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Antagonistic bioaccumulation of waterborne Cu(II) and Cd(II) in common carp (Cyprinus carpio) and effects on ion-homeostasis and defensive mechanisms

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Aquatic Toxicology

Antagonistic bioaccumulation of waterborne Cu(II) and Cd(II) in common carp (Cyprinus carpio) and effects on ion-homeostasis and defensive mechanisms.

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Abstract:	In the aquatic environment, metals are present as mixtures, therefore studies on mixture toxicity are crucial to thoroughly understand their toxic effects on aquatic organisms. Common carp were used to assess the effects of short-term Cu(II) and Cd(II) mixtures, using a fixed concentration of one of the metals, representing 25 % of its individual 96h-LC 50 (concentration lethal for 50 % of the population) combined with a variable concentration of the other metal corresponding to 10, 25 or 50 % of its 96h-LC 50 , and vice versa. Our results showed a fast Cu and Cd bioaccumulation, with the percentage of increase in the order gill > liver > carcass. An inhibitory effect of Cu on Cd uptake was observed; higher Cu concentrations at fixed Cd levels resulted in a decreased accumulation of Cd. The presence of the two metal ions resulted in losses of total Na, K and Ca. Fish tried to compensate for the Na loss through the induction of the genes coding for Na + /K + -ATPase and H + -ATPase. Additionally, a counterintuitive induction of the gene encoding the high affinity copper transporter (CTR1) occurred, while a downregulation was expected to prevent further metal ion uptake. An induction of defensive mechanisms, both metal ion binding protein and antioxidant defences, was observed. Despite the metal accumulation and electrolyte loss, the low mortality suggest that common carp is able to cope with these metal levels, at least during a one-week exposure.	

Common carp were exposed to several binary mixtures of copper (Cu) and cadmium (Cd) at fixed and variable concentrations.

Copper and Cd are quickly accumulated in the gills, even though an antagonistic-like effect on cadmium uptake, caused by Cu was shown.

The metal mixtures led to a drop in sodium (Na) levels already after one day of exposure, whereas calcium (Ca) levels started to decrease only by the end of the experiment.

Induction of defensive mechanisms, such as metallothionein (MT) and glutathione reductase (GR), in order to cope with the increasing amount of metals.

Antagonistic bioaccumulation of waterborne Cu(II) and Cd(II) in common carp (*Cyprinus carpio*) and effects on ion-homeostasis and defensive mechanisms.

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Abstract

In the aquatic environment, metals are present as mixtures, therefore studies on mixture toxicity are crucial to thoroughly understand their toxic effects on aquatic organisms. Common carp were used to assess the effects of short-term Cu(II) and Cd(II) mixtures, using a fixed concentration of one of the metals, representing 25 % of its individual 96h-LC₅₀ (concentration lethal for 50 % of the population) combined with a variable concentration of the other metal corresponding to 10, 25 or 50 % of its 96h-LC₅₀, and vice versa. Our results showed a fast Cu and Cd bioaccumulation, with the percentage of increase in the order gill > liver > carcass. An inhibitory effect of Cu on Cd uptake was observed: higher Cu concentrations at fixed Cd levels resulted in a decreased accumulation of Cd. The presence of the two metal ions resulted in losses of total Na, K and Ca. Fish tried to compensate for the Na loss through the induction of the genes coding for Na⁺/K⁺-ATPase and H⁺-ATPase. Additionally, a counterintuitive induction of the gene encoding the high affinity copper transporter (CTR1) occurred, while a downregulation was expected to prevent further metal ion uptake. An induction of defensive mechanisms, both metal ion binding protein and anti-oxidant defences, was observed. Despite the metal accumulation and electrolyte loss, the low mortality suggest that common carp is able to cope with these metal levels, at least during a one-week exposure.

Keywords: mixture stress, metal pollution, defense mechanisms, ionoregulation, *Cyprinus carpio*.

1. Introduction

The aquatic environment is the main sink of pollutants produced by industries, sewage, agriculture and mining activities. Trace metals are persistent, non-degradable pollutants that can accumulate in the aquatic food web (Díaz-de-Alba et al. 2017). Two groups of metals can be distinguished, namely essential and non-essential. Metal ions belonging to the first group are important for many biological processes but can pose a risk for the organism at concentrations that are too low or too high. Essential metal ions such as those of zinc (Zn), copper (Cu) and iron (Fe) have a well-known role in animal cells. For example, Cu serves as a catalytic cofactor in several enzymes, thus it is essential in cellular respiration, connective tissue formation, melanin production and so on (Zhao et al. 2014). Copper ion uptake can occur through the high-affinity Cu transporter (CTR1) or by the divalent metal ion transporter (DMT1) (Mackenzie et al. 2004, Sevcikova et al. 2011). Furthermore, it can be taken up through a putative apical sodium (Na+)-channel, leading to competition with Na+ at the uptake site. Moreover, the uptake can be facilitated by exchangers, such as the sodium-hydrogen exchangers (NHEs), located in the branchial epithelial cells, through the extrusion of H⁺ (Grosell 2011, Niyogi et al. 2015). Elevated Cu ion levels can be dangerous for aquatic organisms, since they can lead to a disturbance in acid-base balance (Grosell 2011), alter Na+ homeostasis by decreasing the activity of the Na+/K+-ATPase and damage the cells (De Boeck et al. 2001). Moreover, Cu ions can promote oxidative stress (Bopp et al. 2008).

Non-essential metal species such as those of mercury (Hg), lead (Pb) and cadmium (Cd) have no known biological function in vertebrates and are toxic even at low concentrations (Danabas et al. 2018). Cd²⁺ is considered a threat for animals because it can alter ionoregulation, modulate protein structure and generate oxidative stress (Ferain et al. 2018). Toxic effects induced by Cd can result in hypocalcaemia, as a result of the competition between calcium (Ca²⁺) and Cd²⁺ (Cinier et al. 1997, McGeer et al. 2011). Moreover, Cd can cause changes in superoxide dismutase (SOD) and glutathione (GSH) activity in fish liver (Jia et al. 2011). Ultimately, chronic waterborne Cd exposure can induce immunosuppression in common carp leading to death (Zhang et al. 2017).

The antioxidant defence system plays a crucial role in preventing deleterious effects caused by reactive oxygen species (ROS). Enzymes such as SOD, and catalase (CAT) represent the first line of defence, converting superoxide (O_2^{-1}) into hydrogen peroxide (H_2O_2), and H_2O_2 into water (H₂O) and oxygen (O₂) (Pillet et al. 2019). Furthermore, GSH plays an important role in ROS defence and as a chelating agent for metal ions (Lange et al. 2002). Thus, the presence of enzymes such glutathione reductase (GR) and glutathione-S-transferase (GST) is needed for glutathione metabolism. Glutathione-S-transferase metabolizes lipid hydroperoxides (Dautremepuits et al. 2009) and mediates the conjugation reaction of GSH with electrophilic compounds, causing the depletion of GSH (Dickinson and Forman 2002). Glutathione reductase catalyses the reduction of glutathione disulfide (GSSG) in order to maintain a constant ratio of GSH/GSSG (Couto et al. 2016). In addition to the antioxidant system, metallothioneins (MT) play an important role in protecting the organism from metal toxicity. Metallothioneins are cysteine rich proteins, that play a significant role in essential metal ion homeostasis (e.g. Zn and Cu) and binding of non-essential metal ions (e.g. Cd) for sequestration (Atli and Canli 2008, Jakimska et al. 2011). Their levels and activity in tissues can be stimulated by both essential and non-essential metal ions (Hogstrand and Haux 1990, Wu et al. 1999).

Organisms in aquatic ecosystems are generally exposed to a mixture of metals that can be taken up via common uptake routes and interact with each other during uptake (Komjarova and Blust 2009). This interaction can stimulate or inhibit the uptake of particular compounds. For example, a non-competitive interaction can occur between Cu and silver (Ag) in which Ag can stimulate gill-Cu binding but not the other way around (Niyogi et al. 2015). A competitive interaction on the other hand can occur between Cd²+ and Zn²+, since they have a comparable electron configuration and they both have a high affinity for thiol groups (Brzóska and

Moniuszko-Jakoniuk 2001). Cadmium uptake can also be inhibited in presence of Cu as demonstrated in several organisms, such as zebrafish, rainbow trout and freshwater mussel (*Pyganodon grandis*) (Stewart 1999, Franklin et al. 2002, Kamunde and MacPhail 2011b, Komjarova and Bury 2014). Deleterious effects of mixtures of these metal ions have already been studied in different model species such as Mediterranean mussel (*Mytilus galloprovincialis*) (Benali et al. 2017), rainbow trout (*Oncorhynchus mykiss*) (Kamunde and MacPhail 2011a) and zebrafish (*Danio rerio*) (Komjarova and Bury 2014).

The aim of the present study was to investigate bioaccumulation, ionoregulation and responses of defensive mechanisms in common carp after a short-term exposure to waterborne binary mixtures of Cu(II) and Cd(II), using environmentally relevant concentrations. In addition, possible interactions between the two metal ions were investigated. Therefore, common carp were exposed to a series of sublethal mixtures of Cu (nominal concentrations used: 0.07, 0.19 and 0.38 µM) and Cd (nominal concentrations used: 0.026, 0.07, 0.13 µM) using environmentally relevant concentrations. In Flanders (Belgium) where this study was conducted, the water quality guideline for dissolved Cu in surface water is set to 0.11 µM, whereas for Cd the maximum values ranged between 0.004 to 0.013 µM depending on water hardness (Belgian Official Journal, 2015). However, in reality these limits are often exceeded. For instance, in Flanders the Flemish Environmental Agency (VMM) has measured concentration up to 2.05 µM for Cu and 1.06 µM for Cd (VMM 2016). Furthermore, a field study done in Flanders over 14 different locations reported dissolved metal concentrations up to 0.4 µM and 0.2 µM for Cu and Cd, respectively (Bervoets and Blust 2003). Therefore, the metal concentrations used in this study can be considered as environmentally relevant.

