $3000 \times g$ for 10 min. The absorbance of supernatant was measured at 532 nm and corrected for non-specific turbidity by subtracting the absorbance at 600 nm and 450 nm.

Determination of glutathione

Glutathione content was assessed using a commercial assay kit (Nanjing Jiancheng Bioengineering Institute, Nanjing, China). For total glutathione (T-GSH), the supernatant was added to a reaction mixture containing

100 mM sodium phosphate buffer (pH 7.5), 0.6 mM 5,5-dithiobis-2-nitrobenzoic acid (DTNB), 0.25 mM NADPH, and 4 mM EDTA. The reaction was initiated by the addition of glutathione reductase and the absorbance was monitored at 412 nm. For oxidized glutathione (GSSG), the supernatant was incubated with 170 mM 2-vinylpyridine for 1 h at 30 °C to derivatize any reduced glutathione (GSH) present in the sample. GSSG in this portion was quantified using the same method. Reduced glutathione (GSH) content was calculated as the difference between the content of T-GSH and GSSH.

Enzyme assays

The activities of glutathione S-transferase (GST), glutathione peroxidase (GPX), and glutathione reductase (GR) were determined with commercial assay kits (Nanjing Jiancheng Bioengineering Institute, Nanjing, China) according to the manufacturer’s instructions, respectively.

GR activity was measured by the method of Murshed et al. (2008). The reaction was started by adding the supernatant to an assay containing mM sodium phosphate buffer (pH 7.4), 1 mM EDTA, 0.2 mM NADPH, and 5 mM GSSG. GR activity was expressed as units per milligram of protein.

GST activity was measured by measuring the conjugation of GSH with 1-chloro-2,4-dinitrobenzene (CDNB) (Habig et al. 1974). The reaction was started by adding the supernatant to an assay containing 50 mM sodium phosphate buffer (pH 6.5), 1 mM EDTA, 5 mM GSH, and 2 mM CDNB. GST activity was expressed as units per milligram of protein.

GPX activity was measured by the method of Ahmad and Pardini (1988). The reaction was initiated by adding 0.5 mM hydrogen peroxide (H₂O₂) to the assay mixture of supernatant, 50 mM sodium phosphate buffer (pH 7.0), 1 mM EDTA, 2 mM NaN₃, 0.2 mM NADPH, glutathione reductase, and 5 mM GSH. GPX activity was expressed as units per milligram of protein.

Protein amounts were determined by the method of Bradford (1976) with bovine serum albumin (BSA) as the standard.

Total RNA extraction and real-time PCR

Total RNA was isolated from fish tissues using a Trizol kit (TaKaRa, Dalian, China) according to the manufacturer’s instructions and then dissolved in DEPC-treated water. The expression of glutathione S-transferase (*GSTP1*), glutathione peroxidase (*GPXI*), and glutathione reductase (*GR*) was analyzed by quantitative real-time PCR (qPCR). Briefly, 1 µg of total RNA was reverse transcribed into cDNA using the PrimeScript™ RT Master Mix (TaKaRa, Dalian, China), and qPCR was carried out using the SYBR Premix Ex Taq™ II (TaKaRa, Dalian, China), with *β-actin* as the internal control. The primers used for qPCR were listed in Table 1. All the qPCR analyses were performed on an Applied Biosystems 7500 Real-Time PCR System. The cycling procedure for qPCR was 94 °C for 1 min, followed by 40 cycles of 94 °C for 30 s, 60 °C for 30 s, and 72 °C for 1 min.

Three independent biological replicates and three technical replicates of each biological replicate for each sample were analyzed by qPCR. The differences in expression levels were calculated using the 2^{-ΔΔCT} method (Schmittgen and Livak 2008).

Statistical analysis

All data were expressed as mean ± SE (standard error). Statistical analyses were performed using one-way ANOVA followed by LSD test. The significance level was set at *P* < 0.05.

