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# Investigation of Cadmium-Induced Genotoxicity and Oxidative Stress Response in Indian Major Carp, *Labeo rohita*

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# ABSTRACT

We investigated genotoxicity and oxidative stress in the gills of *Labeo rohita* exposed to 33.6, 67.1, and 100.6 mg L<sup>-1</sup> of cadmium chloride at 96 h. Genotoxicity was assessed using single cell gel electrophoresis whereas oxidative stress was monitored through lipid peroxidation induction and antioxidant response parameters, namely reduced glutathione (GSH), glutathione peroxidase, glutathione-S-transferase, superoxide dismutase, and catalase (CAT) activities. Significant (p < .05) effect of both concentration and time of exposure was observed on the extent of DNA damage in treated fish. Similarly, malondialdehyde content, level of GSH, and activities of antioxidant enzymes were significantly elevated in treated groups, except CAT. The increased DNA damage and lipid peroxidation (LPO) content along with fluctuation in antioxidant defense system in fish indicated the interaction of cadmium (Cd) with DNA repair processes and production of reactive oxygen species. Thus, Cd is liable for induction of LPO, alteration of antioxidant defenses, and DNA damage in gills of *L. rohita*.

**Key Words:** cadmium, genotoxicity, *Labeo rohita*, oxidative stress, antioxidant enzymes.

# INTRODUCTION

A variety of contaminants, including toxic heavy metals, are reported to be ubiquitously present in rivers and reservoirs and produce adverse effects on the ecological balance of the recipient environment and harm to a diversity of aquatic organisms (Farombi *et al.* 2007). Cadmium (Cd) is considered as one of the most toxic

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transition elements and a potent industrial hazard, arising primarily from battery, electroplating, pigment, plastics, and fertilizer industries. The dispersion of Cd in the environment has increased in past decades due to its widespread industrial use. Numerous field studies have shown varying degrees of contamination of aquatic systems, via direct or indirect Cd inputs (Hutton 1983; Guinee *et al.* 1999). Lungs, gills, liver, and kidneys are the principal targets of Cd exposure. Moreover, Cd is well known for other various adverse effects (*e.g.*, disturbances of enzyme functions, enhancement of lipid peroxidation influence on mitochondrial functions, and DNA strand break) (Tsuzuki *et al.* 1994; Almeida *et al.* 2002).

Cadmium has many cellular impacts. It triggers induction of some protooncogenes, inhibits the DNA repair system, and alters the activity of several enzymes involved in oxidative stress responses (Waisberg *et al.* 2003; Giaginis *et al.* 2006). The mechanism responsible for the genotoxicity of Cd may involve the direct interaction of Cd<sup>2+</sup> with DNA through the binding of Cd<sup>2+</sup> at guanine, adenine, and thymine bases (Hossain and Huq 2002). Moreover, Cd acts as a mutagen primarily by direct inhibition of an essential DNA mismatch repair, resulting in a high level of genetic instability (Jin *et al.* 2003). Many studies have highlighted the potential role of active oxygen species and free radicals as mediators in the induction of genotoxicity by Cd (Hartwig 1994; Beyersmann and Hechtenberg 1997). Cadmium also has cytotoxic, carcinogenic, and mutagenic effects in test organisms (Monteiro *et al.* 2007; Cambier *et al.* 2010). Cadmium's cellular toxicity and genotoxicity may also be mediated indirectly; cells under oxidative stress display various dysfunctions due to lesions caused by reactive oxygen species to lipids, proteins, and DNA (Risso-de Faverney *et al.* 2001).

The genotoxic effects of environmental pollutants can be monitored using a broad range of both *in vitro* and *in vivo* biomarker assays. However, the comet assay (or single cell gel electrophoresis assay) is gaining popularity over others because it can be performed quickly and provides increased sensitivity for detecting low levels of DNA damage (Nagpure *et al.* 2007; Nwani *et al.* 2010). As a result, the comet assay is being employed to assess the genotoxicity of various chemical compounds in several test organisms (Talapatra *et al.* 2006; Cavas and Konen 2007; Ali *et al.* 2008; Kumar *et al.* 2010).