With this work, together with all the previous studies investigating and showing the complexity of metal mixture scenarios (Komjarova and Blust 2009, Niyogi et al. 2015, Brix et al. 2017, Pillet et al. 2019) we aim to provide new insights into understanding metal accumulation and toxicity in multi-metal exposure scenarios. We hypothesize that metal mixtures would remain sub-lethal, as our exposures were relatively short and maximum exposure concentrations were 25 % + 50 % of the 96h-LC50. Furthermore, we anticipate a quick metal bioaccumulation for both Cu and Cd, even though a reduced accumulation of Cd is expected in presence of Cu. Moreover, we expect that metal accumulation would trigger defensive mechanisms, such as MT and GR to mitigate possible deleterious effects. Regarding ion-homeostasis, in agreement with previous results from our lab, we expect a Na but not a Ca loss (Castaldo et al. 2020, Delahaut et al. 2020)

2. Material and methods

2.1. Experimental animals

Juvenile common carp were obtained from the fish hatchery at Wageningen University, the Netherlands. The fish were kept for several months at the University of Antwerp in a 1000 I aquarium filled with tap water before the experiments started. Fish were fed once a day *ad libitum* with commercial fish food (Hikari® Staple™, Klundert, The Netherlands). Temperature was kept at 20 °C, oxygen was provided with air stones and the photoperiod was set to 12 h light and 12 h dark (12L:12D). A biofilter was provided to maintain water quality. Three weeks before the start of each experiment, 200 fish were transferred and divided over two 200 I polyethylene tanks (100 fish per tank) filled with EPA medium-hard water (Weber 1991). Artificial medium hard water was reconstituted using four salts NaHCO₃ (1.14 mM); CaSO₄·2H₂O (0.35 mM); MgSO₄·7H₂O (0.5 mM) and KCI (0.05 mM) (VWR Chemicals). Oxygen was provided with air stones and the photoperiod was set to 12L:12D. Experimental methods complied with regulations of the Federation of European Laboratory Animal Science Associations (FELASA) and were approved by the local ethics committee of the University of Antwerp (Permit Number: 2015-94, Project 32252).

2.2. Experimental set-up

Two series of one-week waterborne exposures to binary mixtures of Cd and Cu were performed on common carp (length = 58.5 ± 6.8 mm; weight = 2.3 ± 0.9 g mean ± standard deviation (SD)). Besides control groups, treatments consisted of a fixed concentration of one of the metals at 25 % of the 96h-LC₅₀ previously calculated in our lab (Delahaut et al. 2020) combined with 10, 25 and 50 % of the 96h-LC₅₀ of the other metal (indicated as Cu_{fix}/Cd_{var} or Cd_{fix}/Cu_{var}). Exposure tanks consisting of five (plus one as backup in case of mortality) doublewalled polypropylene (PP) containers per treatment, each filled with 9 I of EPA medium-hard water and containing six fish, were set up in the climate chamber at 20°C. In each container oxygen was provided with an air stone. In order to avoid the accumulation of waste products, such as ammonia, 90 % of the water was changed daily. As indicated in Castaldo et al. (2020), aerated EPA medium-hard water was prepared 24 h in advance and kept at 20 °C. Conductivity (275 \pm 6.2 μ S/cm) and pH (8.2 \pm 0.2) were measured daily. Water samples were collected before (N = 120) and after (N = 140) the water change to check stability of the total metal concentrations. The nominal and measured metal concentrations in the water for both exposure series are shown in Table 1 and 2. Metal speciation was calculated with VMinteg (Supplementary information, SI-Table 1 and SI-Table 2).

Table 1: Metal concentrations (mean \pm SD) used for the different treatments in the exposure series Cu_{fix}/Cd_{var} . If the measured concentrations were below the minimum quantification limit of the instrument (BMQL), this was added together with the quantification limit.

	Nominal concentration	Measured concentration
Control	0 μM Cu	0.0031 ± 0.0031 μM Cu
	0 μM Cd	< 0.00089 μM (BMQL) Cd
Treatment Cu _{fix} /Cd ₁₀	25 % LC ₅₀ Cu (0.19 μM)	0.15 ± 0.020 μM Cu
	10 % LC ₅₀ Cd (0.026 μM)	0.025 ± 0.0027 μM Cd
Treatment Cu _{fix} /Cd ₂₅	25 % LC ₅₀ Cu (0.19 μM)	0.16 ± 0.020 μM Cu
	25 % LC ₅₀ Cd (0.065 μM)	0.062 ± 0.0062 μM Cd
Treatment Cu _{fix} /Cd ₅₀	25 % LC ₅₀ Cu (0.19 μM)	0.16 ± 0.024 μM Cu
	50 % LC ₅₀ Cd (0.13 μM)	0.12 ± 0.012 μM Cd

Table 2: Metal concentrations (mean \pm SD) used for the different treatments in the exposure series Cd_{fix}/Cu_{var} . If the measured concentrations were below the minimum quantification limit of the instrument (BMQL), this was added together with the quantification limit.

Nominal concentra	ation N	leasured conce	ntration

Control	0 μM Cu	0.0047 ± 0.0013 μM Cu
	0 μM Cd	< 0.00089 μM (BMQL) Cd
Treatment Cd _{fix} /Cu ₁₀	10 % LC ₅₀ Cu (0.077 μM)	0.068 ± 0.011 μM Cu
	25 % LC ₅₀ Cd (0.065 μM)	0.060 ± 0.0062 μM Cd
Treatment Cd _{fix} /Cu ₂₅	25 % LC ₅₀ Cu (0.19 μM)	0.16 ± 0.020 μM Cu
	25 % LC ₅₀ Cd (0.065 μM)	0.062 ± 0.0062 μM Cd
Treatment Cd _{fix} /Cu ₅₀	50 % LC ₅₀ Cu (0.38 μM)	0.34 ± 0.036 μM Cu
	25 % LC ₅₀ Cd (0.065 μM)	0.061 ± 0.0062 μM Cd

2.3. Metal bioaccumulation and electrolyte levels

At each sampling point (day one, three and seven) 10 fish per treatment (two from each container), were euthanized using an overdose of MS-222 (pH 7.0, ethyl 3-aminobenzoate methane-sulfonic acid, 400 mg/l, Acros Organics, Geel, Belgium). Muscle samples were cut near the caudal fin from individual fish and the following tissues were pooled from two fish in order to obtain enough tissue for ion and metal analysis: the first and the fourth gill arches from both the left and the right side as well as the brain and the liver. The samples were collected in pre-weighed Eppendorf tubes. The remaining carcasses from two fish were pooled and collected in pre-weighed 50 ml Falcon tubes to estimate the whole body accumulation. The samples were stored at -80 °C.

Metal and electrolyte content were determined in five samples of each tissue, at each sampling point. Reference material (SRM-2976, mussel tissue, National Institute of Standards and Technology, Gaithersburg, MD, USA), collected in pre-weighed Eppendorf tubes was included in the analysis as a quality control. The samples and the reference material were dried for 48 h, and the dry weight (dw) was recorded using a precision scale (Sartorius SE2, ultramicrobalance). Briefly, the digestion process (Blust et al. 1988, Reynders et al. 2006a) consisted of 12 h digestion at room temperature using 69 % concentrated HNO₃, followed by three microwave steps. Afterwards, H₂O₂ was added, to destroy the fat tissue, followed by another microwave digestion. Carcass samples, collected in pre-weighed 50 ml Falcon tubes were processed similarly. The samples, after an initial digestion step with 69 % HNO₃ at room temperature for 12 h, were digested using a hot block (Environmental Express, Charleston, SC, USA) for 30 min at 100 °C. At the end of the digestion process, all the samples were diluted using ultrapure Milli-Q (MQ), to reach a final acid volume concentration between 1 and 3 %. Metal and electrolyte content were determined respectively using a 7700x ICP-MS (Agilent Technologies, Santa Clara, CA, USA) and an iCAP 6300 Duo (Thermo Scientific, Waltham, MA, USA). Results obtained with ICP-MS and iCAP refer to the total element content (e.g total Cu, Na). Therefore, charges were only added when relevant for the discussion.

2.4. RNA extraction and real time PCR

The second and the third gill arch of individual fish and an aliquot of the pooled liver samples of two fish were used for RNA extraction and gene expression. Total RNA was extracted from samples (~ 20/30 mg) using Trizol (Invitrogen, Merelbeke, Belgium) following the manufacturer's instructions. Nano-Drop spectrophotometry (NanoDrop Technologies, Wilmington, DE) was used to determine RNA quantity and quality, whereas integrity was evaluated with a 1% agarose gel with ethidium bromide (500 μ g/ml). DNase treatment was performed using the commercial RNase free kit DNase I from Thermo Fisher Scientific (Waltham, MA, USA). Then 1 μ g of RNA was transcribed to cDNA according to RevertAid H minus First strand cDNA synthesis kit protocol (Thermo fisher, Fermentas, Cambridgeshire). According to the OD260/OD280 nm absorption ratio (higher than 1.8), four samples were selected and used for qPCR. Real-time PCR was performed using a Mx3000P QPCR System (Agilent Technologies, Belgium). The assay was performed in duplicate in a final reaction volume of 20 μ l containing 10 μ l of Brilliant III Ultra-Fast QPCR Master Mix (Agilent), 500 nM of each primer (reverse and forward), 5.7 μ l of sterile water, 0.3 μ l of reference dye and 5 ng of cDNA. PCR amplification was carried out following the Brilliant III Ultra-Fast QPCR Master

Mix (Agilent) protocol for Agilent Mx3000P QPCR system. Oligonucleotides primers were taken from literature: elongation factor 1 α (eEF) (Sinha et al. 2012), β -actin (Wu et al. 2014); H⁺-ATPase (Sinha et al. 2016), catalase (CAT) (Wu et al. 2014), superoxide dismutase Cu-Zn (SOD) (Wu et al. 2014), glutathione reductase (GR) (Wu et al. 2014), glutathione S-transferase (GST) (Casatta et al. 2017), metallothionein (MT) (Reynders et al. 2006b), Na⁺/H⁺-exchanger (NHE-2) (Castaldo et al. 2020) and Na⁺/K⁺-ATPase (Castaldo et al. 2020). Primers for CTR1 were designed using NCBI resources Primer blast and synthesized as highly purified salt-free "OliGold" primers by Eurogentec (Eurogentec, Seraing, Belgium). Primer sequences, annealing temperature and primer efficiency are given in SI-Table 10. Primer efficiency was determined based on the slope of the standard curve, using a serial dilution of cDNA.

