

## **Biochemical Responses of Antioxidant Defense System in Crucian Carp (*Carassius Auratus*) Liver after Exposed To Municipal Sewage Treatment Plant Effluent**

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**Abstract:** *The biochemical responses of antioxidant defense system in crucian carp (*Carassius auratus*) liver after exposed to municipal sewage treatment plant (MSTP) effluent were investigated. In this study, *C. auratus* was exposed to the increasing concentrations (v/v, 0%, 5%, 10%, 20%, 30%, 40% and 50%) of MSTP effluents for 12 days, and the biochemical responses of antioxidant defense system in *Carassius auratus* livers were investigated, including malondialdehyde (MDA), superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GPx), glutathione (GSH), and acetylcholinesterase (AChE). Results indicated that MDA contents and GPx activities in *C. auratus* livers showed a universal increase trend ( $P < 0.05$ ), while GSH contents and AChE activities were decreased significantly ( $P < 0.05$ ) as the increasing concentrations of MSTP effluent. In addition, SOD activities in 5% and 10% treatment groups were sharply higher ( $P < 0.05$ ) but the ones in 40% and 50% treatment groups were significantly lower ( $P < 0.05$ ), while CAT activities in 20%, 30% and 40% treatment groups exhibited a tendency to increase ( $P < 0.05$ ), as compared to the freshwater group. Because of the sensitive responses to the increasing concentrations of MSTP effluent after a 12-day exposure, four biomarkers in *C. auratus* liver, namely MDA, SOD, GPx and GSH, can be used as the biomarkers to assess the water quality of the freshwater environment polluted by MSTP effluent.*

**Keywords:** *Biochemical response, Crucian carp, Effluent, Municipal sewage treatment plant*

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### **I. Introduction**

With the rapid development of society and economy, the discharge of sewage is increased greatly in China. Municipal sewage treatment plants (MSTP) plays an important role in purifying the inflows of sewage. At present, the majority of MSTP usually adopts the secondary sewage treatment technology to purify sewage.<sup>1</sup> Although this treatment technology can effectively remove nitrogen, phosphorus, ammonia nitrogen and other pollutants in the sewage, some pollutants such as surfactants, chlorinated hydrocarbons, heavy metals and endocrine disruptors still exist in the effluent discharged from MSTP.<sup>2</sup> It has been reported that the low concentration (<1%, v/v) of these pollutants in MSTP effluent can exert the toxicity effects on aquatic organisms.<sup>3</sup> This complex mix of MSTP effluent is usually directly discharged into the receiving water, e.g. river, lake and sea, which definitely poses a constant threat to aquatic organism and even human health.<sup>4</sup> Nowadays, it is not feasible that chemical analysis can completely allow the qualitative and quantitative measurement for all pollutants in MSTP effluent, because some trace pollutants in the effluent are under the detection limit of chemical analysis.<sup>5</sup>

Biomarker is defined as the biological measurements at the biochemical, cellular or molecular level that result from the interaction between an organism and environmental chemical, physical or biological agents.<sup>6</sup> Researchers have reported that a causal relationship exists between the biomarker response in aquatic organism and the contaminant exposure in aquatic environment.<sup>7-9</sup> Nowadays, the biomarkers in aquatic organisms have been widely used for monitoring the influence of the pollutants in aquatic environment.<sup>8-11</sup>

To our best knowledge, there is no study to utilize the biomarker response of aquatic organism to monitor MSTP pollution levels in the receiving freshwater areas in China. It is known that the antioxidant defense system of fish species, such as the antioxidant enzymes (SOD, CAT, GPx, etc.) and small antioxidant molecules (GSH, vitamin E, and beta-carotene, etc.), has been widely used as the biomarker in the evaluation for the pollutant levels of environmental water quality. Thus in this study, crucian carp (*Carassius auratus*), a

widely distributed fish species in Chinese rivers and lakes, was exposed under laboratory conditions to increasing concentrations of MSTP effluents for 12 days. Subsequently, the biochemical responses of antioxidant defense system in *C. auratus* livers were investigated, including malondialdehyde (MDA), superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GPx), glutathione (GSH), and acetylcholinesterase (AChE). The aim of the study was to provide an effective biological indicator for the evaluation of MSTP effluent pollution in the freshwater environment.