Reactive oxygen species (ROS) like superoxide radicals, hydrogen peroxide, and hydroxyl radicals are continuously being formed during normal aerobic metabolism. These react with cellular components resulting in protein oxidation, oxidative DNA damage, and lipid peroxidation (Vutukuru *et al.* 2006; Li *et al.* 2011). Lipid peroxidation (LPO) as induced by heavy metal pollution in aquatic organisms is expressed by malondialdehyde (MDA) formation, which represents the secondary LPO product in the thiobarbituric acid reactive substances (TBARS) test (Draper *et al.* 1993). ROS are detoxified as antioxidant defense systems including antioxidants like reduced glutathione and a set of antioxidant enzymes such as glutathione peroxidase, glutathione-S-transferase, superoxide dismutase, and catalase that protect macromolecules against oxidative damage (Ozmen *et al.* 2004). Exposure to contaminants, including Cd, in aquatic ecosystems can enhance intracellular formation of ROS, which could cause oxidative damage to biological systems (Ferreira *et al.* 2005). Changes in the level of antioxidant enzyme activities can be used as

biomarkers of possible oxidative damage in different aquatic organisms (Livingstone 2003; Bechard *et al.* 2008).

Labeo rohita is a fish of the carp family SouthAsia and SoutheastAsia. It is the most important among the three Indian major carp species used in carp polyculture systems and it has high consumer preference in India. L. rohita can serve as an excellent indicator of water quality and environmental pollution in aquatic systems (Vutukuru et al. 2007). The gills are the main target of direct contact with contaminants and play a role in metals' uptake, storage, and transfer to the internal compartments via blood transport. Only limited information is available on Cd-induced DNA damage, LPO induction, and alteration in enzymatic as well as non-enzymatic antioxidant activities in gill cells of L. rohita. Thus, the present study investigated oxidative stress and genotoxic effects of Cd in gills of the Indian major carp L. rohita.

#### METHODS AND MATERIALS

#### **Experimental Animal and Chemical**

Freshwater fish *L. rohita* with similar age and relatedness were procured from a local outlet. After transportation to our laboratory, the specimens were given prophylactic treatment by bathing them twice in 0.05% KMnO<sub>4</sub> solution for 2 min to avoid any dermal infections. They had an average (±SD) wet weight and length of  $21.0 \pm 2.5$  g and  $13.0 \pm 1.2$  cm, respectively. They were then acclimatized for 1 month under laboratory conditions before the start of the experiment. Every effort, as suggested by Bennett and Dooley (1982), was made to maintain optimal conditions during acclimatization.

For the present study, technical-grade admium chloride  $(CdCl_2)$  was procured and used for the study. The chemical was in a monohydrate form, A. R.  $(CdCl_2;$ 98.0% EC, Maximum limits of impurities, Iron 0.0005% and Sulphate 0.005%) and was manufactured by HiMedia Laboratories Pvt. Ltd., Mumbai, India.

# Determination of Exposure Concentrations for Assessing Genotoxicity and Biomarker Response

Exposure concentrations of  $CdCl_2$  for use in the assessment of genotoxicity and biomarker response were determined by first conducting acute toxicity bioassays for the test chemical. These were conducted in static systems to determine the 96 h  $LC_{50}$  in the test species *L. rohita*. For these tests, 10 acclimatized fish specimens per  $CdCl_2$  concentration were randomly exposed for 96 h to test concentrations of 100, 110, 120, 130, 140, 150, 160, and 170 mg L<sup>-1</sup> along with a control set of fish that was exposed to tap water. The experiment was repeated twice to obtain the 96 h  $LC_{50}$ value of the test chemical for the species.

The 96 h LC<sub>50</sub> value of CdCl<sub>2</sub> for freshwater carp *L. rohita* was estimated as 134.18 mg L<sup>-1</sup>. Three exposure test concentrations for assessing biomarker response were selected as fractions of the 96 h LC<sub>50</sub> value: low (SL-I; 1/4th of LC<sub>50</sub> = 33.6 mg L<sup>-1</sup>), medium (SL-II; 1/2th of LC<sub>50</sub> = 67.1 mg L<sup>-1</sup>), and high (SL-III; 3/4th of LC<sub>50</sub> = 100.6 mg L<sup>-1</sup>).

# *In Vivo* Exposure Experiment for Assessment of Genotoxicity and Biomarker Response

Healthy fish with average ( $\pm$ SD) wet weight (21.0  $\pm$  2.5 g) and length (13.0  $\pm$  1.2 cm) was randomly selected and 20 fish transferred to each aquarium in 50 L of water. The exposures were continued up to 96 h and tissues were sampled at 24, 48, 72, and 96 h at the rate of five specimens per sampling duration. Fish maintained in water without the test chemical were used as the control. This study was conducted following the OECD guideline No 203 in the static test conditions (OECD 1992).

