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Geno- and cytotoxicity induced on *Cyprinus carpio* by aluminum, iron, mercury and mixture thereof

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ABSTRACT

Metals such as Al, Fe and Hg are used in diverse anthropogenic activities. Their presence in water bodies is due mainly to domestic, agricultural and industrial wastewater discharges and constitutes a hazard for the organisms inhabiting these environments. The present study aimed to evaluate geno- and cytotoxicity induced by Al, Fe, Hg and the mixture of these metals on blood of the common carp *Cyprinus carpio*. Specimens were exposed to the permissible limits in water for human use and consumption according to the pertinent official Mexican norm [official Mexican norm NOM-127-SSA1-1994] Al (0.2 mg L⁻¹), Fe (0.3 mg L⁻¹), Hg (0.001 mg L⁻¹) and their mixture for 12, 24, 48, 72 and 96 h. Biomarkers of genotoxicity (comet assay and micronucleus test) and cytotoxicity (caspase-3 activity and TUNEL assay) were evaluated. Significant increases relative to the control group ($p < 0.05$) were observed in all biomarkers at all exposure times in all test systems; however, damage was greater when the metals were present as a mixture. Furthermore, correlations between metal concentrations and biomarkers of geno- and cytotoxicity were found only at certain exposure times. In conclusion, Al, Fe, Hg and the mixture of these metals induce geno- and cytotoxicity on blood of *C. carpio*.

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1. Introduction

Chemical contaminants reach water bodies through diverse sources of exposure. The relevance of heavy metal studies resides in the characteristics of these elements: high toxicity and persistence, and rapid bioaccumulation by aquatic organisms (Cervantes

and Moreno, 2010; Rosas, 2001).

A heavy metal is any metallic element having a relative high density and toxicity even at very low concentrations. Such metals include Al, Fe and Hg, among others (Lucho-Constantino et al., 2005). Aluminum is one of the most abundant metals in Earth's crust and is used in diverse anthropogenic activities (García-Medina et al., 2010). Iron is a micronutrient; it is essential for all living organisms and plays a major role in vital biochemical activities such as oxygen transport to tissue, electron transfer, and catalysis (Aisen et al., 2001; Pérez et al., 2005). It is also involved in DNA synthesis and is a component part of hemoglobin, myoglobin, cytochromes and diverse enzymes (Huang et al., 2015). Iron is naturally present in ground and surface water. In aquatic ecosystems, Hg presence results mainly from its atmospheric deposition due to anthropogenic activities (Chan et al., 2003; Morel et al.,

Abbreviations: AAS/AES, atomic absorption/emission spectrophotometry; ICP-MS, inductively coupled plasma-mass spectrometry; MeHg, methylmercury; MNi, micronuclei; TUNEL, terminal deoxynucleotidyl transferase-mediated dUTP nick-end labeling; pNA, p-nitroaniline; ROS, reactive oxygen species; SEM, standard error of the mean

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1998; Pacyna and Pacyna, 2002). In water, inorganic Hg is transformed by bacterial action to methylmercury (MeHg), a highly toxic compound which accumulates in sediment (Chan et al., 2003; Driscoll et al., 2007; Morel et al., 1998; Ullrich et al., 2001).

Heavy metals enter the aquatic ecosystem through hospital, municipal and industrial residues which are discharged without prior treatment, resulting in the presence of high concentrations (Costa et al., 2015; Fernández-Dávila et al., 2012; García and Dorronsoro, 2005; Kim et al., 2013).

In Mexico, research on the environmental occurrence of this type of contaminants includes studies by Avila-Pérez et al. (1999), who detected Fe (880–33080 mg L⁻¹) and Hg (17–181 mg L⁻¹), among other metals, in surface water and the water column of José Antonio Alzate Reservoir, in the State of Mexico. In Aguascalientes, Al and Fe concentrations were measured in the Río San Pedro, concentrations of 0.06–62.60 and 1.04–22.60 mg L⁻¹ respectively were detected (Guzmán et al., 2010). The presence of Fe (1.09–1.80 mg L⁻¹) has been reported in Río Chihuahua water (Gutiérrez et al., 2008), while in the State of Mexico Al (6.04–24.45 mg L⁻¹), Fe (1.37–5.10 mg L⁻¹) and Hg (<0.001 mg L⁻¹) were found in Madín Dam (González-González et al., 2014), and Hg (0.021 mg L⁻¹) was detected in wastewater from a public hospital (Neri-Cruz et al., 2015).