2.5. Statistical analysis

All data were presented as mean values \pm S.D. For the statistical analyses, normality of the data was tested with the Shapiro-Wilk test. Two-way analyses of variance (ANOVA) were performed on all accumulation and gene expression data, followed by Tukey test. Data were considered statistically significant when p-value < 0.05. All statistical tests were performed with GraphPad Prism version 8.02 for Windows (GraphPad Software, La Jolla California USA). According to Custer et al. (2000), for metal concentrations below the minimum quantification limit (BMQL), a value of MQL/2 was assigned. If > 50% of the observations were BMQL, no statistical tests were conducted. Data presented in the supplementary information, such as curve fitting the metal bioaccumulation (non-linear, Michaelis-Menten, one-phase decay) and sodium loss (two-phase decay) as a function of time and exposure concentration, were analysed using the same software.

3. Results

3.1. Dynamics of Cu and Cd bioaccumulation

3.1.1. Copper bioaccumulation

Copper accumulation in gill tissue showed a similar pattern for both the exposure scenarios (Fig. 1.1A and 1.2A). In general Cu is always higher in the treatment compared with the control. This increase seems relatively independent from waterborne Cu concentrations. However, at day seven fish in treatment Cd_{fix}/Cu_{25} accumulated more Cu in comparison with treatment Cd_{fix}/Cu_{10} . After seven days of exposure, a strong increase in Cu was observed in all the treatments, compared with the previous sampling day (ranging from \simeq 67 % to \simeq 87 %). In the liver (Fig. 1.1B and 1.2B), for both the experimental series, Cu content showed almost no differences between treatment and control; however at day seven the metal content increased in similar amounts for all the treatments compared with the controls (ranging from \simeq 60% to \simeq 135 %). In the remaining carcasses (Fig. 1.1C and 1.2C), by the end of the exposure period, a significant increase was observed in the treatments Cu_{fix}/Cd_{10-25} compared with the control. In the treatment Cd_{fix}/Cu_{50} , Cu increased significantly compared with the control from day three onwards. Moreover, at day seven, the metal content was higher in the treatments Cu_{fix}/Cd_{25} and Cd_{fix}/Cu_{25} compared to day one (Fig. 1.1C and 1.2C).

For both experimental series, a fast Cu accumulation was observed in the gills during the first day in all treatments. From day one onwards, Cu accumulation continued at a slightly lower pace, increasing linearly in time (See supplementary information, SI-Fig 1 A and B). During experimental series Cu_{fix}/Cd_{var} , there seems to be a steady Cu net accumulation rate ($\simeq 2.6$, 1.5 and 1.4 nmol g⁻¹ dw h⁻¹ for day one, three and seven respectively), which is not affected by Cd levels in the water (SI-Fig 1 A-1 and SI-Table 3). However, looking at experimental series Cd_{fix}/Cu_{var} , the accumulation appears to reach a limiting value at the highest Cu exposure concentration during the first days, which is less pronounced at day seven (SI-Fig 1 A-2 and SI-Table 4). By the end of the experiment, Cu accumulation in both the experimental series in terms of percentage of increase was in the order gills > liver > carcass, whereas in

terms of absolute values the order was liver > gills > carcass. In muscle and brain tissue, no statistically significant accumulation of Cu was observed for both exposures (SI-Table 5 and 6).

3.1.2. Cadmium bioaccumulation

Cadmium concentrations in the gills were nearly always significantly higher in the treatments compared to the control for the exposure series Cu_{fix}/Cd_{var} (Fig. 2.1A). Moreover, Cd accumulation showed a concentration dependent increasing trend linked to the waterborne metal concentrations. Furthermore, Cd in the gills significantly increased in all the treatments compared to the previous sampling day from day three onwards (Fig. 2.1A). Throughout exposure Cd_{fix}/Cu_{var} (Fig. 2.2A), Cd concentrations in the gills were significantly elevated in almost all the treatment groups compared to the control from day one onwards. A significantly lower Cd accumulation was observed with increasing waterborne Cu concentrations. Cadmium concentrations increased significantly in the treatments Cd_{fix}/Cu_{10-25} at day three compared with day one, while for treatment Cd_{fix}/Cu_{50} this only occurred after seven days (Fig. 2.2A).

For liver and carcass similar, but less pronounced Cd accumulation trends were observed in both exposure series (Fig 2.1B, 2.2B, 2.1C and 2.2C). For the exposure Cu_{fix}/Cd_{var} , significantly elevated Cd levels in the liver (Fig. 2.1B) compared to the control were observed in treatment Cu_{fix}/Cd_{50} from day one onwards and treatment Cu_{fix}/Cd_{25} from day three onwards. A significantly higher Cd accumulation in the liver of fish exposed to higher Cd concentrations was most evident at the end of the exposure. Cd content significantly increased compared to day one from day three onwards for treatment Cu_{fix}/Cd_{50} and at day seven for treatment Cu_{fix}/Cd_{25} . For exposure Cd_{fix}/Cu_{var} (Fig. 2.2B), significantly elevated liver Cd levels compared to the control were observed in treatment Cd_{fix}/Cu_{10} from day three onwards and Cd_{fix}/Cu_{25} after seven days. A significant difference in Cd accumulation among treatments was only observed after seven days. Cadmium levels significantly increased compared to day three in treatment Cd_{fix}/Cu_{10} and Cd_{fix}/Cu_{25} after seven days.

During exposure Cu_{fix}/Cd_{var} , the Cd concentration in the carcass (Fig. 2.1C) of treatment Cu_{fix}/Cd_{25} and Cu_{fix}/Cd_{50} showed a significant increase compared to the control from day three onwards. At the end of the exposure, Cd concentrations in the carcass were significantly elevated in treatment Cu_{fix}/Cd_{25} and Cu_{fix}/Cd_{50} compared to day one (Fig. 2.1C). Regarding exposure Cd_{fix}/Cu_{var} , Cd concentrations in the carcass were significantly elevated compared to the control in treatment Cd_{fix}/Cu_{10} from the first day onwards and from day three onwards in treatment Cd_{fix}/Cu_{25} and Cd_{fix}/Cu_{50} (Fig. 2.2C). Significantly higher Cd accumulation was observed in treatment Cd_{fix}/Cu_{10} compared to Cd_{fix}/Cu_{25} and Cd_{fix}/Cu_{50} from day one onwards (Fig. 2.2C). A significant increase in Cd content for all treatments was observed after three days compared to day one with a further increase after seven days for treatment Cd_{fix}/Cu_{10} (Fig. 2.2C).

Cadmium accumulation increased both through time and among the different exposure levels without reaching steady-state in the Cu_{fix}/Cd_{var} exposures (SI-Fig 2 A-1 and B-1). Although metal concentrations increased with time in fish exposed to Cd_{fix}/Cu_{var} (SI-Fig 2 A-2), a clear dose dependent inhibition of Cu on Cd levels can be observed, with a fast reduction in Cd uptake at the highest Cu exposure level. The accumulation rates for Cd in fish exposed to $Cu_{fix}/Cd_{10-25-50}$ were $\simeq 0.3$, 0.6 and 1/0.8 nmol g^{-1} dw h^{-1} , respectively for day one and three, whereas the accumulation rates dropped to 0.1, 0.3 and 0.4 nmol g^{-1} dw h^{-1} , respectively at the end of the experiment. In fish exposed to the $Cd_{fix}/Cu_{10-25-50}$ scenario, the accumulation rates expressed in nmol g^{-1} dw h^{-1} were 1.1, 0.6 and 0.4, respectively for day one and 0.9, 0.6

and 0.2, respectively at day three. A further decrease was observed at day seven were the accumulation rates were 0.5, 0.3 and 0.2 nmol g⁻¹ dw h⁻¹ respectively.

During the first day of exposure Cd_{fix}/Cu_{var} , Cd concentrations seem to level off into a plateau with almost equal inhibition at 25 % and 50 % LC_{50} for Cu (SI-Fig 2 B-2). However, this effect disappeared from day three onwards with a dose dependent inhibition observed for all Cu concentrations used (SI-Fig 2 B-2).

Similar to the results for Cu, the observed accumulation pattern in both exposures was gills > liver > carcass both in terms of relative and absolute values. In muscle and brain tissue, Cd levels stayed below the detection limit in both exposures (SI-Table 5 and 6).

3.2. Expression of MTs and antioxidant enzymes

A clear increased expression of the gene coding for metallothionein in the gills was observed in all treatments compared to the control from day one onwards in both exposure series (Fig 3.1A and 3.2A). In experimental series Cu_{fix}/Cd_{var} , a significantly higher MT gene expression was observed in treatment Cu_{fix}/Cd_{50} compared to Cu_{fix}/Cd_{25} after three days and Cu_{fix}/Cd_{10} after seven days (Fig. 3.1A). For exposure series Cd_{fix}/Cu_{var} no significant differences in MT gene expression were observed among treatments (Fig. 3.2A). The MT gene expression in the liver showed a statistically significant increase compared to the control after three days for all treatments in the Cu_{fix}/Cd_{var} series (Fig. 3.1B). A significant increase in liver MT gene expression compared to day one was observed for all treatments after three days with a subsequent significant decrease at day seven (Fig. 3.1B). For experimental series Cd_{fix}/Cu_{var} (Fig. 3.2B), a significantly higher transcription of the MT gene in the liver compared to the control was observed for treatment Cd_{fix}/Cu_{25} after three days and Cd_{fix}/Cu_{50} after seven days. Gene expression significantly increased compared to day one for treatment Cd_{fix}/Cu_{25} and Cd_{fix}/Cu_{50} after three days (Fig. 3.2B). After seven days a significant increase compared to day three was observed for treatment Cd_{fix}/Cu_{50} (Fig. 3.2B).