## II. Materials and Methods

### Materials

MSTP effluent was collected from a draining exit of a sewage treatment plant by Zhulong riverside (Zibo, Shandong province, China). The collected MSTP effluents were transferred to the laboratory within 30 min and kept at 4°C until use.

*C. auratus* were purchased from the local market (Zibo, Shandong, China), kept in 150 L aquarium supplied with aerated freshwater, fed with commercial fish food pellets every day. The average length and weight of fish were  $12.17 \pm 2.51$  cm and  $20.53 \pm 2.78$  g, respectively. The water temperature was kept at  $25 \pm 1$ °C. After acclimated under laboratory conditions for 7 days in aerated freshwater, the healthy and dynamic crucian carps were selected as the targets operated in exposure experiment. The freshwater used in this study was tap water that exposed under the sunshine for at least 3 days for removing the chlorine in tap water completely.

### Exposure experiment

MSTP effluents were diluted by freshwater to increasing concentrations (v/v) of 0%, 5%, 10%, 20%, 30%, 40% and 50%, respectively. For each group, 15 crucian carps were exposed to 100 L of MSTP effluent-aerated freshwater mixture for 12 days. The clean effluent-freshwater was replaced daily during the exposure experiment.

### Preparation of the supernatant of *C. auratus* liver

*C. auratus* were stunned and their livers were collected immediately. *C. auratus* livers were homogenized with cold (4°C) Tris-HCl buffer solutions (100 mM, pH 7.4) (1:9, w (g)/v (mL)) containing 1 mM EDTA-2Na, 10 mM sucrose and 0.8% NaCl. The homogenates were centrifuged at 3,000 r/min for 15 min at 4°C. The supernatants were collected and used for biochemical assays. All operations were performed below 5°C.

### Biochemical assays

The activities of SOD, CAT, GPx, AChE and the contents of MDA, GSH in the supernatants of *C. auratus* livers were investigated by using the commercially-available assay kits (Nanjing Jiancheng Bioengineering Institute, China).

### Data analysis

All experiments were conducted in triplicate. Analysis of variance (ANOVA) was performed using a SPSS package (SPSS 17.0 for Windows, SPSS Inc., Chicago, IL, USA) according to the method of Wan et al.<sup>2</sup> Two-tailed Pearson's correlation analysis was applied to determine the correlation between the concentrations of MSTP effluent and the biochemical responses of antioxidant defense system in *C. auratus* liver, and the statistical significance levels were set as  $P < 0.05$ .

## III. Results And Discussion

The changes for MDA content in *C. auratus* livers were shown in Fig. 1. The MDA content in the treatment groups showed a significant increase compared with the control group ( $P < 0.05$ ) (Fig. 1). MDA is generally considered as a biomarker of lipid peroxidation caused by an excessive accumulation of reactive oxygen species (ROS, such as superoxide radical and peroxide, et al).<sup>12</sup> It has been reported that MSTP effluent can give rise to oxidative stress through the over-production of ROS, which can eventually lead to the deterioration of protein and lipid in organism.<sup>4,13</sup> Therefore, the increasing MDA contents in *C. auratus* liver after exposed to MSTP effluent suggested that MSTP effluent could cause oxidative damage to *C. auratus* liver.<sup>11</sup>

As shown in Fig. 2 and 3, SOD activities in 5% and 10% treatment groups were sharply higher ( $P < 0.05$ ) but the ones in 40% and 50% treatment groups were significantly lower ( $P < 0.05$ ), CAT activities in 20%, 30% and 40% treatment groups exhibited a tendency to increase ( $P < 0.05$ ) while there were no significant differences found in other treatment groups ( $P > 0.05$ ), as compared to the control group.