On each sampling day, gill tissues were collected from fish for single cell gel electrophoresis assay and biochemical investigations. Temperature, pH, conductivity, dissolved oxygen, chloride, total hardness, and total alkalinity were determined for water on each sampling day by standard methods (APHA 2005).

# Genotoxicity Assay: Single Cell Gel Electrophoresis (SCGE)

The alkaline single cell gel electrophoresis or comet assay (CA) was performed as a three-layer procedure (Singh *et al.* 1988) with slight modifications (Klaude *et al.* 1996). Gill tissue was removed with the help of scissors and 100 mg tissue was homogenized in an ice-cold homogenization buffer (1× Hanks' balanced salt solution [HBSS], 20 mM EDTA, 10% dimethyl sulfoxide [DMSO], pH 7.0–7.5) followed by centrifugation at 3000 rpm at 4°C for 5 min. The cell pellet was then suspended in chilled phosphate buffered saline (PBS). Viability of gill cells was evaluated by the trypan blue exclusion test method (Anderson *et al.* 1994) and the cell suspensions showing viability >84% were further processed for SCGE according to the method described by Ali *et al.* (2008).

Two slides per specimen were prepared and 20 cells per slide (200 cells per concentration) were scored randomly and analyzed using an image analysis system (Komet-5.5 Kinetic Imaging, UK) attached to a fluorescent microscope (Leica) equipped with appropriate filters. The parameter selected for quantification of DNA damage was percent tail DNA (% tail DNA = 100-% Head DNA), as determined by the imaging software (Komet-5.5 Kinetic Imaging, UK).

# **Biochemical Assays**

# Preparation of post-mitochondrial supernatant (PMS)

Gills tissue of 200 mg weight was homogenized in chilled sodium phosphate buffer (0.1 M, pH 7.4) using a Potter-Elvehjem homogenizer. The homogenate was centrifuged at 10,000 g for 10 min at 4°C using a Sigma refrigerated centrifuge (model 3K30) and the resulting supernatant was centrifuged again at 12,500 g for 10 min at 4°C to prepare PMS, which was further used for various biochemical analyses.

# Lipid peroxidation (LPO)

Tissue LPO was measured using the method of Ohkawa *et al.* (1979). Tissue homogenate was incubated with 8.1% SDS (w/v) for 10 min followed by addition of 20% acetic acid (pH 3.5). The reaction mixture was incubated with 1.2% thiobarbituric acid (w/v) for 1 h in boiling water bath. The reaction product was extracted in

butanol-pyridine solution and read at 532 nm. The amount of TBARS was calculated using a molar extinction coefficient of  $1.56 \times 10^5$ /M per cm. The extent of LPO was expressed as nM TBARS formed/h/mg protein.

# Reduced glutathione (GSH)

GSH concentration was measured in gill tissue using the method of Ellman (1959). Samples were treated with trichloroacetic acid (TCA, 10% w/v) and centrifuged at 8900 rpm for 10 min. The supernatant (50  $\mu$ L) was mixed with Tris-HCl buffer (230  $\mu$ L, 0.8 M Tris/HCl, 0.02 M EDTA, pH 8.9) and 20  $\mu$ L of 0.01 M DTNB (2,2'-dinitro-5,5'-dithiobenzoic acid, Ellman's reagent). The reaction mixture was incubated for 5 min at room temperature. The absorbance of GSH-DNTB conjugate was determined at 412 nm and the concentration (nM GSH/mg protein) was calculated using standard calibration.

# Glutathione peroxidase (GPx)

The GPx activity was measured using the procedure of Flohe and Gunzler (1984). One mL of reaction mixture was prepared that contained 0.3 mL of phosphate buffer (0.1 M, pH 7.4), 0.2 mL of GSH (2 mM), 0.1 mL of azide (10 mM), 0.1 mL of  $H_2O_2$  (1 mM), and 0.3 mL of tissue supernatant. After incubation at 37°C for 15 min, reaction was terminated by addition of 0.5 mL of 5% TCA. Tubes were centrifuged at 3000 × g for 5 min and supernatant was collected. 0.2 mL phosphate buffer (0.1 M, pH 7.4) and 0.7 mL of DTNB (0.4 mg/mL) was added to 0.5 mL of reaction supernatant. After mixing, absorbance was recorded at 420 nm. The GPx activity was expressed as nM GSH consumed/min/mg protein.

# **Glutathione S-transferase (GST)**

The activity of GST was determined spectrophotometrically at 25°C by following the formation of GSH conjugate with 1-chloro-2,4-dinitrobenzene (CDNB) at 340 nm using extinction coefficient of 9.6 mM<sup>-1</sup> cm<sup>-1</sup> (Habig *et al.* 1974). The reaction mixture contained 3 mL volume of: 0.1 M sodium phosphate buffer, pH 6.5; 30 mM GSH; 30 mM CDNB in ethanol and the enzyme solution. GST activity was expressed as nM CDNB conjugates/min/mg of protein.