Relative GR mRNA abundance in the gills was almost doubled throughout experimental series Cu_{fix}/Cd_{var} in all treatments compared to the control (Fig. 4.1A). Moreover, no statistically significant differences in relative GR gene expression among treatments was observed. During exposure Cd_{fix}/Cu_{var} (Fig. 4.2A), a significantly increased relative GR mRNA abundance compared to the control was observed on nearly all sampling days. After seven days, relative GR mRNA abundance in treatment Cd_{fix}/Cu_{50} was significantly elevated compared to the control and both other treatments. Moreover, a significant increase compared to day three and one was also observed (Fig. 4.2A).

Regarding the expression of SOD in the gills, a significant increase in relative mRNA abundance compared to the control was observed from day three onwards in treatment Cu_{fix}/Cd_{10} and on day seven for treatment Cu_{fix}/Cd_{25} (Fig. 4.1B). During exposure Cd_{fix}/Cu_{var} a significant increase in relative SOD mRNA was observed after seven days in treatment Cd_{fix}/Cu_{25} and Cd_{fix}/Cu_{50} (Fig. 4.2B). Moreover, for both the exposure scenarios, no significant differences were observed among the treatments during the same sampling day. Furthermore, the gene expression of treatment Cd_{fix}/Cu_{50} at day seven is significantly higher compared with day one.

Concerning the expression of CAT in the liver, no significant differences were observed between the control and the treatment for fish exposed to Cu_{fix}/Cd_{var} (Fig. 4.1C). A significant decrease in relative mRNA abundance compared to the control was observed on day one and three for treatment Cd_{fix}/Cu_{50} ; however, by day seven the levels were similar to the control (Fig. 4.2C).

Finally, for GST expression in the liver, a significant decrease in relative mRNA abundance compared to the control was observed for treatment Cu_{fix}/Cd₁₀ after three days (Fig. 4.1D).

During exposure Cd_{fix}/Cu_{var} (Fig. 4.2D), a significant increase compared to the control was observed in treatment Cd_{fix}/Cu_{25} after one day and a significant decrease in treatment Cd_{fix}/Cu_{50} after three days. The GST mRNA abundance significantly decreased compared to the first sampling day, in treatment Cd_{fix}/Cu_{10} and Cd_{fix}/Cu_{25} after three days (Fig. 4.2D).

In gills, no statistically significant changes in relative GST and CAT mRNA abundance were observed between treatments and control during both exposure series (SI-Table 8 and 9). In liver, no statistically significant changes in relative SOD and GR mRNA abundance between treatments and control were observed during both exposure series (SI-Table 8 and 9).

3.3. Effects of metal exposure on ionoregulation

3.3.1. Sodium

The Na concentration in the gills showed a similar trend in the two experimental series, with a significant loss in the treatments from day one onwards (Fig. 5.1A and 5.2A). No significant difference in Na loss among treatments was observed. Changes in the electrolyte content for fish exposed to the same treatments started to become evident after day three, compared with the previous sampling day for the exposure Cu_{fix}/Cd_{var} (Fig. 5.1A), whereas for fish exposed to a variable concentration of Cu, the decrease was more accentuated at day seven compared with the previous days (Fig. 5.2A). In the liver a significant Na decrease compared to the control was observed from day one onwards for nearly all the treatments for fish exposed to a fixed amount of Cu (Fig. 5.1B). After seven days, a significantly lower Na concentration compared to day one was observed in treatment Cufix/Cd₁₀ and Cufix/Cd₂₅ (Fig. 5.1B). In the second experimental series, a significant Na loss compared to the control was observed in treatment Cd_{fix}/Cu₂₅ and Cd_{fix}/Cu₅₀ after seven days (Fig. 5.2B). A significant sodium decrease compared to day one was observed in treatment Cd_{fix}/Cu₅₀ at day seven (Fig. 5.2B). In the muscle (Fig. 5.1C and 5.2C) a comparable trend can be observed between the two experimental series, with a significant Na loss in the treatments compared with the control only by the end of the experiment. A significantly lower Na concentration compared to day one was only observed in fish from treatment Cu_{fix}/Cd₁₀ and Cu_{fix}/Cd₅₀ after seven days (Fig. 5.1C). In the brain a significant Na decrease was observed for all the treatments of exposure Cufix/Cd_{var} from day three onwards (Fig. 5.1D) which was also significant compared to day one (Fig. 5.1D). For experimental series Cd_{fix}/Cu_{var} a Na loss was observed only for fish exposed to the highest concentration of Cu at day seven (Fig. 5.2D). Finally, for the carcass (Fig. 5.1E and 5.2E), a significant decrease in Na levels was observed for all treatments from day three onwards in exposure series Cufix/Cdvar (Fig. 5.1E) and from day one onwards in exposure series Cd_{fix}/Cu_{var} (Fig. 5.2E).

3.3.2. Potassium

Potassium concentrations in the liver during experimental series Cu_{fix}/Cd_{var} significantly decreased for all treatments compared to the control from day three onwards (Fig. 6.1B). A significantly lower liver K concentration compared to day one was observed after three days in treatment Cu_{fix}/Cd_{10} and Cu_{fix}/Cd_{50} and after seven days in treatment Cu_{fix}/Cd_{25} . Furthermore, after seven days a further K decrease compared to day three was observed in treatment Cu_{fix}/Cd_{10} . During experiment Cd_{fix}/Cu_{var} (Fig. 6.2B), liver K concentrations were significantly lower in treatment Cd_{fix}/Cu_{25} and Cd_{fix}/Cu_{50} compared to the control from day three onwards. For treatment Cd_{fix}/Cu_{10} this decrease became significant at the end of the experiment. Potassium content in the liver was significantly lower compared to day one in treatment Cd_{fix}/Cu_{10} after three days and treatment Cd_{fix}/Cu_{50} after seven days (Fig. 6.2B).

In the brain, K content significantly decreased compared to the control group for all treatments from day one onwards during exposure Cu_{fix}/Cd_{var} (Fig. 6.1D). A significant decrease in K concentrations compared to day one was observed in treatment Cu_{fix}/Cd_{10} and Cu_{fix}/Cd_{25} after seven days (Fig. 6.1D). For exposure Cd_{fix}/Cu_{var} (Fig. 6.2D), K concentrations significantly decreased compared to the control in all treatments after seven days. Moreover, for treatment Cd_{fix}/Cu_{50} the decrease was already significant after one day (Fig. 6.2D). A significantly lower brain K concentration compared to day one was observed in treatment Cd_{fix}/Cu_{50} after three days and Cd_{fix}/Cu_{25} after seven days (6.2D).

For the remaining carcasses a significant K loss was only observed for all treatments compared to the control group at the end of exposure Cu_{fix}/Cd_{var} (Fig. 6.1E), while in the Cd_{fix}/Cu_{var} (Fig. 6.2E), a significant loss compared to the control was observed for treatment Cd_{fix}/Cu_{50} only after seven days.

Gills and muscle samples did not show any significant differences in K content compared to the control throughout both experiments (Fig. 6.1A, 6.2A, 6.1C and 6.2C).

3.3.3. Calcium and Magnesium

Calcium concentration in gills significantly decreased compared to the control after seven days in treatment Cu_{fix}/Cd_{50} (Fig. 7.A). For the liver, a significant decrease compared to the control was observed in treatment Cu_{fix}/Cd_{50} after three days, but not after seven days (Fig. 7.C). Regarding the Mg content in the gills, no significant changes were observed during exposure Cu_{fix}/Cd_{var} (Fig. 7.B). For the liver, a significant decrease for all treatment groups compared to the control was evident after seven days (Fig. 7.D). Moreover, the Mg content declined significantly compared to day one, in treatment Cu_{fix}/Cd_{10} and Cu_{fix}/Cd_{50} at day three and treatment Cu_{fix}/Cd_{25} at day seven (Fig. 7.D). A further liver Mg concentration decrease compared to day three was observed in treatment Cu_{fix}/Cd_{10} after seven days (Fig. 7.D). For all other tissues, no significant changes in the Ca or Mg content were observed during both experimental series (SI-Table 5 and 6).

3.3.4. Gene expression of ion channels in the gills

The expression of the CTR1 gene in the gills increased compared to the control in all treatments at day one and day seven during exposure Cu_{fix}/Cd_{var} (Fig. 8.1A). After three days, the expression significantly decreased compared to day one in treatment Cu_{fix}/Cd_{10} and Cu_{fix}/Cd_{25} (Fig. 8.1A). During exposure Cd_{fix}/Cu_{var} the CTR1 gene expression significantly increased in all treatments compared to the control for all sampling days, except for treatment Cd_{fix}/Cu_{50} on day one and treatment Cd_{fix}/Cu_{10} on day three (Fig. 8.2A). After seven days, a significantly higher CTR1 gene expression was observed in treatment Cd_{fix}/Cu_{25} and Cd_{fix}/Cu_{50} compared to treatment Cd_{fix}/Cu_{10} (Fig. 8.2A). Furthermore, after seven days, the expression significantly increased compared to day three in treatment Cd_{fix}/Cu_{25} and treatment Cd_{fix}/Cu_{50} (Fig. 8.2A).