SOD and CAT are the major antioxidant enzymes that are responsible for the removal of superoxide radicals ( $O_2^{\cdot -}$ ). Specifically, SOD is the enzyme that can catalyze the reaction of  $O_2^{\cdot -}$  to  $H_2O_2$ , and then  $H_2O_2$  is

transformed to H<sub>2</sub>O by CAT.<sup>8</sup> The rises in SOD activities suggested that the pollutants in 5% and 10% of MSTP effluents could produce a certain amount of O<sub>2</sub><sup>•-</sup> and the SOD activity in the cells therefore needs to be enhanced to catalyze the O<sub>2</sub><sup>•-</sup> into H<sub>2</sub>O<sub>2</sub>.<sup>2</sup> However, SOD activities were universally weakened by 40% and 50% of MSTP effluent. Xie et al. found that the excess production of ROS in organism could cause damage to the SOD protein, thus leading to the sharp decrease in SOD activity.<sup>14</sup> In the present study, the results observed in MDA contents in 40% and 50% of MSTP effluent could be ascribed to the over production of ROS induced by MSTP effluent (Fig. 1). Kim and Jung also reported that the over production of O<sub>2</sub><sup>•-</sup> or their transformation production hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) induced by MSTP effluent could inhibit the activity of SOD in pale chub.<sup>15</sup> Thus, it was believed that SOD in *C. auratus* liver is prone to oxidative modification and even inactivation by the excess accumulation of ROS.

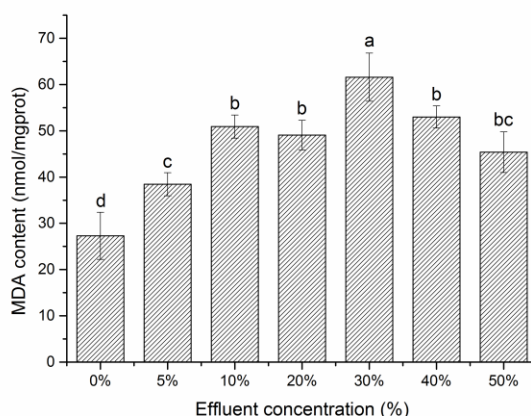
CAT is mainly located in the peroxisomes, and is responsible for the reduction of hydrogen peroxide produced from the metabolism of long chain fatty acids in peroxisomes.<sup>8</sup> The insignificant variation of CAT activities in 5% and 10% of MSTP effluent suggested that the lower concentration of pollutants in MSTP effluent could not completely stimulate the activity of CAT. The increase of CAT activity in 20%~40% treatment groups could reflect that the production of H<sub>2</sub>O<sub>2</sub> was still within the scavenging capacity of CAT.<sup>14, 16</sup> The reduced enzyme activity of CAT in 50% of MSTP effluent indicated that the detoxification capacity of CAT was surpassed by the amount of hydroperoxide products lipid peroxidation.<sup>7</sup>

GPx plays a detoxification role by catalyzing the H<sub>2</sub>O<sub>2</sub> to H<sub>2</sub>O and meanwhile oxidizing glutathione (GSH) to oxidized glutathione (GSSG).<sup>17</sup> In Fig. 4 and 5, GPx activities of *C. auratus* livers in all treatment groups showed a universal increase trend ( $P < 0.05$ ) while GSH contents was decreased gradually as the increasing concentrations of MSTP effluent. Thus, it was believed that GPx activity in *C. auratus* liver was completely induced by the pollutants in MSTP effluent and played a vital role in the remove of peroxide by transforming GSH into GSSG. The different responses of CAT and GPx may indicate different mechanisms for removal of ROS for the two enzymes. CAT is responsible for the reduction of hydrogen peroxide, while GPx catalyzes the reduction of both hydrogen peroxide and lipid peroxides.<sup>18</sup> In addition, besides as the substrate for GPx, GSH also acts as a direct scavenger of oxyradicals and plays an important role in detoxification reaction for ROS.<sup>19</sup> Reduced GSH depletion is often considered a biomarker of environmental stress as observed in fish stressed by environmental pollutants.