# Superoxide dismutase (SOD)

Tissue SOD activity was analyzed using the method of Kakkar *et al.* (1984). The reaction mixture contained 1.2 mL (0.052 mM) sodium pyrophosphate buffer, 0.1 mL (186 mM) phenazine methosulphate, and 0.3 mL (300 mM) nitro blue tetrazolium. Reaction was initiated by adding 0.2 mL NADH (780 mM) and stopped by adding 1 mL glacial acetic acid. Color intensity of the chromogen was measured at 560 nm. The SOD activity was expressed as units/min/mg of protein.

# Catalase (CAT)

The activity of CAT was measured according to the method of Aebi (1984). The reaction mixture (3 mL volume) contained: 1.98 mL of 50 mM sodium phosphate buffer (pH 7.0), and 1 mL of 30 mM  $H_2O_2$ . The reaction was initiated by adding

20  $\mu$ L of enzyme extract. The CAT activity was determined by measuring the rate of disappearance of H<sub>2</sub>O<sub>2</sub> at 240 nm for 1 min. Enzyme activity was expressed as  $\mu$ M H<sub>2</sub>O<sub>2</sub> decomposed/min/mg protein.

# **Protein estimation**

Protein content in tissue samples was determined using a colorimetric method and BSA as standard (Lowry *et al.* 1951).

#### **Statistical Analyses**

The 96-h LC<sub>50</sub> for CdCl<sub>2</sub> was calculated using IBM SPSS Statistics (version 19.0). One-way analysis of variance (ANOVA) was used to compare the means obtained for various parameters among different concentrations and durations. A *p*-value less than .05 was considered statistically significant. The values for all the biomarkers are expressed as mean  $\pm$  SE (n = 5). Regression analysis was performed to analyze relationships between genotoxicity and various oxidative stress response parameters by using MS Excel.

#### RESULTS

#### **Physicochemical Properties of the Test Solution**

The physicochemical characteristics of the test water measured during experimentation were: temperature 28.6–29.4°C, dissolved oxygen 6.6–7.2 mg L<sup>-1</sup>, and pH 7.4–7.6. The conductivity of the water ranged from 252–284  $\mu$ M cm<sup>-1</sup> while the chloride, total hardness, and total alkalinity varied from 42–48, 160–170, and 220–240 mg L<sup>-1</sup> as CaCO<sub>3</sub>, respectively.

# **DNA Damage**

DNA damage was measured as % tail DNA in gill cells (Figure 1A–C) and indicated that fish exposed to different concentrations of CdCl<sub>2</sub> exhibited significantly higher DNA damage than control specimens. DNA damage in gill tissue was dose and time dependent. The highest level of DNA damage was observed at SL-III concentration ( $20.68 \pm 0.59\%$  tail DNA), followed by SL-II ( $17.24 \pm 0.51\%$  tail DNA), and SL-I concentrations ( $14.51 \pm 0.45\%$  tail DNA) at 96 h post exposure. Lower exposure concentrations or durations resulted in lower amounts of DNA damage (statistically significant at p < .05). The lowest level of DNA damage in gills was observed at 24 h. There was a gradual non-linear increase in the level of DNA damage with progression of the experiment.

# Lipid Peroxidation in Gills

LPO in gill cells increased significantly from 2.57 to 3.78 nM TBARS formed/ h/mg protein (p < .05) at different treatment concentrations of CdCl<sub>2</sub> when compared with the control group (Figure 2A). The maximum increase of LPO was observed at SL-III concentration of CdCl<sub>2</sub> after 96 h post exposure. The amount of

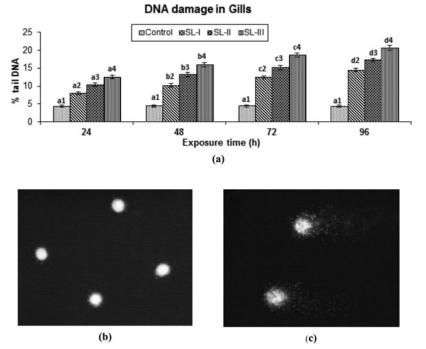


Figure 1. (A) DNA damage in gills of *L. rohita* due to CdCl<sub>2</sub> exposure. Values with different superscript letters (lowercase) differ significantly between exposure durations within concentration. Values with different numeric superscripts differ significantly between concentrations within exposure duration. (B) Control gill cells. (C) Cadmium-exposed gill cells at highest concentration and 96 h post exposure.