Regarding H⁺-ATPase gene expression, a significant increase in the transcription was observed in all treatments at day one and seven during exposure Cu_{fix}/Cd_{var} (Fig. 8.1B). During exposure Cd_{fix}/Cu_{var}, a significant increase was observed for all treatments compared to the control after seven days (Fig. 8.2B). Moreover, for treatments Cd_{fix}/Cu₂₅ this increase was already significant from day one onwards and for treatment Cd_{fux}/Cu₂₅ it was also significant on day one (Fig. 8.2B). Considering Na⁺/K⁺-ATPase gene expression, a significant increase in mRNA abundance compared to the control was observed in treatment Cu_{fix}/Cd₁₀ after one day (Fig. 8.1C). After three days, the relative Na⁺/K⁺-ATPase mRNA abundance significantly decreased in all treatments compared to day one (Fig. 8.1C). During exposure Cd_{fix}/Cu_{var}, a significant increase in relative Na⁺/K⁺-ATPase mRNA abundance compared to the control was

observed after one day in all treatments (Fig. 8.2C). After three days, relative Na⁺/K⁺-ATPase mRNA abundance significantly decreased in treatment Cd_{fix}/Cu₁₀ and Cd_{fix}/Cu₂₅ compared to day one (Fig. 8.2C).

Regarding NHE-2, a significant decrease in relative mRNA abundance compared to the control was observed from day three onwards for all treatments during exposure Cu_{fix}/Cd_{var} (Fig. 8.1D). During exposure Cd_{fix}/Cu_{var} , a significant decrease of the relative NHE-2 mRNA abundance compared to the control was observed on day three for treatment Cd_{fix}/Cu_{25} and from day three onwards in treatment Cd_{fix}/Cu_{50} (Fig. 8.2D).

4. Discussion

We hypothesized that metal bioaccumulation and induction of protective mechanisms would occur. Results show that defensive mechanisms in common carp were able to respond adequately to minimize adverse effects and mortality. As expected throughout the experiment mortality was limited to treatment Cd_{fix}/Cu_{50} and only three fish (\simeq 8% of the population of one experimental series) died. The relatively low mortality could be explained by the short exposure period, the relatively tolerable Na loss and the activation of defensive mechanisms.

4.1. Dynamics of Cu and Cd bioaccumulation

4.1.1. Copper bioaccumulation

Not surprisingly, our data confirmed our initial hypothesis that metals would accumulate faster in gills and liver compared to other tissues. As expected, in fish exposed to a fixed amount of Cu, the content of this metal in the gill tissue, increased in comparable amounts for all the treatments (Cu_{fix}/Cd₁₀₋₂₅₋₅₀) at each sampling day, showing the importance of the exposure time on metal accumulation. After one week of exposure, for the Cu_{fix}/Cd_{var} series, the variable amount of Cd in the water showed relatively little effect on gill Cu accumulation (net Cu content by the end of the experiment was approximately 230, 240 and 213 nmol/g dw for treatment Cu_{fix}/Cd₁₀₋₂₅₋₅₀ respectively). For the experimental series Cd_{fix}/Cu_{var} a more marked Cu net accumulation, proportional to the metal exposure concentration in the water, was expected by the end of the experiment. Probably this discrepancy was due to the presence of Cd, which seemed to stimulate Cu accumulation at the lower Cu exposure concentrations.

When comparing the results obtained in the single exposure scenario using comparable metal concentrations, in which the net accumulated values after one week were approximately 112, 182 and 308 nmol/g dw for Cu, and 81, 137 and 267 nmol/g dw for Cd (Castaldo et al. 2020, Castaldo et al., under review) with results obtained in the binary mixture, it can be noticed that Cu content is slightly higher in the treatments Cd_{fix}/Cu_{10-25} (respectively $\simeq 44-32$ %) but that is not for the case of exposure to Cd_{fix}/Cu₅₀. Also in the Cu_{fix}/Cd_{var} exposure series, Cu accumulated to values slightly higher than those found for 25% of the 96 h-LC₅₀ in the single Cu exposures (~ 17-32%). This confirms that there was no systematic inhibiting effect of Cd on Cu accumulation, if anything, it was slightly stimulating (except at the highest Cu exposure). Results on Cu accumulation inhibition/stimulation by Cd are often inconsistent. A reduction of Cu uptake in presence of Cd was demonstrated in water flea (Daphnia magna) and in rainbow trout (Komjarova and Blust 2008, Niyogi et al. 2015). However, in another study, in which zebrafish were exposed to an increasing concentration of Cd (0.01, 0.05, 0.2 μM) plus a fixed concentration of Cu (0.02 µM), the presence of Cd did not alter Cu uptake (Komjarova and Blust 2009). Moreover, in rainbow trout a stimulation of Cu uptake in presence of Cd occurred (Brix et al. 2016). Therefore, the explanation of shared transporters and non-specific competition for binding sites seems reasonable (Niyogi et al. 2015). In fact, even though Cd and Cu are considered to be respectively the Ca2+ and Na+ antagonists (Grosell et al. 2002, Nivogi and Wood 2004), several studies provided evidence of shared uptake routes of these metals in fish gills via the ECaC, DMT1 and Zip-8 (Cooper et al. 2007, Alsop and Wood 2011, Komjarova and Bury 2014, Niyogi et al. 2015). Alternatively, effects might go unnoticed, since

after the first day with faster accumulation rates, the Cu accumulation rate seems to attain a steady-state value which is similar over all exposure conditions.

In gills, the fast Cu accumulation during the onset of the exposure was expected, considering that gills are in direct contact with the water, and the exposure medium is non-complexing (see SI-Table 1 and 2). Similarly, a previous study on rainbow trout and European eel (Anguilla anguilla), reported a rapid accumulation of Cu in the gills already after a few hours of exposure (Grosell et al. 1998, Kamunde et al. 2002). Moreover, such a fast accumulation is consistent with the high conditional equilibrium constant for metal ion binding sites on the gill surface (log $K_{\rm cond} = 7.4 - 7.8 \, (\text{dm}^3 \, \text{mol}^{-1})$, calculated at pH 6.2 and 7.9 with ionic strength of $\simeq 1.\text{e}\text{-}04$ and 3.20e-03) (Playle et al. 1993, Brix et al. 2016) and the capacity thereof. However, we have to consider that gills are only a temporary target organ for metal toxicity, as metal ions are subsequently transferred to the liver and kidney for the excretion via the hepatobiliary system (Grosell 2011, Kondera et al. 2014). In fish exposed to Cufix/Cdvar and Cdfix/Cuvar, a significant Cu accumulation in the liver was only observed at day seven. During experimental series Cd_{fix}/Cu_{var}, this accumulation appeared to be directly proportional to the external Cu concentration, even though differences among treatments were not statistically significant. Moreover, the pronounced increase in Cu content in both liver and gills after seven days could be related to the hepatobiliary excretion no longer being able to compensate for the increased metal bioaccumulation through the gills. Regarding Cu accumulation in the carcass, the transient increase reported for both the exposure scenarios seems to follow the pattern observed for liver with a slight delay in the accumulation, supporting the hypothesis that regulatory mechanism are struggling to keep up with the continuous Cu uptake via the gills. No significant accumulation of Cu in muscle tissue was observed in the present study. A lack of Cu accumulation in the muscle of common carp was also observed by De Boeck et al. (1997), suggesting that the metal accumulation in the muscle becomes significant when the storage capacity of the liver is exceeded (Laurén and McDonald 1987). In the brain no Cu accumulation was reported, this is in accordance with what observed by Shaw et al. (2012) and with the thought that Cu accumulation from metal salts in fish brain is slow (Handy 2003, Shaw et al. 2012).

Several mechanisms in vertebrates are known to play a role in Cu homeostasis, such as the CTR1 and the Cu-ATPase (Anni et al. 2019). The CTR1 has been proposed as a Cu+ transporter which is insensitive to external Na+ concentrations (Mackenzie et al. 2004, Craig et al. 2010, Komjarova and Bury 2014). According to Grosell and Wood (2002) the copper uptake pathway which is sensitive to external Na⁺ concentration dominates in environments with a Na⁺ deficiency, whereas the Na⁺ insensitive copper uptake pathway dominates when Na⁺ concentrations are above 200 μmol I⁻¹ (Grosell and Wood 2002). The regulation of this transporter is unclear and results are often contradictory (Boyle et al. 2011). The transcript level of CTR1 is downregulated in the intestine of sea bream in response to a high copper diet, whereas that is not the case during waterborne Cu exposure (concentration representing the 25 % 96 h LC₅₀) (Minghetti et al. 2008). Similar to what was found by Komjarova and Bury (2014) in zebrafish exposed to Cu, we observed an increased CTR1 gene expression in common carp. In Cufix/Cdvar series a significant increase in CTR1 mRNA abundance occurred at day one and seven, whereas in the Cdfix/Cuvar series the gene expression was almost continuously increased in all treatments when compared to the control. The observed increment, especially at day seven, seems to be dependent on exposure media Cu levels. Also in zebrafish gills exposed to 0.016 µM of Cu, an increase in CTR1 gene expression was reported (Leung et al. 2014). Similarly, in yeast the transcript of CTR-type transporters are regulated by Cu levels (Labbé et al. 1997), whereas this is not observed in mammals (Lee et al. 2000). According to our results, in addition to the influence of the Cu concentrations, one can assume that the observed increased expression could be related to changes in internal electrolyte concentrations. Moreover, in fasted fish, an increase in cortisol levels can occur (Vijayan and Moon 1992, Hashemi et al. 2008), playing a role in up-regulating CTR1 mRNA expression as suggested by Tellis et al. (2012). However, a downregulation of the Cu

transporter gene would be expected to slow down metal accumulation and prevent potential toxic effects.