Results

Ammonia-induced H₂O₂ generation and lipid peroxidation

Exposure to ammonia induced a production of H₂O₂ in both the liver and gills of FFRC strain common carp (Fig. 1). In the liver, H₂O₂ content showed no significant change after exposure to 10 mg/L ammonia, but increased gradually within 48 h of exposure to 20 and 30 mg/L ammonia (Fig. 1A). In the

Table 1 Primer sequences used in qPCR

Gene name	Sequence of primers (5’-3’)	Accession no.
<i>β-actin</i>	Forward: GCAAGAGAGGTATCCTGACC Reverse: CCCTCGTAGATGGGCACAGT	XM_019103102.1
<i>GPXI</i>	Forward: AGGAGAATGCCAAGAATG Reverse: GGGAGACAAGCACAAGG	GQ376155.1
<i>GR</i>	Forward: CCACCCGTCCACTTT Reverse: CACCTCTACCGACCATAG	JN126053.1
<i>GSTP1</i>	Forward: ACTACAACCTGTTCGACCTT Reverse: CCTATTCTAACGACGGG	DQ497597.1

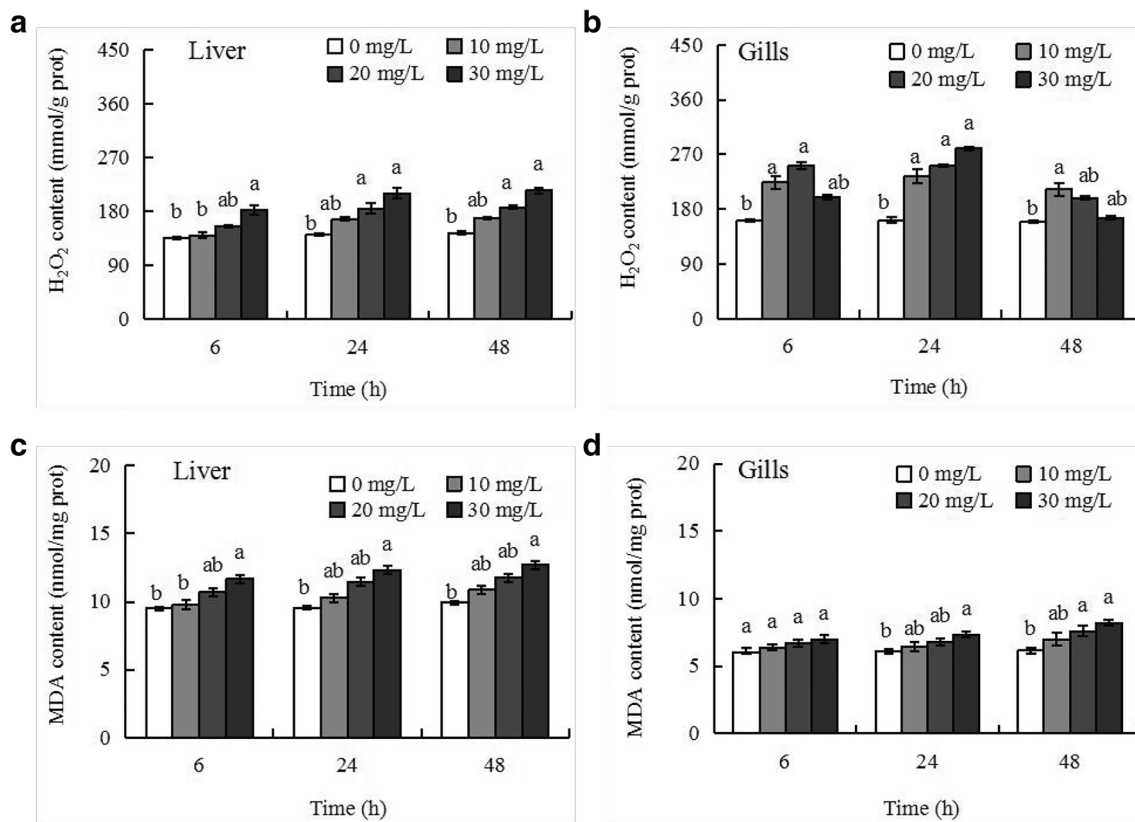


Fig. 1 The content of H₂O₂ (A and B) and MDA (C and D) in the liver and gills of FFRC strain common carp during ammonia exposure. Values are mean \pm SE. Different letters indicate significant difference ($P < 0.05$) among treatments