The constant release of contaminants into the aquatic ecosystem implies the presence of a mixture of these. Trace amounts or minimum quantities of metals can have positive or negative effects on aquatic organisms. Slight variations in their concentrations, decreases as well as increases, can have toxic consequences on aquatic organisms (Wittmann, 1981). In fish, Al is neurotoxic; it induces gill damage due to increased mucus production which affects osmoregulation and respiration (Exley et al., 1997; Ward et al., 2006), causing hypoxia, hypercapnia, metabolic acidosis and eventually respiratory insufficiency (Allin and Wilson, 2000; Røyset et al., 2005). Furthermore, it elicits diverse hematological changes (Bhagwant and Bhikajee, 2000; García-Medina et al., 2010). While Fe has diverse biological functions, at high concentrations this metal may induce DNA damage, hemochromatosis and carcinogenesis (Huang, 2004; Mello-Filho and Meneghini, 1991), the principal organs so damaged being the heart, liver and endocrine glands (Italia et al., 2015). In fish, the main damage induced by Hg is at the central nervous system level (Berntssen et al., 2003).

Several studies have shown that exposure to contaminants, including heavy metals, stimulates ROS (reactive oxygen species) production in the cell (Li et al., 2006; Sinha et al., 2007), contributing to oxidative stress generation and, consequently, DNA damage. Metallic ions interact with cellular components such as nuclear proteins and DNA, inducing DNA damage and conformational changes which may lead to cell cycle modulation, carcinogenesis or apoptosis (Beyersmann and Hartwig, 2008; Chang et al., 1996; Wang and Shi, 2001).

Fish are used as sensitive indicators of genotoxic and mutagenic changes since they bioaccumulate contaminants present in water (Yadav and Trivedi, 2009) and their response to chemical exposure is similar to response in higher vertebrates (Al-Sabti, 1991). The common carp *Cyprinus carpio* is a commercial species which, due to its economic importance and wide geographic distribution, has been proposed as a test aquatic organism in toxicological assays (De Boeck et al., 2007; Oruç and Usta, 2007). In Mexico, this species is consumed by humans and is frequently cultured in water bodies contaminated with diverse xenobiotics including heavy metals.

Therefore, the aim of the present study was to evaluate the geno- and cytotoxicity induced by Al, Fe, Hg and the mixture of these metals on blood of *C. carpio*.

2. Materials and methods

2.1. Specimen procurement and maintenance

Common carp (*Cyprinus carpio*) specimens were obtained from the aquaculture facility in Tiacaque (State of Mexico). Polyethylene bags containing water and oxygen were used to transport the fish to the Environmental Toxicology Laboratory at the Department of Chemistry (Universidad Autónoma del Estado de México). Carp were maintained for a 15-d acclimation period in 160-L fish tanks (a fish for each two liters of water was placed) with synthetic culture medium (pH 7.4), at room temperature, with constant aeration and a 12:12 h light/dark photoperiod, and were fed Pedregal Silver™ fish food.

2.2. Experimental design

Specimens 20.15 ± 0.28 cm long and weighing 55.67 ± 6.2 g were maintained under conditions similar to those used for acclimation. To test systems with six carp each was added Al (0.2 mg L⁻¹), Fe (0.3 mg L⁻¹) or Hg (0.001 mg L⁻¹) according to the permissible limits in water for human use and consumption in the official Mexican norm (NOM-127-SSA1-1994). The assay also included a test system to which was added the mixture of all three metals at the same concentrations, as well as a control system. Exposure times were 12, 24, 48, 72 and 96 h. Static systems were used without renewal of test solution. The assay was performed in triplicate, using a total of 450 fish. At the end of the exposure time, carp were removed from the systems and placed in a fish tank containing a xylocaine solution (0.02 mg mL⁻¹, AztraZeneca, Naucalpan, State of Mexico), to anesthetize them prior to collecting a blood sample (2 mL) from the caudal vein, using a heparinized 2-mL hypodermic syringe.

2.2.1. Comet assay

The procedure proposed by Tice et al. (2000) was used. Previously prepared microscope slides were used to obtain the sample. Frosted slides were covered with 200 µL of 1% agarose (Sigma-Aldrich, St Louis, MO) and maintained at room temperature until dry. Whole blood samples (10 µL) were mixed with 75 µL of 0.7% agarose (Sigma-Aldrich, St Louis), and 50 µL of this mixture was spread on the initial agarose layer and solidified on ice. The slides were placed inside a Coplin jar with lysis solution [2.5 M NaCl, 100 mM EDTA, 10 mM Tris (all Sigma-Aldrich, Toluca, Mexico), 10% dimethyl sulfoxide (DMSO, J.T. Baker, Center Valley, PA) and 1% Triton X-100 (Sigma-Aldrich, St Louis)] pH 10, for 1 h at 4 °C. The slides were then placed in the electrophoresis chamber with alkaline solution [300 mM NaOH and 1 mM EDTA (both Sigma-Aldrich, Toluca)] at pH 13 for 20 min. Electrophoresis was performed at 300 mA and 25 V (4 °C, 20 min, field strength: 0.8 V/cm) and was halted with a neutralization buffer [0.4 M Trizma base (Sigma-Aldrich, St Louis) pH 7.4]. The DNA was stained with 50 µL ethidium bromide (10 mg mL⁻¹; Sigma-Aldrich, St Louis) and examined in an epifluorescence microscope (Motic BA410) equipped with a Moticam Pro CCD digital camera.