4.1.2. Cadmium bioaccumulation

Unlike Cu, Cd is a xenobiotic, which does not fulfil any known metabolic role and is considered as a non-essential metal (Matsuo et al. 2005). In the gills a time and dose dependent increase can be observed. Furthermore, an antagonistic-like interaction between accumulation of the two metal ions was obvious, since higher water Cu concentrations resulted in lower levels of Cd accumulation (Cd content by the end of the experiment was approximately 79, 50 and 29 nmol/g dw for treatment $Cd_{fix}/Cu_{10-25-50}$ respectively).

Regarding Cd in presence of Cu, the accumulation was up to 4 times lower compared to the single exposure scenario. These results are also reflected in the decreasing accumulation rates for Cd with increasing Cu exposure concentrations. A similar inhibition compared to the single exposures was also observed in an earlier study, in which common carp were exposed to a ternary mixture of 10% of the 96 h-LC50 of Cu, Zn and Cd (Castaldo et al. 2020). Moreover, an antagonistic inhibition of Cd uptake in the presence of Cu was reported in several other fish species such as Nile tilapia (*Oreochromis niloticus*), rainbow trout and zebrafish exposed to a Cd/Cu mixture (Eroglu et al. 2005, Komjarova and Blust 2009, Brix et al. 2017). The presence of shared uptake routes for Cd²⁺ and Cu²⁺, and non-specific competition for binding sites in fish gills likely explain the antagonistic like effect of Cu on Cd uptake (Cooper et al. 2007, Alsop and Wood 2011, Komjarova and Bury 2014). In general, Cd accumulated fast in the gills (Vinodhini and Narayanan 2008) and considering the very low background levels, accumulation was significant from the beginning, with the exception of fish exposed to Cd_{fix}/Cu_{50} .

In contrast to Cd accumulation in the gills, accumulation in the liver was more differentiated across treatments: Cd increased in the liver from day one in the treatment Cu_{fix}/Cd_{50} , followed by the Cu_{fix}/Cd_{25} treatment from day three onwards. Similar to a previous study with common carp exposed to a ternary mixture (Castaldo et al. 2020), no differences were observed in fish exposed to the lowest Cd concentration. Probably, this is due to efficient excretion processes (faeces, mucosal sloughing and hepatobiliary excretion) (McGeer et al. 2011). In contrast to exposure Cu_{fix}/Cd_{var} , fish exposed to Cd_{fix}/Cu_{var} mainly showed a significant Cd accumulation in the liver at day seven. However, Cd accumulation in the liver occurred only for the treatments Cd_{fix}/Cu_{10-25} reflecting the accumulation pattern of the gills. Regarding the Cd accumulation in the carcasses, a similar pattern compared to gills and liver was reported for both experimental series. The metal concentration seems to follow a transient increase starting at day three for almost all the treatments. However, we have to take into account that no Cd accumulation was reported in the muscle and in the brain. Therefore, one can hypothesize that this limited Cd accumulation could be explained by metal adsorption to the skin.

The rapid and substantial Cd accumulation in the gills reflects their role as the primary uptake site of metal ions during waterborne exposures, and indicates the vulnerability of these tissues, (Benhamed et al. 2016). The rapid Cd accumulation, similar to Cu, is consistent with the very high affinity that Cd has for gill binding sites (conditional log *K* of 8.6 (dm³ mol⁻¹), calculated at pH 6.2 and ionic strength of 1.e-04) (Playle et al. 1993, Playle 2004) and the capacity thereof, together with the non-complexing nature of our exposure media. Considering the binding constants, one can assume that relatively more Cd, rather than Cu, should bind to the gill surface. However, our results showed a higher Cu-compared to Cd accumulation. Therefore, we can hypothesize that both metals entered the cell, but Cu displaced Cd from MTs due to a higher affinity for the protein (Vašák 1991). Thus, Cd will be subsequently flushed away into the kidney for excretion processes, whereas Cu will remain into the tissue bound to MTs. Moreover, we can also consider firstly, that an antagonistic-like effect, of Cu on Cd uptake due to the shared branchial uptake routes will occur (Alsop and Wood 2011, Komjarova and Bury 2014), secondly that at the same equitoxic concentrations, Cd levels were almost three times

lower compared to Cu, and thirdly that our fish were fasted. In fact a higher Cu accumulation was reported in fish exposed to a reduced food ratio (Hashemi et al. 2008). Therefore, we hypothesize that due to the absence of food, fish tried to compensate for the electrolyte losses, by enhancing the uptake of essential elements from the water, thereby promoting Cu uptake. Moreover, it has been suggested that Cu binding capacity (B_{max}) can vary reflecting changing requirements of this essential metal in growing juvenile fish (Brix et al. 2016).

No significant accumulation of Cd in muscle tissue was observed in the present study. This is in accordance with previous studies that reported a significant Cd accumulation only after several months of exposure (Cinier et al. 1999, Benhamed et al. 2016). Similar to the muscle, no Cd accumulation was observed in the brain. This was unexpected since the potential of Cd to accumulate in the brain of freshwater fish, such as silver catfish and zebrafish was pointed out in previous studies (Pretto et al. 2010, Al-sawafi et al. 2017). The lack of both Cu and Cd accumulation in these tissues indicates on the one hand that storage organs such as the liver were not saturated and on the other hand the ability of common carp to handle metal excesses. However, a longer exposure is needed to validate this thought.

4.2. Defensive mechanisms

Metallothioneins (MTs) are cysteine rich proteins which play an important role in metal ion homeostasis: their binding affinity for metal ions can reduce intracellular free metal ion concentrations thereby providing a protective role (Hamilton and Mehrle 1986. De Boeck et al. 2003). The metal binding strength of metallothioneins follows the order Hg²⁺ > Cu⁺ > Cd²⁺ $> Pb^{2+} > Zn^{2+} > Co^{2+}$ (Vašák 1991). In the present study, all the exposure conditions showed MT gene induction. In general, the MT mRNA expression was always increased in the gills, whereas that was not the case in the liver. In fact, the MT gene expression was delayed in the liver until day three. The increase in MT gene expression occurred concurrently with a significant metal accumulation for both the tissues. Several studies have pointed out the important role of MTs as metal scavengers and the relationship between metal accumulation and MT levels in different tissues (De Smet et al. 2001, De Boeck et al. 2003). The fast accumulation of Cu and Cd in the gills may have triggered the induction of the MT gene. Considering that background MT levels differ between the different tissues, with lower values in the gills compared to the liver (Hashemi et al. 2008), we can assume that the rapid increase in gene expression in the gills was a response of the fish to induce the synthesis of MTs in order to increase the protein levels. This fast response in common carp, as suggested by De Boeck et al. (2003), is clearly an advantage considering that extensive damage is usually caused by Cu toxicity during the first hours and days of exposure (McDonald and Wood 1993). Regarding the liver, the MT gene expression peak, observed at day three, followed by a decrease at day seven, could suggest that a temporary elevated protein synthesis was sufficient to cope with the metals, at least during this one week exposure. Common carp is known to guickly adapt to prolonged metal ion exposures, only increasing defensive mechanisms when needed (Martinez et al. 2004, Pillet et al. 2019).

As already mentioned, in addition to MTs, various antioxidant enzymes are present in cells to cope with deleterious effects caused by ROS (Wang et al. 2010). In the present work, we investigated the relative gene expression of SOD, CAT, GR and GST both in the liver and in the gills. The results obtained in the gills for both the exposure series are quite similar. Glutathione reductase is an enzyme involved in the renewal of GSH, and together with GST, it plays a key role in the GSSG/GSH balance (Dautremepuits et al. 2009). We observed an upregulation of the GR gene, similar to what was reported for sea bream exposed to Cu (Minghetti et al. 2008) and obscure pufferfish (*Takifugu obscurus*) exposed to Cd (Kim et al. 2010). Moreover, Kim et al. (2010), proposed that GR is one of the main antioxidants against Cd-induced oxidative stress in *obscure pufferfish*, since its transcript had the highest expression level in all examined tissues. It is known that common carp rely on GSH, which can bind metal ions, as a first line of defence (Eyckmans et al. 2011). Therefore, an increase in the GR gene expression is expected to reduce the level of oxidized glutathione. The obtained results, similar to what observed in a previous study (Castaldo et al. 2020),

suggested that a saturation of base levels of MT and GSH occurred, due to their ability to bind metal ions (Lange et al. 2002), and that a continuous production of the proteins is needed to handle the increasing amount of metals in the gills.

Regarding the SOD, metal ions such as Cd, Cu and Zn have been demonstrated to increase the SOD gene transcript (Sanchez et al. 2005, Cho et al. 2006). In our study, a general increase of the SOD gene occurred in all treatments compared to the control after seven days of exposure. Superoxide dismutase, together with CAT are considered as the first line of defence against ROS (Atli and Canli 2010, Weydert and Cullen 2010, Pillet et al. 2019). Therefore, we can assume that this transcription increase, which occurred concurrently with a drastic metal accumulation, is an attempt by the fish to boost its internal defences.

In the liver, no differences were observed in the expression of the GR and SOD genes. Considering that metal levels in the liver only started to increase after three or seven days, this could be an indication that metal levels remained below the threshold required to significantly increase ROS production. Regarding the CAT mRNA abundance, a decrease was observed during the first three days, followed by a subsequent recovery. This shows, on the one hand the susceptibility of common carp to elevated Cu and Cd concentrations and on the other hand the ability to quickly adapt to a stressful situation. In other studies, increases as well as decreases in CAT activity have been reported (Jia et al. 2011, Díaz-de-Alba et al. 2017, Pan et al. 2018). However, our trend is in accordance with Pillet et al. (2019) and indicates that common carp are able to adapt rapidly to metal ion exposure.

From the obtained results, we can conclude that gills, as expected, are the most vulnerable tissue. In fact the gills were the tissue with the highest percentage of metal accumulation, therefore a higher demand of defensive proteins is needed to counteract the high metal accumulation. Moreover, the gene expression responses of defensive mechanisms suggest fast adaptability of common carp towards oxidative stress.