gills, H₂O₂ content increased rapidly at 6 h of ammonia exposure and reached a maximum at 24 h in all treatment groups of fish ($P < 0.05$), and then recovered to the control level at 48 h in the 20 and 30 mg/L groups (Fig. 1B).

MDA content significantly increased with increasing ammonia concentration and exposure duration in both the liver and gills of FFRC strain common carp (Fig. 1). In the liver, MDA content showed no significant change within 48 h of exposure to 10 and 20 mg/L ammonia, but increased significantly after exposure to 30 mg/L ammonia as a function of time ($P < 0.05$) (Fig. 1C). In the gills, MDA content did not

have significant change within 48 h of exposure to 10 mg/L ammonia, but increased significantly after exposure to 20 and 30 mg/L ammonia for 48 h ($P < 0.05$) (Fig. 1D).

Ammonia-induced changes in GSH content and GSH to GSSG ratio

GSH content increased significantly within 24 h of ammonia exposure, following by a reduction at 48 h of exposure in both the liver and gills of FFRC strain common carp (Fig. 2). In the liver, GSH content increased rapidly at 6 h ($P < 0.05$) and

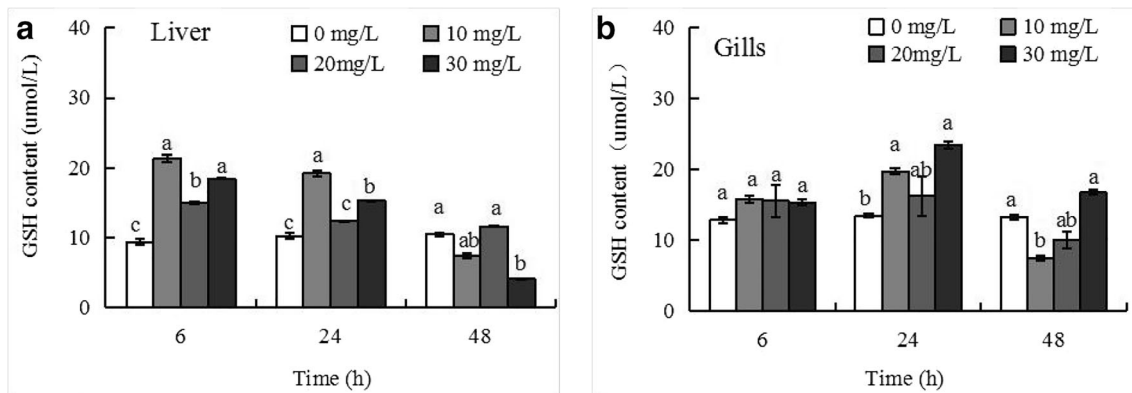


Fig. 2 GSH content in the liver (A) and gills (B) of FFRC strain common carp during ammonia exposure. Values are mean \pm SE. Different letters indicate significant difference ($P < 0.05$) among treatments

remained at a high level after 24 h of ammonia exposure ($P < 0.05$). After exposure to ammonia for 48 h, GSH content recovered to the control level in the 10 and 20 mg/L ammonia treatment groups, but decreased significantly in the 30 mg/L treatment group ($P < 0.05$) (Fig. 2A). In the gills, GSH content did not have significant change during 48 h of exposure to 20 mg/L ammonia. However, a significant increase in GSH content occurred after exposure to 10 and 30 mg/L ammonia for 24 h, respectively ($P < 0.05$). After exposure to ammonia for 48 h, GSH content recovered to the control level in the 30 mg/L treatment group, but decreased significantly in the 10 mg/L group ($P < 0.05$) (Fig. 2B).