2.2.2. Micronucleus test

Whole blood from each specimen was fixed with pure ethanol (Mallinckrodt Baker, State of Mexico) on a slide for 5 min, then stained with 10% Giemsa (Hycel, Mexico City, Mexico) for 9 min. A light microscope was used to examine a total of 1000 cells per sample. Results were expressed as the total number of micronucleated cells per 1000 cells (Çavaş and Ergene-Gözükara, 2005).

2.2.3. Caspase-3 activity

2.2.3.1. Cellular extract preparation.

Jurkat cells (ATCC # TIB-152)

were cultured in RPMI-1640 medium containing 10% fetal bovine serum in a humidified, 5% CO₂ atmosphere at 37 °C, as recommended by ATCC. Cell density was initially adjusted to 10⁶ cells mL⁻¹, and anti-Fas mAb (50 ng mL⁻¹, MBL International) was added to the cells as a positive (apoptosis inducer) control. To inhibit apoptosis samples were added 125 µL of the inhibitor Z-VAD-FMK (carbobenzoxy-valyl-alanyl-aspartyl-(O-methyl)-fluoromethylketone, 20 mM) at the same time as the anti-Fas mAb. The samples were incubated overnight for 16 h at 37 °C in a humidified, 5% CO₂ atmosphere, then centrifuged at 450g and 4 °C for 10 min, and the cells were harvested. The cell pellet was maintained on ice, washed with cold phosphate-buffered saline (PBS) and resuspended in cell lysis buffer. The final cell concentration was 10⁸ cells mL⁻¹. Cells were lysed by freeze-thaw, then incubated on ice for 15 min. The cell lysate was centrifuged at 15,000g and 4 °C for 20 min, and the supernatant was collected.

2.2.3.2. Colorimetric assay. A CaspACE™ colorimetric assay kit (Promega, Madison, WI) and an UltraCruz™ microplate with flat-bottom wells were used. The blank was prepared white reaction mixture [32 µL caspase buffer (312.5 Mm HEPES pH 7.5, 31.25% saccharose, 0.3125% CHAPS (3-[(3-cholamidopropyl)-dimethylammonio]–1-propane-sulfonate)), 2 µL DMSO, 10 µL dithiothreitol (DTT, 100 mM)] and 54 µL deionized water. For the control group and groups exposed to Al, Fe, Hg or the mixture, the following were used: reaction mixture, 20 µL blood and 54 µL deionized water; for the positive control and for inhibited apoptosis samples: reaction mixture, 20 µL cellular extract and 34 µL deionized water. After all solutions had been transferred, 2 µL of substrate DEVD-pNA was added to each well. The microplate was covered with parafilm and incubated for 4 h at 37 °C. Absorbance was read at 405 nm and the specific activity of caspase-3 was calculated. Results were expressed as nM free pNA h⁻¹ µg⁻¹ protein. Total protein content was determined by the Bradford (1976) method.

2.2.4. Determination of total protein content

To 25 µL of supernatant was added 75 µL deionized water and 2.5 mL Bradford's reagent [0.05 g Coomassie blue (Sigma-Aldrich, Toluca), 25 mL of 96% ethanol (Sigma-Aldrich, Toluca) and 50 mL H₃PO₄ (Vetec, Mexico City) in 500 mL deionized water]. The tubes were shaken and allowed to rest for 5 min prior to reading absorbance at 595 nm. Results were interpolated onto a standard bovine serum albumin curve.