LPO in gills of treated groups increased significantly with exposure concentration and time.

#### **Induction of Reduced Glutathione**

Exposure of *L. rohita* to CdCl<sub>2</sub> for 48, 72, and 96 h caused significant (p < .05) increase in GSH level in the gills at all concentrations as compared to controls (Figure 3). GSH level in gill tissue was dose-dependent with the highest GSH level observed at SL-III concentration ( $5.11 \pm 0.29$  nM GSH/mg protein) at 96 h post exposure. No significant effects among durations of exposure were observed in fish specimens exposed except at the SL-III test concentration and 96 h post exposure.

# **Antioxidant Enzymes Activities**

Significant (p < .05) effect of GPx activities was observed in gills of *L. rohita* at all three test concentrations of CdCl<sub>2</sub> as compared to control (Table 1). The highest activities of GPx in gill tissue (44.41 ± 2.05 nM GSH consumed/min/mg protein) were observed at the SL-III concentration and 96 h post exposure. Significantly less GPx activity was observed at the SL-II (medium) concentration and 24 h post

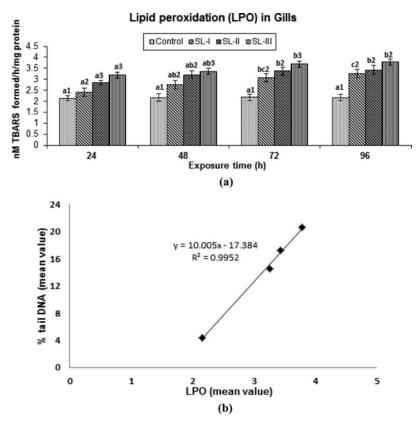
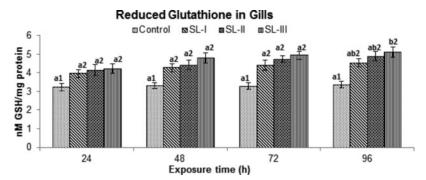


Figure 2. (A) Lipid peroxidation induced by CdCl<sub>2</sub> in gills of *L. rohita*. The values are expressed in nanomoles of TBARS formed/h/mg of protein. Values with different superscript letters (lowercase) differ significantly between exposure durations within concentration. Values with different numeric superscripts differ significantly between concentrations within exposure duration. (B) Regression analysis of mean data of lipid peroxidation (LPO) and % tail DNA in fish exposed to CdCl<sub>2</sub>. Values comprise control and exposed groups at 96 h post-exposure.

exposure. GPx activity in gills of treated groups increased in a dose-dependent manner.

Cadmium chloride induced a significant (p < .05) increase in GST activity at most test concentrations and exposure durations; significant increase did not occur for the SL-I concentration and 24 h post exposure (Table 1). The highest increased GST activity was observed in gills (234.04 ± 8.58 nM CDNB conjugates/min/mg of protein) at the SL-III concentration of CdCl<sub>2</sub> and 96 h post exposure. No significant effect among durations of exposure was observed in GST activity except at 96 h post exposure and the SL-III test concentration.

A significant (p < .05) elevation in SOD activities was observed at all three test concentrations of CdCl<sub>2</sub> as compared to control group except at the SL-I concentration



**Figure 3.** Reduced glutathione level in gill tissues of *L. rohita* as influenced by CdCl<sub>2</sub> exposure. The values are expressed in nanomoles of GSH/mg of protein. Values with different superscript letters (lowercase) differ significantly between exposure durations within concentration. Values with different numeric superscripts differ significantly between concentrations within exposure duration.

and 24 h post exposure (Table 2). SOD activity in gills of *L. rohita* increased significantly from 4.02 to 5.32 units/min/mg of protein with highest elevation at SL-III and 96 h post exposure. SOD activity increased with dose and exposure duration.