GSH/GSSG ratio displayed a comparable pattern as seen for GSH content in both the liver and gills of FFRC strain common carps (Table 2). In the liver, GSH/GSSG ratios increased significantly at 6 and 24 h of ammonia exposure ($P < 0.05$). However, a significant increase occurred after 24 h of ammonia exposure in the gills. After exposure to ammonia for 48 h, GSH/GSSG ratio recovered to the control level in both the liver and gills.

Ammonia-induced responses of glutathione-related enzymes

The glutathione-related enzymes in the glutathione redox system responded differently to ammonia exposure in the liver and gills of FFRC strain common carp (Fig. 3). In the liver, the activities of GST and GR showed a gradual increasing trend after exposure to 30 mg/L ammonia, but no significant change was found during exposure to 10 or 20 mg/L ammonia (Fig. 3A and E). The GSH-Px activity had no significant change in any of the treatment groups (Fig. 3C).

In the gills, GST activity increased significantly within 24 h of ammonia exposure, and then recovered to the control level at 48 h (Fig. 3B). The GSH-Px activity increased gradually and reached maximum values at 48 h of ammonia exposure (Fig. 3D). However, there were no significant changes in GR

activity during the 48 h of ammonia exposure in any of the treatment groups (Fig. 3F).

Ammonia-induced changes in the transcription levels of glutathione-related enzymes

The transcriptional levels of genes encoding for the GSH-related enzymes (*GSTP1*, *GPX1*, and *GR*) were analyzed at 24 h of ammonia exposure, which seemed to be the most responsive time point for most of the doses of ammonia (Fig. 4). In the liver, the expression of *GSTP1* was significantly upregulated in the 30 mg/L treatment group, and the expression of *GR* was upregulated in a dose-dependent manner, but the expression of *GPX1* was not changed during ammonia exposure. In the gills, the expression of *GSTP1* was significantly higher as compared with the control in all treatment groups, and the expression of *GPX1* was increased in a dose-dependent manner, while the expression of *GR* did not show a significant difference between the control and all the treatment groups.

Discussion

Oxidative stress is one of the toxicity mechanisms of environmental pollutants on aquatic organisms (Benli et al. 2008; Liew et al. 2013). In this study, ammonia induced significant increases in H_2O_2 content in the liver and gills of FFRC strain common carp and caused oxidative stress, which was manifested by an increase in MDA content. However, there were differences in cellular response to ammonia stress between the liver and gills of fish. In the gills, H_2O_2 content increased rapidly at 6 h, and the content of GSH, the levels of GST, and GSH-Px increased significantly during ammonia exposure. In the liver, H_2O_2 content enhanced after exposure to ammonia for 24 h, and GSH content increased significantly at 6 h and maintained at a high level at 24 h. The rapid increase in GSH level might be due to the fact that this peptide was synthesized in the liver and then transported to other organs of fish (Kaplowitz et al. 1985). Meanwhile, the levels of GST and GR increased gradually after exposure to high concentration of ammonia (30 mg/L). Based on the accumulation of H_2O_2 content and change in the glutathione redox system, it can be deduced that the gills experienced oxidative stress earlier than the liver of fish during ammonia exposure. This result was consistent with the previous report that exposure to high concentration ammonia induced oxidative stress in the gills earlier than that in the brain of the mudskipper (Ching et al. 2009). This might be due to that the gills, but not other organs, were exposed to environmental ammonia instantly under ammonia stress. Moreover, fish gills serve as the dynamic respiratory and osmoregulatory organ, and are likely to possess a high capacity to produce ROS (Sinha et al. 2015). Therefore, it is

Table 2 GSH/GSSG ratio in the liver and gills of FFRC strain common carp during ammonia exposure