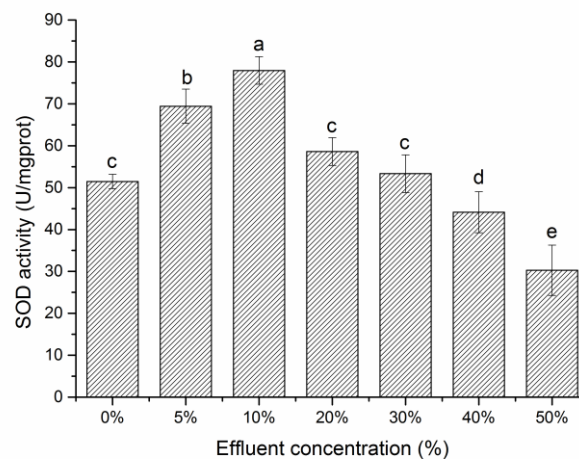
In this study, the AChE activity of *C. auratus* liver was significantly inhibited by MSTP effluent at all exposure concentrations (Fig. 6). AChE is usually considered as a biomarker for reflecting organophosphate/carbamate pollution in environment,<sup>20</sup> because organophosphate/carbamate pesticides have been found to inhibit AChE activity in aquatic organisms.<sup>21</sup> It has been reported that MSTP effluent contains organophosphorus/carbamate pesticides.<sup>22</sup> Consequently, the reduction of AChE activity in this study could be probably attributed to the existence of organophosphorus/carbamate pesticides in MSTP effluent.

Two-tailed Pearson's correlation analysis showed that the biochemical responses of the four antioxidant defense parameters, namely MDA, SOD, GPx and GSH, in *C. auratus* livers showed significant linear relationships with the increasing concentrations of MSTP effluent after a 12-day exposure (Table 1). Thus, MDA, SOD, GPx and GSH in *C. auratus* liver can be used as biomarkers to assess the water quality of the freshwater environment polluted by MSTP effluent.

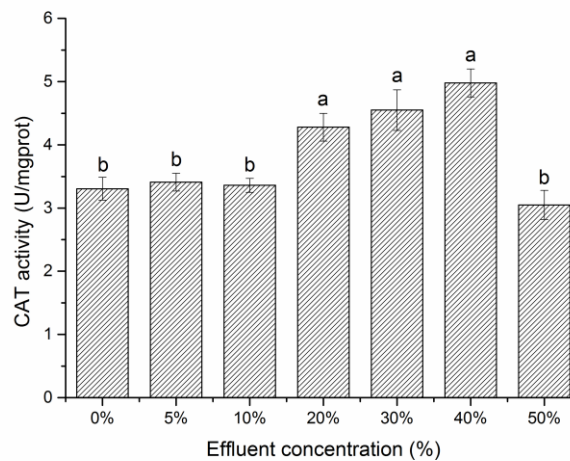
#### IV. Figures And Tables



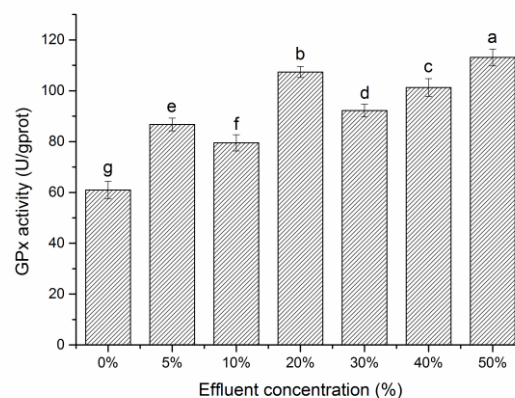
**Figure 1.** Variation of MDA contents in *C. auratus* livers as exposed to MSTP effluent. The lowercase letter represents significance in MDA contents of *C. auratus* liver. Means in columns with different letters were significantly different ( $P < 0.05$ ).