CAT activity in gills decreased by 171.69 to 141.29  $\mu$ M H<sub>2</sub>O<sub>2</sub> decomposed/ min/mg protein at different treatment concentrations of CdCl<sub>2</sub> when compared with control group. The smallest significant (p < .05) decrease in CAT activity was observed at the SL-III concentration and 24 h post exposure (Table 2). The CAT activity in gills depleted in a dose-dependent manner and highest depletion observed

A1 .		Exposure time (h)					
Antioxidant enzymes	Conc.	24	48	72	96		
GPx GST	Control SL-I SL-II SL-III Control SL-I	$\begin{array}{c} 27.16 \pm 1.71^{\mathrm{al}} \\ 30.84 \pm 1.18^{\mathrm{al}} \\ 34.31 \pm 1.46^{\mathrm{a2}} \\ 37.86 \pm 1.10^{\mathrm{a2}} \\ 167.79 \pm 4.59^{\mathrm{a1}} \\ 187.09 \pm 8.31^{\mathrm{a12}} \end{array}$	$\begin{array}{c} 28.35\pm1.41^{a1}\\ 34.39\pm1.65^{ab2}\\ 36.75\pm1.95^{ab23}\\ 40.28\pm1.32^{ab3}\\ 168.41\pm6.56^{a1}\\ 105.28\pm5.25a^2 \end{array}$	$\begin{array}{c} 29.73 \pm 1.58^{\mathrm{al}} \\ 36.57 \pm 1.87^{\mathrm{ab2}} \\ 39.28 \pm 1.74^{\mathrm{ab2}} \\ 41.62 \pm 2.15^{\mathrm{ab2}} \\ 166.92 \pm 5.87^{\mathrm{al}} \\ 205.46 \pm 8.01^{\mathrm{a2}} \end{array}$	$28.84 \pm 1.36^{a1}$ $39.04 \pm 2.16^{b2}$ $42.73 \pm 1.87^{b2}$ $44.41 \pm 2.05^{b2}$ $165.62 \pm 6.09^{a1}$ $211.66 \pm 7.44^{a2}$		
	SL-II	$196.75 \pm 9.45^{\mathrm{a2}}$		$203.40 \pm 8.01^{42}$ $211.74 \pm 7.39^{a2}$ $226.44 \pm 8.10^{ab2}$	$211.00 \pm 7.44$ $218.14 \pm 5.95^{a2}$ $234.04 \pm 8.58^{b2}$		

Table 1.Effect of CdCl2 on glutathione peroxidase (nM GSH consumed/min/<br/>mg protein) and glutathione-S-transferase (nM CDNB conjugates/<br/>min/mg of protein) enzyme activities in gills of *L. rohita*.

Values with different superscript letters (lowercase) differ significantly between exposure durations within concentration. Values with different numeric superscripts differ significantly between concentrations within exposure duration.

**Table 2.** Effect of  $CdCl_2$  on superoxide dismutase (units/min/mg of protein)and catalase ( $\mu M H_2O_2$  decomposed/min/mg protein) enzymeactivities in gills of L. rohita.

Antioxidant		Exposure time (h)					
enzymes	Conc.	24	48	72	96		
SOD	Control	$3.18\pm0.14^{\rm al}$	$3.26\pm0.16^{\rm al}$	$3.32\pm0.20^{\mathrm{a}1}$	$3.29\pm0.21^{\rm al}$		
	SL-I	$3.69 \pm 0.20^{\mathrm{a}12}$	$3.91 \pm 0.21^{ m ab2}$	$4.19 \pm 0.25^{\mathrm{ab2}}$	$4.43 \pm 0.25^{\mathrm{b2}}$		
	SL-II	$4.02 \pm 0.24^{\mathrm{a}2}$	$4.26\pm0.25^{\rm ab2}$	$4.61 \pm 0.22^{ m ab2}$	$4.95 \pm 0.27^{\mathrm{b2}}$		
S	SL-III	$4.25 \pm 0.23^{\mathrm{a2}}$	$4.62 \pm 0.22^{ m ab2}$	$4.89 \pm 0.28^{ m ab2}$	$5.32 \pm 0.30^{\mathrm{b2}}$		
CAT	Control	$184.38 \pm 6.16^{\rm al}$	$182.13 \pm 4.84^{\rm a1}$	$180.92 \pm 3.91^{\mathrm{al}}$	$183.45 \pm 4.35^{\rm al}$		
	SL-I	$171.69 \pm 4.69^{\rm a12}$	$164.85 \pm 6.01^{\rm ab12}$	$157.42 \pm 5.14^{\rm ab2}$	$151.96 \pm 3.15^{\rm b2}$		
	SL-II	$165.82 \pm 6.32^{\rm al2}$	$157.74 \pm 4.15^{\rm ab2}$	$152.45 \pm 4.32^{\mathrm{ab2}}$	$147.94 \pm 4.08^{\rm b2}$		
	SL-III	$161.69 \pm 5.66^{\rm a2}$	$153.51 \pm 3.22^{\rm a2}$	$148.59 \pm 3.51^{\rm ab2}$	$141.29 \pm 3.67^{\rm b2}$		

Values with different superscript letters (lowercase) differ significantly between exposure durations within concentration. Values with different numeric superscripts differ significantly between concentrations within exposure duration.

at the SL-III test concentration and 96 h post exposure. A significant depletion among exposure durations was observed at 96 h post exposure.