		0 mg/L	10 mg/L	20 mg/L	30 mg/L
Liver	6 h	3.25 ± 0.15 ^c	11.51 ± 0.42 ^a	6.00 ± 0.25 ^b	10.39 ± 0.41 ^a
	24 h	4.10 ± 0.21 ^b	10.11 ± 0.35 ^a	3.26 ± 0.08 ^b	9.56 ± 0.32 ^a
	48 h	4.67 ± 0.23 ^a	3.64 ± 0.11 ^{ab}	4.80 ± 0.43 ^a	2.39 ± 0.06 ^b
Gills	6 h	5.47 ± 0.16 ^a	6.74 ± 0.31 ^a	7.06 ± 0.34 ^a	6.97 ± 0.43 ^a
	24 h	5.74 ± 0.21 ^b	8.98 ± 0.27 ^a	5.21 ± 0.25 ^b	11.33 ± 0.39 ^a
	48 h	5.63 ± 0.32 ^a	3.00 ± 0.12 ^b	4.19 ± 0.27 ^{ab}	5.44 ± 0.41 ^a

Different letters indicate significant difference ($P < 0.05$) among treatments

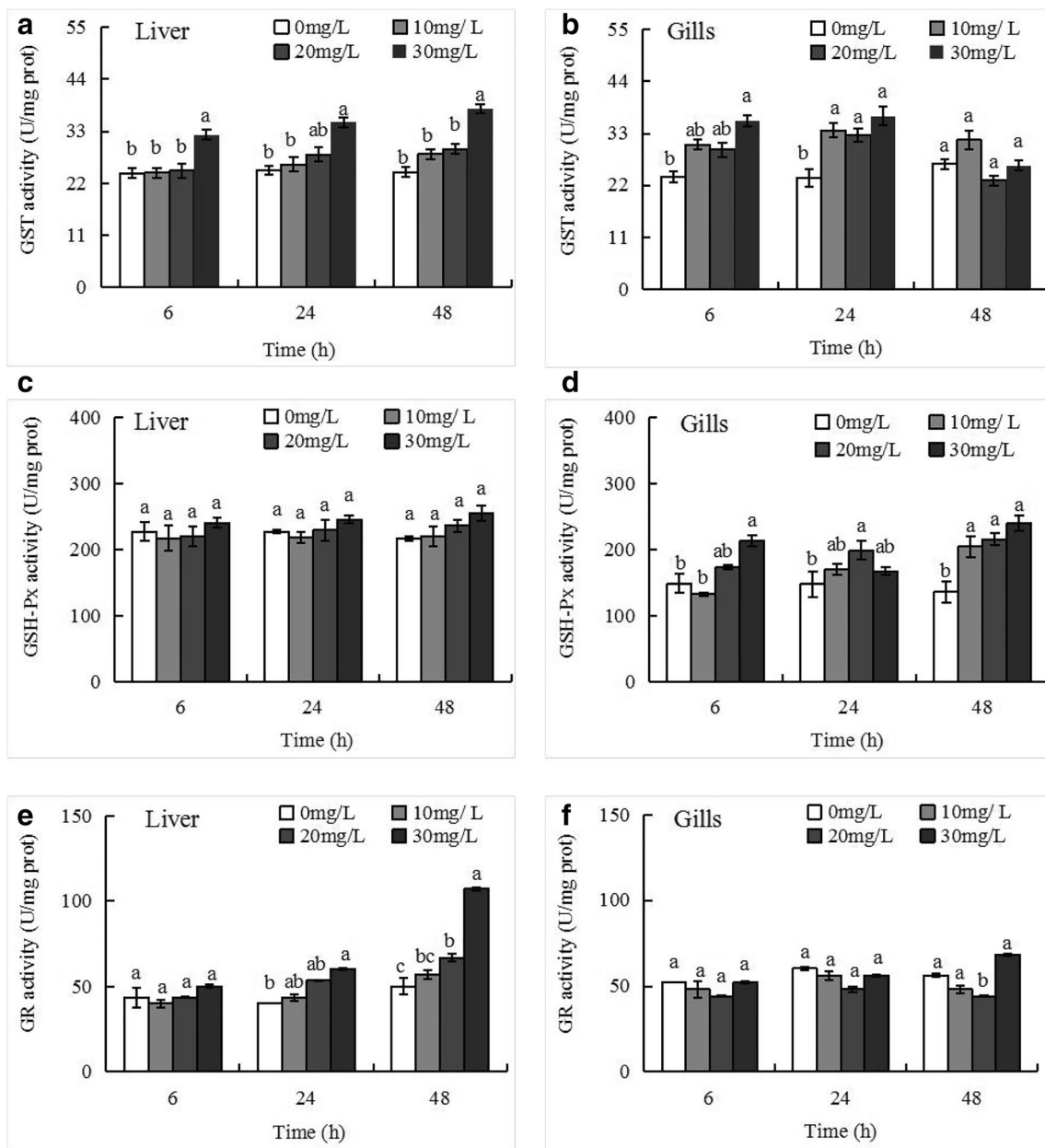


Fig. 3 Activities of enzymes involved in the glutathione redox system during ammonia exposure. Values are mean \pm SE. Different letters indicate significant difference ($P < 0.05$) among treatments

necessary to investigate the injury of gills in fish under various environmental stresses.

Ammonia exposure induces oxidative stress, and further leads to alterations in antioxidant defense system in fish (Sinha et al. 2014; Cong et al. 2018). The glutathione redox system, including GSH and glutathione-related enzymes, has been reported to provide protection against ROS in fish exposed to varied environmental pollutants (Anjum et al. 2014; Radovanovic et al. 2015). Hegazi et al. (2010) have found that glutathione-related enzymes played important roles in preventing ammonia-induced oxidative stress in the liver and white muscle of Nile tilapia juveniles. In the present study,

the glutathione redox system was found to be involved in response to ammonia stress in both the liver and gills of FFRC strain common carp. The GSH level and GSH/GSSG ratio increased rapidly after exposure to ammonia. Meanwhile, the expression levels and activities of GSH-Px, GR, and GST enhanced significantly, suggesting a transcriptional activation of genes for antioxidant enzymes in response to ammonia stress. This result was different from a previous study in which the glutathione redox system was nearly unaffected in carp exposed to environmental ammonia (Sinha et al. 2014). A possible explanation for this discrepancy is that fish used in the present study is a new strain of common carp with