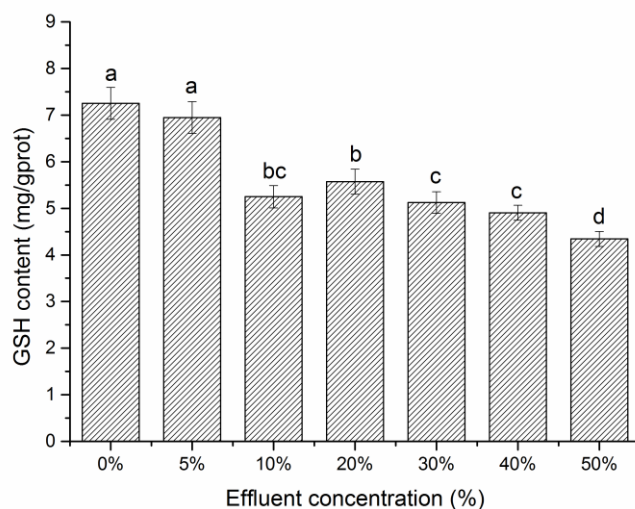
**Figure 2.** Variation of SOD activities in the livers of *C. auratus* exposed to MSTP effluent. The lowercase letter represents significance in SOD activities of *C. auratus* liver. Means in columns with different letters were significantly different ( $P<0.05$ ).



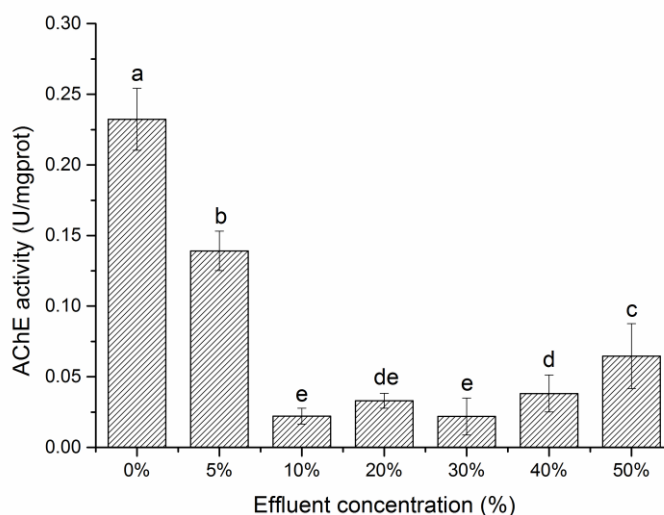
**Figure 3.** Variation of CAT activities in the livers of *C. auratus* exposed to MSTP effluent. The lowercase letter represents significance in CAT activities of *C. auratus* liver. Means in columns with different letters were significantly different ( $P<0.05$ ).



**Figure 4.** Variation of GPx activities in the livers of *C. auratus* exposed to MSTP effluent. The lowercase letter represents significance in GPx activities of *C. auratus* liver. Means in columns with different letters were significantly different ( $P<0.05$ ).



**Figure 5.** Variation of GSH contents in the livers of *C. auratus* exposed to MSTP effluent. The lowercase letter represents significance in GSH contents of *C. auratus* liver. Means in columns with different letters were significantly different ( $P<0.05$ ).



**Figure 6.** Variation of AChE activities in the livers of *C. auratus* exposed to MSTP effluent. The lowercase letter represents significance in AChE activities of *C. auratus* liver. Means in columns with different letters were significantly different ( $P<0.05$ ).

**Table 1** Pearson's correlation coefficients among four biomarkers in *C. auratus* liver after a 12-day exposure to MSTP effluent.

| Biomakers | Correlation coefficients |
|-----------|--------------------------|
| MDA       | 0.869*                   |
| SOD       | -0.765*                  |
| CAT       | 0.336                    |
| GPx       | 0.825*                   |
| GSH       | -0.883*                  |
| AChE      | -0.587                   |

\* $P<0.05$  according to Pearson's correlation analysis.

## V. Conclusion

The present study clearly demonstrated that MSTP effluent could induce oxidative stress in *C. auratus* liver, because the biochemical responses of antioxidant defense system, namely SOD, CAT, GPx, GSH, AChE and MDA, in *C. auratus* liver were significantly altered by MSTP effluent over the expose period. Result



showed MDA, SOD, GPx and GSH in *C. auratus* liver could be applied as the biomarkers to assess the water quality of the receiving freshwater environment polluted by MSTP effluent due to their sensitive responses to the increasing concentrations of MSTP effluent. The present study provides a basis for the establishment of the evaluation system of MSTP effluent-freshwater pollution by applying the four sensitive biomarkers.

### Acknowledgements

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