# **Regression Analysis**

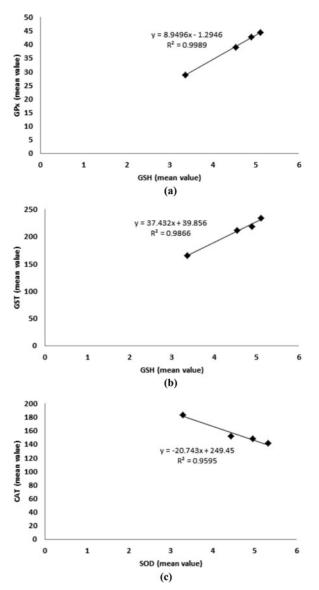
Regression analysis (Figure 2B) showed a significant positive correlation ( $r^2 = 0.995$ ) between mean values of% tail DNA and LPO data at 96 h post exposure. This shows that there exists a relationship between genotoxic and oxidative stress-inducing effects of Cd and oxidative stress might be contributing to gentoxicity of Cd in *L. rohita*.

Significant positive correlations were estimated between GSH and both glutathione dependent antioxidant enzyme GPx ( $r^2 = 0.999$ ) and GST ( $r^2 = 0.987$ ) at 96 h post exposure (Figures 4A and 4B). Regression analysis between reduced glutathione and both enzymes (GPx and GST) indicates that GSH has a role in detoxification mechanism along with GPx and GST. A negative correlation ( $r^2 =$ 0.9595) between SOD and CAT shows that Cd might be directly responsible for suppressing CAT activity and production of free radicals (Figure 4C).

# DISCUSSION

The present study investigated genotoxicity and oxidative stress responses in the gill of *L. rohita* exposed to Cd. In aquatic organisms, the gill is the osmoregulatory surface tissue and primary site of uptake of waterborne pollutants and it is the major target organ for acute copper, zinc, cadmium, and silver toxicity to freshwater fish (Karlsson-Norrgren *et al.* 1985; Olsson *et al.* 1998). Failure of the gill to function during acute exposure to Cd can lead to the death of fish. Previous studies show that Cd possibly moves across the epithelial layer of fish gills via calcium channels (Farag *et al.* 1994). Calcium exerts considerable control over permeability of the gills and displacement of calcium could stimulate ion loss and water uptake (Verbost *et al.* 





**Figure 4.** Regression analysis between mean data of (A) reduced glutathione (GSH) and glutathione peroxidase (GPx), (B) reduced glutathione (GSH) and glutathione-S-transferase (GST), and (C) superoxide dismutase (SOD) and catalase (CAT) in fish exposed to CdCl<sub>2</sub> at 96 h post exposure.

1989). Cadmium affects calcium balance in rainbow trout (Verbost *et al.* 1987) and induces damage in gill structure of zebra fish, rainbow trout, and tilapia (Karlsson-Norrgren *et al.* 1985; Pratap and Wendelaar Bonga 1993).

The comet assay indicates that Cd exposure of *L. rohita* gill cells elicits DNA damage leading to DNA strand breaks. The finding from our study demonstrates that % tail DNA in gill cells increases with increasing Cd concentration and exposure duration. The highest level of DNA damage observed at the highest Cd concentration after 96 h post exposure. This may be due to continuous exposure to Cd, which leads to the enhancement of ROS formation in gill cells. ROS can react with susceptible biological macromolecules and cause LPO, DNA damage, and protein oxidation (Li *et al.* 2005; Lesser 2006; Cambier *et al.* 2010). Thus, oxidative DNA damage may be the most likely mechanism of genotoxicity of Cd. Moreover, Cd also alters the repair mechanism of DNA by inhibition of an essential DNA mismatch repair (Shi *et al.* 2005). Both the interaction with DNA repair processes and the induction of oxidative DNA damage may account for Cd's genotoxicity (Hartwig 1994).