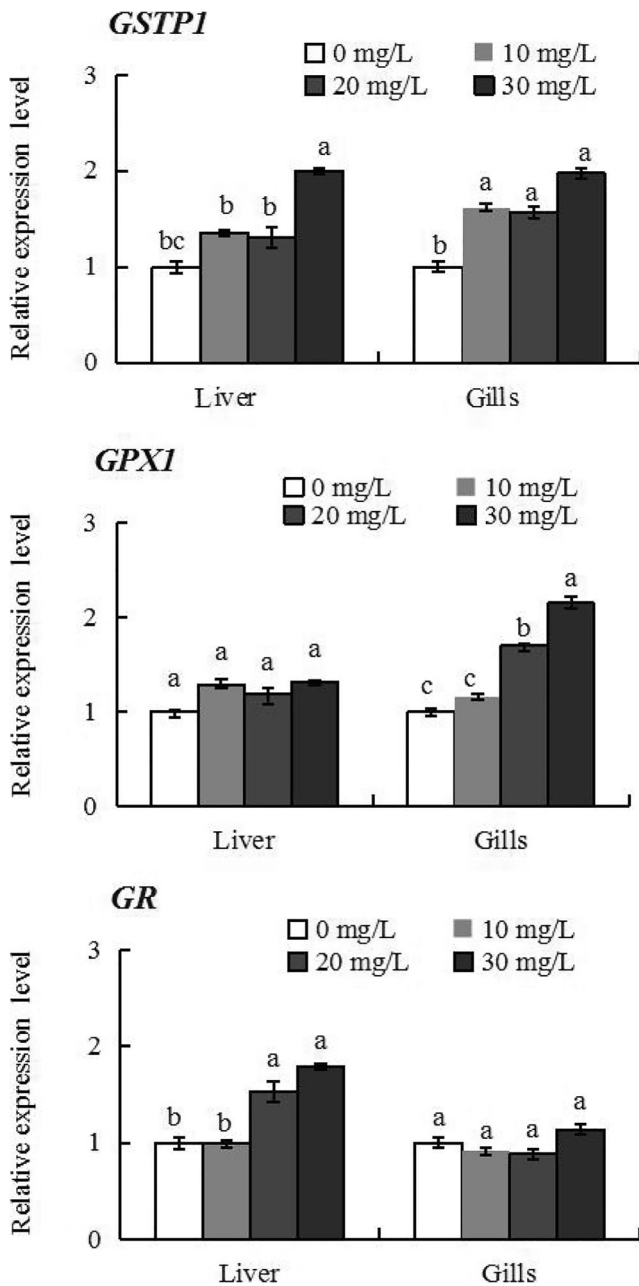


Fig. 4 Relative gene expression levels of *GSTP1*, *GPX1*, and *GR* in the liver and gills of FFRC strain common carp after exposure to ammonia for 24 h. Values are mean ± SE. Different letters indicate significant difference ($P < 0.05$) among treatments

a well-developed defense system. Present finding suggested that FFRC strain common carp might have an effective antioxidant system to deal with ammonia exposure.

GSH-Px catalyzes the reduction of H_2O_2 and organic hydroperoxides, and GST detoxifies metabolites from oxidative reactions (Elia et al. 2006). The strong increase in the activities of GSH-Px and GST is expected to play important roles in protection of fish against ammonia-induced oxidative stress, which might lead to a decrease in cellular GSH content.

However, both the liver and gills maintained a high level of GSH during ammonia exposure (24 h). In the liver, parallel increment in GR activity provided an efficient replenishment of GSH. However, it could maintain a relatively high GSH content in spite of a lack of increment in GR activity in gills. Likewise, an increase in GSH content and GSH/GSSG ratio with a decline in GR activity was reported in the brain of mudskippers exposed to ammonia (Ching et al. 2009). It has been found that the activity of cysteine synthetase, an enzyme producing GSH in cells, was increased under acute hyperammonemic conditions (Murthy et al. 2000). Presumably, these additional routes could have contributed to the increase in GSH levels in FFRC strain common carp exposed to ammonia.

MDA, an end product of lipid peroxidation, is a marker of radical-induced tissue damage (Papadimitriou and Loumbourdis 2002; Del Rio et al. 2005; Lushchak 2011). The content of MDA showed no significant change in both the liver and gills during exposure to 10 mg/L ammonia, which may be due to the antioxidant defense of the glutathione redox system to ammonia-induced oxidative stress. However, after exposure to 30 mg/L ammonia for 48 h, MDA content increased significantly along with the decrease in GSH content and GSH/GSSG ratio. It is possible that an increase in both GSH-Px and GST activities continuously utilized the GSH to scavenge ROS, leading to a depleted GSH pool, resulting in oxidative damage in the liver and gills of FFRC strain common carp. These data showed that the glutathione redox system could be induced under ammonia stress, but the antioxidant response was not sufficient to prevent oxidative damage from the increasing ammonia concentrations.

In conclusion, this is a systematical study which provided a particular insight into the change in H_2O_2 production and the response of the glutathione redox system in the liver and gills of FFRC strain common carp under ammonia stress. Ammonia induced fine-tuning in the levels of GSH, and the transcription and activities of glutathione-related enzymes, which played important roles in scavenging H_2O_2 and preventing lipid peroxidation during exposure to low concentration of ammonia. However, high concentration of ammonia disrupted the glutathione redox system, and the GSH pool and the glutathione-related enzymes could not fully counteract ammonia-induced oxidative damage in both the liver and gills of fish. The glutathione redox system may play an important role in the defense against ammonia stress in the new strain of common carp.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval All applicable international, national, and/or institutional guidelines for the care and use of animals were followed.

Informed consent Informed consent was obtained from all individual participants included in the study.

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