2.2.5. TUNEL assay

The ApopTag S7110 fluorescein kit (Chemicon, Temecula, CA) was used. Whole blood samples (300 µL) were centrifuged at 800g and 4 °C for 5 min; the cell pellet was resuspended with 50 µL of the mounting medium and 1 µL of cells was placed on a slide with poly-L-lysine (Sigma-Aldrich, St Louis), dried at 60 °C for 5 min,

fixed in cold acetone for 10 min, and hydrated in successive changes (from 100% to 50%) of xylene, ethanol (both Mallinckrodt Baker) and water. The cells were treated with proteinase K (20 µg mL⁻¹, Fluka-Sigma-Aldrich, Toluca) for 10 min, then washed with PBS (NaCl 0.138 M, KCl 0.0027 M, Sigma-Aldrich, Toluca) pH 7.4. Next, 60 µL equilibrium buffer was added, followed by incubation with 65 µL TdT enzyme for 60 min at 37 °C. The cells were washed with PBS prior to adding the anti-FITC conjugate that was used as a cell marker, then maintained at room temperature for 30 min, washed with PBS, stained with propidium iodide (1.5 µL mL⁻¹, Sigma-Aldrich, St Louis) and examined in the epifluorescence microscope with digital camera. The negative control sample consisted of cells treated as described but without addition of TdT, while the positive control sample consisted of cells treated with DNase I (1 µL mL⁻¹, Sigma-Aldrich, St Louis). A total of 100 cells were examined per specimen, and apoptosis was expressed as the percentage of TUNEL-positive cells per 100 cells.

2.3. Determining concentrations of Al, Fe, Hg and mixture thereof

Whole blood samples (0.2–0.4 mL) were digested with 8 mL HNO₃ (65% Suprapur-grade) and 2 mL H₂O₂ (30% v/v Suprapur-grade, Merck Millipore, Barcelona, Spain) in a Milestone Ethos Plus microwave oven (Monroe, CT) for 10 min at 180 °C after 15-min ramping. Digested samples were diluted with 4 mL ultrapure water (18.2 MΩ cm) and refrigerated until analysis. The total Al and total Hg analyses were performed by inductively coupled plasma-mass spectrometry (ICP-MS) using a Thermo Electron XSeries II system with collision cell technology (CCT), operated in standard mode to obtain data on ²⁷Al and ²⁰²Hg. Iron was analyzed by atomic absorption/emission spectrophotometry (AAS/AES) using a Varian SpectraAA 50 system equipped with air-acetylene flame and a hollow-cathode lamp. Detection limits (µg L⁻¹) were: Al 1.185, Hg 0.058, and Fe 0.016. Quantification limits (µg L⁻¹) were: Al 3.951, Hg 0.083, and Fe 0.057.

As reference materials, muscle and liver of dogfish were analyzed; the percentage of recovery in muscle was 95.9 ± 9.2% for Al, 102.3 ± 7.5% for Hg and 92.9 ± 2.9% for Fe, while in liver it was 95.4 ± 12.4% for Al, 102.4 ± 24.2% for Hg and 106.2 ± 6.2% for Fe.

2.4. Statistical analysis

Results were statistically evaluated by one-way analysis of variance (ANOVA). Data normality and homoscedasticity were verified by Shapiro-Wilk and Levene tests, respectively. The Bonferroni *post hoc* test was used to evaluate significant differences, with *p* set at < 0.05. Pearson's correlation analysis was performed to find potential correlations of concentrations of Al, Fe, Hg and the mixture with biomarkers of geno- and cytotoxicity. Sigstastat v2.03 was used.

Table 1
Concentrations of Al, Fe, Hg and mixture thereof in blood of *C. carpio*.

Metals	12 h	24 h	48 h	72 h	96 h
Al (mg L ⁻¹)	0.153 ± 6 × 10 ⁻³ ,b,c,d,e	0.129 ± 6 × 10 ⁻³ ,a,c,d,e	0.137 ± 4 × 10 ⁻³ ,a,b,d,e	0.186 ± 9 × 10 ⁻³ ,a,b,c,e	0.0729 ± 8 × 10 ⁻⁴ ,a,b,c,d
Fe (mg L ⁻¹)	0.03 ± 2 × 10 ⁻⁴ b,c,d,e	0.01 ± 5 × 10 ⁻⁴ a,d,e	0.01 ± 5 × 10 ⁻⁴ a,d,e	0.02 ± 5 × 10 ⁻⁴ a,b,c,e	0.04 ± 1 × 10 ⁻⁴ ,a,b,c,d
Hg (mg L ⁻¹)	0.00014 ± 5 × 10 ⁻⁵ *	0.00017 ± 6 × 10 ⁻⁶	0.00024 ± 1 × 10 ⁻⁶ *	0.00017 ± 9 × 10 ⁻⁶ *	0.00015 ± 1 × 10 ⁻⁵ *
Mixture					
Al (mg L ⁻¹)	0.100 ± 7 × 10 ⁻³ b,c,d,e	0.0415 ± 3 × 10 ⁻⁴ a,c,d,e	0.0643 ± 7 × 10 ⁻⁴ a,b,d,e	0.0597 ± 3 × 10 ⁻⁴ a,b,c,e	0.0519 ± 3 × 10 ⁻⁴ a,b,c,d
Fe (mg L ⁻¹)	0.03 ± 4 × 10 ⁻⁵ b,c,d,e	0.01 ± 3 × 10 ⁻⁴ a,c,d,e	0.01 ± 3 × 10 ⁻⁴ a,b,d	0.02 ± 5 × 10 ⁻⁴ a,b,c	0.02 ± 3 × 10 ⁻⁴ a,b
Hg (mg L ⁻¹)	0.0003 ± 1 × 10 ⁻⁶ c,e	0.00014 ± 1 × 10 ⁻⁶ c,d,e	0.0008 ± 9 × 10 ⁻⁶ a,b	0.0005 ± 1 × 10 ⁻⁶ b	0.0007 ± 4 × 10 ⁻⁶ a,b