4.3. Changes in electrolyte levels and organism responses

Metal ions are known to interfere with electrolyte ion homeostasis, due to competition at the uptake sites (Čelechovská et al. 2007, Niyogi et al. 2015). Among the electrolytes, Na+ is the major cation of the extracellular fluid (Sathya et al. 2012). In agreement with our initial hypothesis, decreased levels of Na were observed within the first day of exposure for both the experimental series. The net Na loss observed in the gill by the end of the experiment, in fish exposed to Cu_{fix}/Cd₁₀ and Cd_{fix}/Cu₁₀ was respectively around 159 and 117 μmol/g dw. This Na loss was slightly higher than in fish exposed to a ternary mixture of 10% 96 h LC₅₀ Cu, Zn and Cd, in which the Na loss was 78 µmol/g dw (Castaldo et al. 2020). Considering that Cd alone did not alter Na+ influx in rainbow trout (Birceanu et al. 2008), nor Na levels in common carp gills (Delahaut et al. 2019), this electrolyte decrease cannot be explained by additive effects. It is known that both Cu and Cd can inhibit the Na+/K+-ATPase activity through the binding at the Mg²⁺ binding sites and the -SH groups, respectively (Lionetto et al. 2000, Grosell et al. 2002, Handy et al. 2002). Therefore, it is likely that the inhibition of Na+/K+-ATPase activity caused by both metal ions, gradually lead to an increased Na+ content in the gill intracellular fluid (ICF), resulting in a reduced water-gill ICF electrochemical gradient for Na+ entry via apical channels (Birceanu et al. 2008).

As reported in previous studies, and in different species such as Nile tilapia and rainbow trout, gills are the most affected tissue in waterborne exposures (Grosell and Wood 2002, Mackenzie et al. 2004, Atli and Canli 2011, Niyogi et al. 2015). A drastic loss in total Na can result in death, however different species can tolerate different percentages of loss. For example the threshold for rainbow trout and yellow perch is set to 30 % and 40 % loss of whole body electrolyte content, respectively (Taylor et al. 2003), whereas gibel carp can tolerate losing up to the 45% of their whole body total Na content (De Boeck et al. 2010). In the present study the gill loss of total Na at day seven ranged between 33 and 45 %, whereas in carcass the Na decrease ranged between 24 and 36 %. However, Na levels in the gill of control animals

slightly increased from day one to seven (approximately 15%) which explains the high percentage (45 %) of Na loss encountered in fish gills exposed to Cu_{fix}/Cd_{var}.

Sodium can enter the gills through a putative Na $^+$ -channel powered by H $^+$ -ATPase, a Na $^+$ /Cl cotransporter and the NHEs (McCormick 2001, Kumai and Perry 2012). In order to understand mechanisms that act to maintain Na $^+$ -homeostasis, we analysed the expression of genes coding for Na $^+$ /K $^+$ -ATPase, H $^+$ -ATPase and the NHE-2 in the gills. An upregulation of the Na $^+$ /K $^+$ -ATPase gene expression was recorded in fish exposed to Cd_{fix}/Cu_{var} after one day for all the treatments compared to the control. In agreement with Boyle et al. (2011), we hypothesize that the increase in the Na $^+$ /K $^+$ -ATPase gene expression could be a compensatory mechanism to counteract the inhibition of the enzyme activity, caused by metals as already reported by several authors (De Boeck et al. 2001, Eyckmans et al. 2011) for the inhibition of the enzyme activity.

The active uptake of Na⁺ in freshwater fish is necessary for ionic homeostasis, and as already mentioned, Na+ can be taken up via a Na+/H+ exchanger or via a putative Na+-channel coupled with H⁺-ATPase (Wilson et al. 2000). Several NHEs have been identified and NHE-2 has been proposed as a candidate for the Na+-sensitive component of Cu+ uptake (Mackenzie et al. 2004, Craig et al. 2010, Komjarova and Bury 2014). In contrast with the observations reported by Komiarova and Bury (2014), our results showed a decreasing trend in NHE-2 mRNA abundance in both the exposure series. If NHE-2 can be proposed as a Cu uptake component, this reduction can be interpreted as an attempt by the fish to reduce the direct Cu uptake, and/or as a way to cut down the extrusion of protons in order to minimize the influx of Cu. Nevertheless, the latter assumption is unlikely as the H+-ATPase expression was upregulated. Alternatively, as described by Grosell (2011), it can be an indirect effect caused by the inhibition of the carbonic anhydrase, which leads to a depletion of the necessary substrate for the exchanger. Theoretically, the lower amount of substrate could lead to a reduced efficiency of the Na⁺/H⁺- exchanger but not of the H⁺-ATPase which is an active transporter. However, in both cases one of the outcomes is a drop of internal Na⁺ content, which the organisms try to compensate for by increasing the expression of the gene coding for H+-ATPase. Nonetheless, the Na⁺ loss in the gills has repercussions on the whole-body Na levels.

In contrast to Na $^+$, K $^+$ is the major cation of the intracellular environment (Sathya et al. 2012). Potassium homeostasis has been investigated in different species with contradictory results. A whole body K $^+$ reduction was reported in zebrafish larvae exposed to Cu (1.57 μ M) and Cd (10.7 μ M) (Alsop and Wood 2011). In contrast, higher levels of K were reported in major carp (*Catla catla*) and *Prochilodus scrofa* exposed to Cu and Cd (Cerqueira and Fernandes 2002, Hassan et al. 2018). Potassium is an important component of the Na $^+$ /K $^+$ -ATPase system, which is involved in maintaining the transepithelial membrane potential. It allows active transepithelial transport and regulates the cell volume (Skou and Esmann 1992). Moreover, the ion exchange mediated by Na $^+$ /K $^+$ -ATPase is crucial to prevent cell swelling (Lodish et al. 2000). The loss of total K, observed in liver, brain and carcass could be related to the ability of metal ions to inhibit the ion-transporting enzymes, leading to cell damage (Suresh et al. 1995, McGeer et al. 2000, Matsuo et al. 2005).

According to previous results obtained in our lab with common carp (Delahaut et al. 2019) we did not expect a loss in total Ca. Nonetheless, after one week of exposure in fish exposed to Cu_{fix}/Cd₅₀ an electrolyte decrease occurred in the gills, suggesting a synergistic-like effect between metal ions on Ca loss. For the liver, a significant Ca decrease was observed for treatment Cu_{fix}/Cd₅₀ after three days, but not after seven days. This trend is likely due to the considerable variation in the Ca concentration of the liver in fish from the control group throughout the exposure. A competition between Cd²⁺ and Ca²⁺ at the apical and basolateral Ca²⁺-channel has been reported resulting in hypocalcaemia (McGeer et al. 2000, Niyogi et al. 2015). The significant decrease observed only after one week indicates that both time and the exposure concentrations play a crucial role. Moreover, one can assume that the presence of Cu contributed to this loss. Considering that previous studies on zebrafish showed that Cu

decreases Ca²⁺ uptake and vice versa (Craig et al. 2010, Alsop and Wood 2011), we hypothesize that a competition between Ca²⁺ and Cu²⁺ at the uptake site occurred (Grosell 2011). Moreover, it has been suggested that Cu²⁺ epithelial transport may be via a Ca²⁺ pathway (Alsop and Wood 2011, Komjarova and Bury 2014).

Magnesium, together with Ca^{2+} , is another important cation found in bony tissue and is involved in the activation of numerous enzymes (Bijvelds et al. 1996). A decreased liver Mg content was reported in all the treatments of fish exposed to Cu_{fix}/Cd_{50} after seven days. Since Mg^{2+} is needed for energy metabolism and protein synthesis, a deficiency could occur after prolonged metal exposure due to the higher energy demands or due to competition with metal ions for the protein binding sites (Bijvelds et al. 1996, Matović et al. 2010, Pillet et al. 2019).

5. Conclusions

The main aim of the present study was to investigate the effects of a binomial waterborne metal mixture on bioaccumulation, defensive mechanisms, ion-homeostasis and survival in common carp. One of our initial hypothesis that was confirmed by our data was a quick metal bioaccumulation. Moreover, a dose dependent, non-mutual, antagonistic-like interaction between Cu and Cd uptake was observed. On the one hand Cu showed a marked inhibitory effect on Cd uptake, whereas on the other hand Cd showed a relatively small effect on Cu uptake. Another hypothesis tested in our experiment was the activation of defensive mechanisms, which were activated to a different extent in both gills and liver to protect the fish. In gills genes involved in defensive mechanisms against metal ion toxicity, such as MT and GR, were continuously upregulated compared to the control in order to mitigate possible deleterious effects. In the liver only transient increases in defence mechanisms were observed. As gills are continuously in contact with water, they are more vulnerable to the metal exposure compared to the liver, thus a non-stop production of defences is necessary. Regarding our initial hypothesis on electrolytes levels, a Na loss was confirmed in this study. The loss through the gills affected the whole body Na content, and an impairment of Na homeostasis affected K levels in several tissues as well. Nonetheless, the fish tried to cope with this situation by increasing the expression of the H⁺-ATPase and Na⁺/K⁺-ATPase genes. which are involved in Na+ homeostasis. In contrast with our expectations, a Ca decrease was reported in the gills, suggesting a synergic-like effect between the two metals on ionhomeostasis. A final hypothesis was that metal mixtures would remain sub-lethal to common carp which was confirmed since only few fish died in the different exposure scenarios. This low mortality rate can be linked to activation of the defence mechanisms present in the fish. In addition, the relatively limited Na loss and the short exposure period could have played a role in fish survival. In conclusion, we can affirm that the negligible mortality, together with the fish responses to a stressful situation, indicate the ability of common carp to cope with these levels of metal pollution, at least for one week. Further studies, in a longer exposure scenario, could provide new insights for unveiling the long-term biological effects of metal pollution on common carp.