Lipid peroxidation is acknowledged as being highly deleterious and deriving from attacks by hydroxide free radical, resulting in oxidative damage to organs or tissues (Shi et al. 2005; Pandey et al. 2008). Hydroxide free radical is produced from H<sub>2</sub>O<sub>2</sub> via the Fenton reaction (Evans and Halliwell 1994). CAT is the primary enzyme in scavenging  $H_2O_2$ , so when CAT activity is inhibited, more  $H_2O_2$  is available for production of hydroxide free radical. In this study, the decrease of CAT activity in gills of L. rohita, together with the increase of SOD activity, could explain the increased MDA level in gills. The elevated MDA level directly reflected that the Cd may cause lipid peroxidation due to production of free radicals. Both the LPO levels and the % Tail DNA increased simultaneously. It is well established that Cd with oxidative stress has the potential to attach on lipids and DNA resulting in clastogenic and molecular damage (Raisuddin and Jha 2004; Jha 2008). Certain fish and non-piscine species (mollusks) have shown a good correlation between level of pollution and extent of genetic damage (Cavas and Konen 2007; Jha 2008). Increase in DNA damage and induction of lipid peroxidation in Cd-treated fish was positively correlated ( $r^2 = 0.9952$ ) at 96 h post exposure.

Superoxide dismutase catalyzes conversion of the superoxide anion radical to molecular oxygen and hydrogen peroxide and this may protect against superoxide-induced oxidative damage (Vutukuru *et al.* 2006). Contaminants such as heavy metals including Cd (Almeida *et al.* 2002; Asagba *et al.* 2008), oils, and phenols (Zhang *et al.* 2005) affect SOD activity in fish (Oruc *et al.* 2004). In the present study, SOD activity in the gills of Cd-treated fish was significantly higher than control fish, possibly in response to toxic effect of superoxide anion radical.

CAT catalyses the conversion of  $H_2O_2$  to water and molecular oxygen and protects biological systems against ROS (Romeo *et al.* 2000). In this study, CAT in Cd-treated groups decreased significantly compared to controls with increased concentration and exposure duration, which indicated that the abilities to protect against  $H_2O_2$ were reduced (Vutukuru *et al.* 2006). The explanation for reduction in CAT activity by Cd exposure may be related to the direct binding of the metal to –SH groups of the enzyme molecule. The reduced CAT activity in gill may also be associated with the compensatory high activity of GPx, which acts as a defense against the formation of  $H_2O_2$  or effective antioxidant responses due to a higher renovation of gill epithelium (Ferreira *et al.* 2005). The negative correlation between increased activity of SOD and decreased activity of CAT suggest that superoxide radicals neutralized by SOD protects the biological system from free radicals while catalase levels may not be adequate to neutralize hydrogen peroxides, resulting induction of lipid peroxidation.

Glutathione plays an important role in modulation of metal-induced LPO and works as a reducing substrate in oxidative reactions (Schlenk and Rice 1998; Stohs *et al.* 2000). Cadmium exposure caused an increase in GSH content in gill cells of *L. rohita.* Some investigators opined that such Cd-mediated induction of cystolic thiol-rich proteins (GSH and metallothionein) could protect against secondary-induced oxidative stress, probably by maintaining a reduced state in the cell (Tort *et al.* 1996). Transient increase in GSH in our study may be a primary protective response of cells against Cd-induced oxidative stress while increased levels of GSH were probably not sufficient to get rid of oxidative stress, as concomitant increase in LPO amount has been observed in gills of *L. rohita.* 

Both GST and GPx are glutathione dependent enzymes and with the assistance of GSH, they play an important role in protecting organs from oxidative stress (Hayes and Strange 1995). GST catalyzes conjugation of many xenobiotics to the tripeptide glutathione while GPx provides optimal protection against oxidative stress (Hermes-Lima 2004) by scavenging  $H_2O_2$  or ROOH, and has been considered to be complementary to CAT (Halliwell and Gutteridge 1999). In the present study, GPx and GST activity increased significantly in gills with increased concentration and exposure duration, which was consistent with most investigations (Atif *et al.* 2005; Bouraoui *et al.* 2008), suggesting that GPx and GST activity could be elevated to resist the Cd-induced toxicity. The positive correlation between GSH and both enzymes (GPx and GST) also shows that increased amount of GSH as well as elevated activity of GPx and GST in Cd-treated fish groups may protect biological systems from oxidative stress.

# CONCLUSION

The present study revealed that Cd is responsible for inducing DNA damage, lipid peroxidation, and oxidative stress in *L. rohita* gills. The effect of Cd on DNA damage, lipid peroxidation, non-enzymatic antioxidant, and antioxidant enzymes in gill tissue were concentration and time dependent. The tissue antioxidants seem to counteract the impact of reactive oxygen species while the antioxidant enzymes were unable to prevent DNA damage and lipid peroxidation in the gills due to acute exposure of the metal.

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