Values are the mean of three replicates ± SEM. Significantly different from: *mixture; ^a12 h; ^b24 h; ^c48 h; ^d72 h; ^e96 h. *p* < 0.05. Nominal concentration in water for Al (0.2 mg L⁻¹), Fe (0.3 mg L⁻¹) and Hg (0.001 mg L⁻¹).

3. Results and discussion

Anthropogenic activity has resulted in the presence of high concentrations of heavy metals in water bodies, soil and air (Andreu et al., 2016). In Mexico, the concentrations of Al, Fe and Hg detected in diverse water bodies exceed the maximum permissible limits in the official norm on water for human use and consumption (NOM-127-SSA1-1994).

Heavy metal distribution and transport in water depends on the presence of species of these metals in the water column as well as on their environmental availability (González-González et al., 2014). After entering the body, these contaminants are

transported in the blood to the different organs and tissues where they bioaccumulate since they are not metabolized. Table 1 shows the concentration of Al (0.0729–0.186 mg L⁻¹), Fe (0.01–0.04 mg L⁻¹), Hg (0.00014–0.00024 mg L⁻¹) and the mixture [Al (0.0415–0.1 mg L⁻¹); Fe (0.01–0.03 mg L⁻¹); Hg (0.00014–0.0008 mg L⁻¹)] in blood of *C. carpio* exposed for 12, 24, 48, 72 and 96 h. In mixtures of contaminants with diverse specific modes of action, may occur interactions and the changes are basically modifications in the concentration of the contaminants. Heavy metals can alter physiologic and biochemical parameters in blood and tissue of fish (Vinodhini and Narayanan, 2009). Damage to DNA has been proposed as a useful biomarker for evaluating the

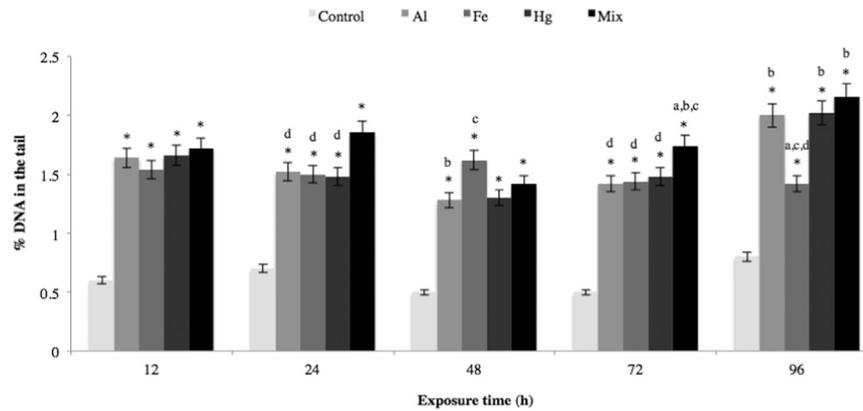


Fig. 1. Comet assay determination of DNA damage in blood of *C. carpio* exposed to Al, Fe, Hg and mixture thereof for 12, 24, 48, 72 and 96 h. Values are the mean of three replicates \pm SEM. Significantly different from: *control group; ^aAl; ^bFe; ^cHg; ^dMix. (Bonferroni *post hoc*, $p < 0.05$).

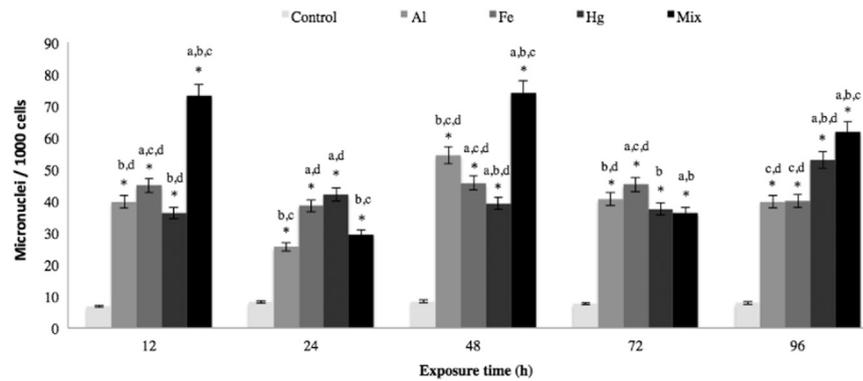


Fig. 2. Frequency of micronuclei in blood of *C. carpio* exposed to Al, Fe, Hg and mixture thereof for 12, 24, 48, 72 and 96 h. Values are the mean of three replicates \pm SEM. Significantly different from: *control group; ^aAl; ^bFe; ^cHg; ^dMix. (Bonferroni *post hoc*, $p < 0.05$).

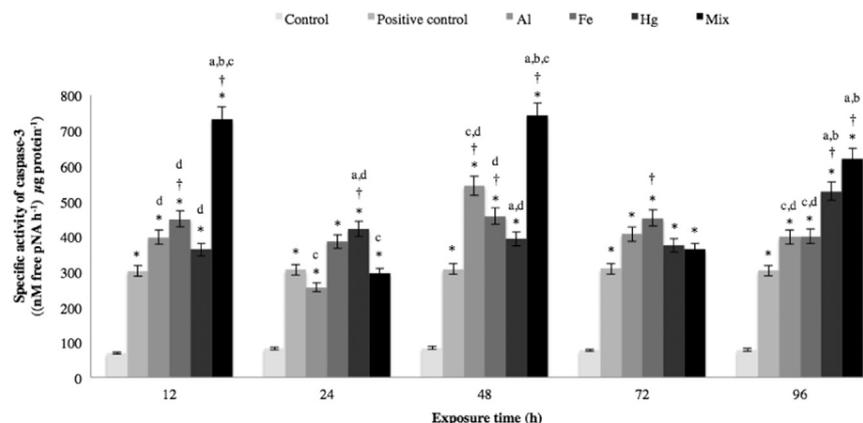


Fig. 3. Specific activity of caspase-3 in blood of *C. carpio* exposed to Al, Fe, Hg and mixture thereof for 12, 24, 48, 72 and 96 h. Values are the mean of three replicates \pm SEM. Significantly different from: *control group; †positive control; ^aAl; ^bFe; ^cHg; ^dMix. (Bonferroni *post hoc*, $p < 0.05$).

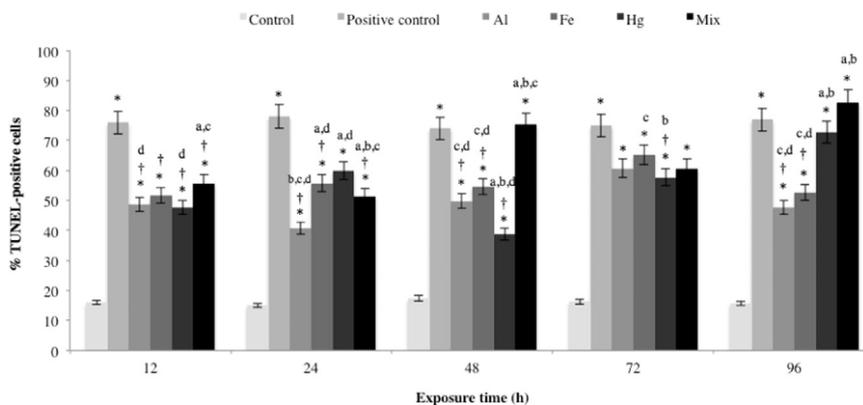


Fig. 4. TUNEL assay determination of the percentage of apoptotic cells in blood of *C. carpio* exposed to Al, Fe, Hg and mixture thereof for 12, 24, 48, 72 and 96 h. Values are the mean of three replicates ± SEM. Significantly different from: *control group; †positive control; ^aAl; ^bFe; ^cHg; ^dMix. (Bonferroni post hoc, *p* < 0.05).

Table 2
Pearson's correlation analysis of metal concentrations (Al, Fe, Hg and mixture thereof) with biomarkers of geno- and cytotoxicity in blood of *C. carpio*.

Biomarkers	Time (h)	Metals					
		Al	Fe	Hg	Al (mixture)	Fe (mixture)	Hg (mixture)
Comet assay	12	-0.671	0.468	-0.823	-0.209	0.395	-0.189
	24	0.142	0.645	0.492	0.306	0.792	0.753
	48	-0.217	0.873	-0.698	-0.343	0.052	0.462
	72	0.506	-0.753	-0.165	-0.594	0.431	-0.508
	96	0.274	-0.218	0.099	0.509	0.662	0.385
Micronucleus test	12	0.165	-0.308	-0.412	0.829	0.408	-0.625
	24	0.044	-0.327	-0.025	-0.314	-0.836	-0.793
	48	-0.946	-0.560	0.020	0.153	0.209	-0.229
	72	-0.932	0.827	-0.078	-0.319	0.039	-0.121
	96	-0.818	-0.456	0.829	-0.332	-0.754	-0.536
Caspase-3 activity	12	0.273	0.719	0.387	0.293	0.023	-0.137
	24	-0.461	0.859	0.625	-0.853	-0.354	-0.475
	48	-0.439	-0.019	-0.129	-0.679	-0.359	0.654
	72	-0.693	-0.309	-0.006	-0.656	0.536	-0.609
	96	0.012	0.168	-0.160	-0.884	-0.919	-0.156
TUNEL assay	12	-0.945	0.189	-0.826	-0.569	0.189	0.073
	24	1.000	-0.945	0.327	0.923	0.277	0.442
	48	-0.821	-0.693	-0.997	0.929	0.756	-0.987
	72	0.722	0.945	-0.881	-0.412	0.918	-0.839
	96	0.934	0.756	0.899	-0.487	-0.803	-0.727

Correlation coefficients > 0.5 are statistically significant (shown in bold).

genotoxicity of environmental contaminants in biomonitoring studies (Bombail et al., 2001), these contaminants include heavy metals. The alkaline comet assay is a reliable and sensitive early biomarker of genotoxicity in aquatic organisms since it detects DNA branch migration caused by strand breaks, alkali-labile sites and transient repair sites (Frenzilli et al., 2009; Kadam et al., 2013; Møller, 2006; Villarini et al., 2011).

As can be seen in Fig. 1, DNA damage increased significantly with each of the metals as well as the mixture in comparison to the control group (*p* < 0.05). This is probably explained by the fact that by products of normal mitochondrial metabolism and homeostasis include the buildup of potentially damaging levels of ROS (Zorov et al., 2014), in addition, MeHg also induces production of these species (Farina et al., 2013) that can induce intracellular Fe release (Flint et al., 1993; Halliwell and Aruoma, 1991). The superoxide anion (O₂•) releases Fe from ferritin (Bolann and Ulvik, 1990; Halliwell and Gutteridge, 1990) while hydrogen peroxide (H₂O₂) releases Fe from hemoproteins (Gutteridge, 1986; Halliwell and Gutteridge, 1990). Via the Fenton reaction, this metal produces the hydroxyl radical (OH•) which is highly reactive and binds to DNA bases, forming adducts (Valko et al., 2005). Aluminum itself can displace Fe in diverse biomolecules and increase intracellular

Fe levels, thus promoting the Fenton reaction (Amador et al., 2001; Dua and Gill, 2001; Yousef, 2004). This metal can also damage the mitochondrion and affect electron transport in the respiratory chain (Bondy and Campbell, 2001; García-Medina et al., 2010), increasing ROS production and thereby inducing genotoxicity. In mixture, Fe concentration did not increase, this may be explained by the fact that, free Fe can be buffered by a specific ferritin, which has an important physiological role as an antioxidant (Campanella et al., 2009; Santambrogio et al., 2007).

The main damage induced on DNA includes DNA base and sugar damage, protein-DNA and DNA-DNA crosslinks, single- and double-strand breaks, and abasic site formation (Bolognesi and Cirillo, 2014; Medeiros, 2008).

Micronuclei (MNi) are chromatin masses which form from chromosomal fragments or intact whole chromosomes and appear as small nuclei within the cytoplasm of cells in interphase (Cavas et al., 2005). Fig. 2 shows MNi frequencies in our study. Significant increases (*p* < 0.05) occurred at all exposure times and in all test systems (Al, Fe, Hg and the mixture). Such increases may be due to chromosomal breakage which yields acentric fragments, a resultant event of the clastogenic effects induced by some xenobiotics (Samanta and Dey, 2012), including metals.

These genotoxicity results are consistent with those of other authors. García-Medina et al. (2011, 2013) reported DNA damage and higher MNi frequencies in *C. carpio* exposed to Al (0.05, 120 and 239 mg L⁻¹), while Yadav and Trivedi (2009) concluded that exposure to Hg (0.081 mg L⁻¹) induces MNi formation in *Channa punctata*. Similarly, Bolognesi et al. (1999) found that Hg (32 µg L⁻¹) induced DNA damage and higher MNi frequencies in *Mytilus galloprovincialis*.

Cytotoxicity is defined as the pre-lethal changes and events which occur in cells prior to necrosis (Vasquez, 2012). Lesions to DNA and inefficient repair mechanisms are crucial in the unleashing of apoptosis (Roos and Kaina, 2006). Apoptosis is induced by the activation of intracellular proteases known as caspases (Boatright and Salvesen, 2003). Caspase-3 is an effector caspase in which both apoptosis pathways (intrinsic and extrinsic) converge. As shown in Fig. 3, significant increases ($p < 0.05$) with respect to the control group were observed in the specific activity of caspase-3 at all exposure times in all test systems. This may be explained by the fact that, since ROS are produced, a change in membrane potential takes place with a consequent decrease in ATP production and an increase in calcium levels (Bondy and Campbell, 2001; Flora et al., 2003; Verstraeten et al., 2008; Yamamoto et al., 2002). Damage to DNA and increased cytosolic calcium can initiate the intrinsic pathway. In this pathway, cytochrome c and other apoptogenic proteins in the cytosol are released by the mitochondrion via the opening of mitochondrial pores regulated by the Bcl-2 family; this release can be ROS induced. Free cytochrome c binds to Apaf1 (apoptotic protease activating factor 1) forming a multimeric complex that recruits and activates procaspase-9. This activated caspase-9 in turn activates the effector procaspase-3 (Reed and Pellecchia, 2005). Caspase-3 is responsible for proteolytic cleavage of a wide range of cellular targets, leading ultimately to cell death; increased activity of this caspase is seen in apoptotic processes (Gulbins et al., 2003; Taylor et al., 2008). The metals analyzed in our study are reported to affect calcium homeostasis. Calcium is transported to the cell nucleus where it activates nucleases which induce DNA strand breaks and elicit cytotoxicity (Meneghini, 1997; Valko et al., 2005).

Antibodies against specific neopeptides are used as markers for specific early events in apoptosis (O'Brien et al., 2001). In our study, anti-Fas mAb was used as the positive control since it is an apoptosis inducer (Watanabe et al., 2004). A significant increase ($p < 0.05$) in caspase-3 activity was observed in the positive control with respect to the control group at all exposure times, while similar increases with respect to the positive control were observed at 12 h with Fe and the mixture; at 24 h with Hg; at 48 h with Al, Fe and the mixture; at 72 h with Fe; and at 96 h with Hg and the mixture.

Apoptosis is characterized by cellular volume reduction, protein degradation, DNA fragmentation, chromatin condensation, blebbing of the cell surface and formation of apoptotic bodies (Thornberry, 1998).

One method used for *in situ* detection of DNA damage is the TUNEL assay, since it is able to detect cells in the initial stages of apoptosis and those in which morphologic changes, including apoptotic bodies, have already occurred (Gavrieli et al., 1992; Ky-larová et al., 2002). Significant increases ($p < 0.05$) in this type of damage with respect to the control group are seen in Fig. 4 at all exposure times in all test systems (Al, Fe, Hg and the mixture). This is consistent with our caspase-3 activity results, since the metals analyzed induce DNA damage.

DNase I is an endonuclease that induces DNA fragmentation (Nagata, 2000). Significant increases ($p < 0.05$) in the percentage of TUNEL-positive cells were observed in the positive control relative to the control group at all exposure times (Fig. 4). However, significant decreases with respect to the positive control were

found with Al and Fe (12, 24, 48 and 96 h), Hg (12, 48, 72 h) and the mixture (12 and 24 h).

Our cytotoxicity results are consistent with those of other authors, including García-Medina et al. (2011, 2013), who found an increase in the percentage of TUNEL-positive cells in *Cyprinus carpio* exposed to Al. Similarly, Voccia et al. (1994) described Hg-induced cytotoxicity in *Oncorhynchus mykiss* while Dayeh et al. (2005) and Rau et al. (2004) reported Fe-induced cytotoxicity in the latter species and *Poeciliopsis lucida*, respectively.

Table 2 shows the correlation of metal concentrations with biomarkers of geno- and cytotoxicity; as can be seen, correlations exist at certain exposure times.

Most ecotoxicological studies focus on the effects of exposure to single contaminants, but aquatic organisms in a contaminated environment are usually exposed to a mixture of contaminants (De Zwart and Posthuma, 2005). Toxicity of the metal mixture depends on metal concentration and specific composition as well as duration of exposure of the fish (Vosylienė et al., 2003).

4. Conclusions

The metals Al, Fe, Hg, and the mixture thereof induce cyto- and genotoxicity on blood of *C. carpio* exposed to the permissible concentrations for these metals in water for human use and consumption provided in the official Mexican norm NOM-127-SSA1-1994. The maximum permissible limits for these metals should be revised and updated since current values induce major damage on common carp and probably other aquatic species also, which, if consumed, could transfer these contaminants to humans.

Conflict of interest

The authors declare they have no actual or potential competing financial interests.

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