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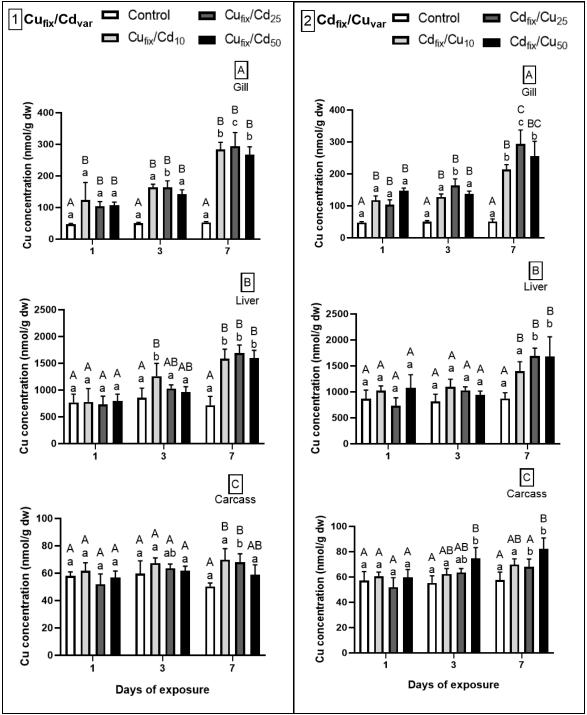


Fig. 1. Copper (Cu) concentration (nmol/g dry weight) in gills (A), liver (B) and carcass (C) of *Cyprinus carpio* exposed to Cu_{fix}/Cu_{var} (1) or Cd_{fix}/Cu_{var} (2) mixtures for 1, 3 and 7 days. Mean \pm SD, N=5. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p <0.05) among treatments within the same sampling day.

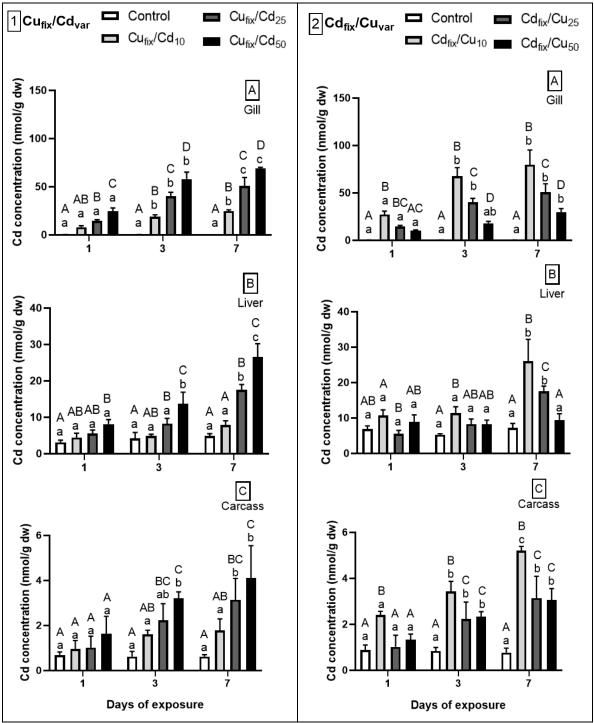


Fig. 2. Cadmium (Cd) concentration (nmol/g dry weight) in gills (A), liver (B) and carcass (C) of *Cyprinus carpio* exposed to Cu_{fix}/Cd_{var} (1) or Cd_{fix}/Cu_{var} (2) mixtures for 1, 3 and 7 days. Mean \pm SD, N=5. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p <0.05) among treatments within the same sampling day.

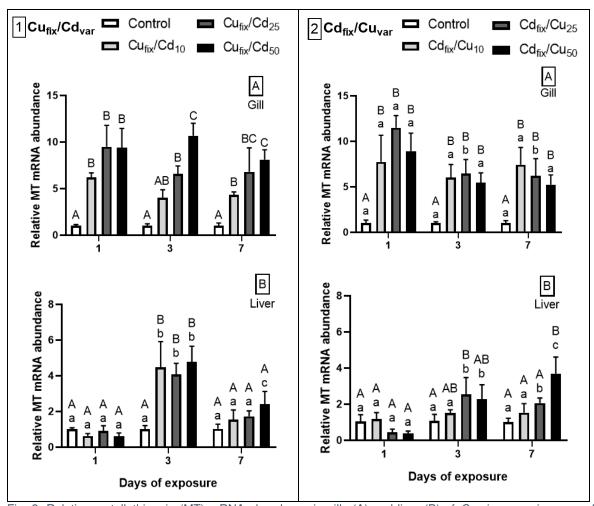


Fig. 3. Relative metallothionein (MT) mRNA abundance in gills (A) and liver (B) of *Cyprinus carpio* exposed to Cu_{fix}/Cd_{var} (1) or Cu_{var}/Cd_{fix} (2) mixtures for 1, 3 and 7 days. Mean \pm SD, N=4. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p < 0.05) among treatments within the same sampling day.

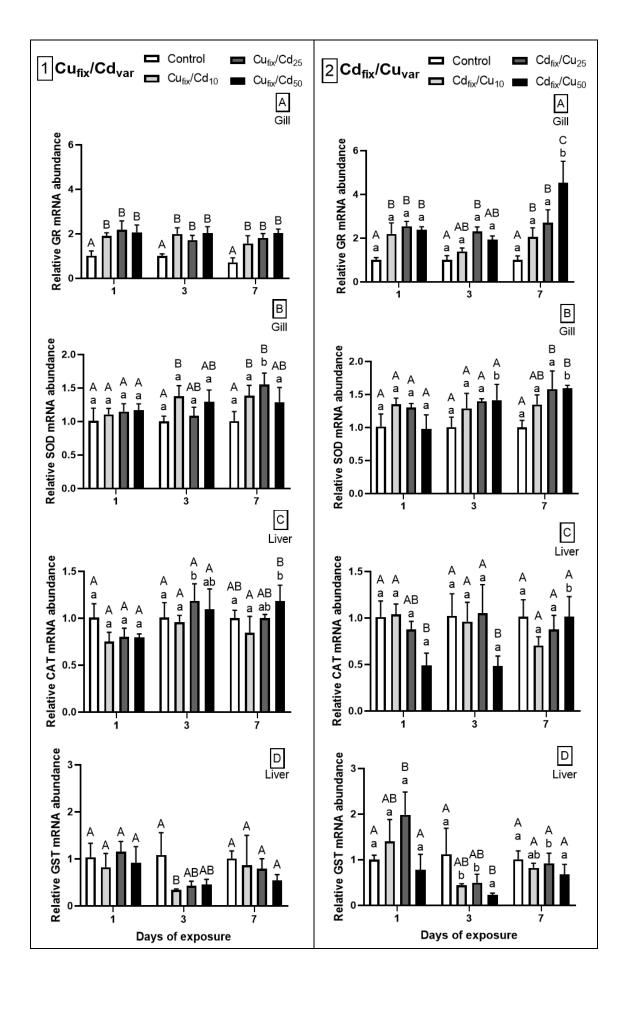


Fig. 4. Relative glutathione reductase (A), superoxide dismutase Cu-Zn (B), catalase (C) and glutathione S-transferase (D) mRNA abundance in gills and liver of *Cyprinus carpio* exposed to Cu_{fix}/Cd_{var} (1) or Cd_{fix}/Cu_{var} (2) mixtures for 1, 3 and 7 days. Mean \pm SD, N=4. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p <0.05) among treatments within the same sampling day.

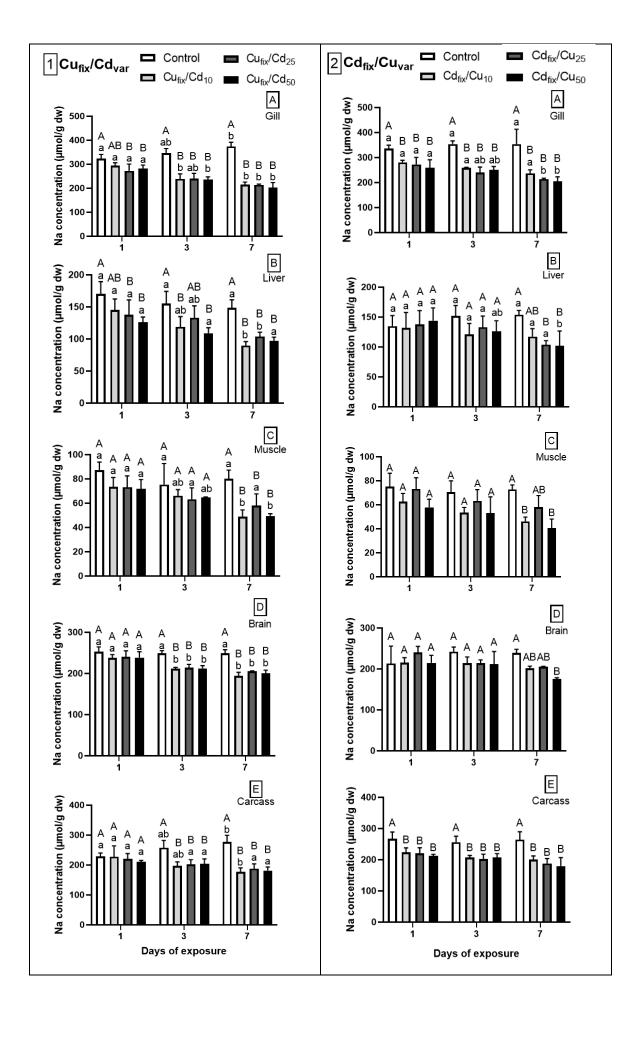


Fig. 5. Sodium (Na) concentration (μ mol/g dry weight) in gills (A), liver (B), muscle (C), brain (D) and carcass (E) of *Cyprinus carpio* exposed to Cu_{fix}/Cd_{var} (1) or Cd_{fix}/Cu_{var} (2) mixtures for 1, 3 and 7 days. Mean \pm SD, N=5. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p <0.05) among treatments within the same sampling day.

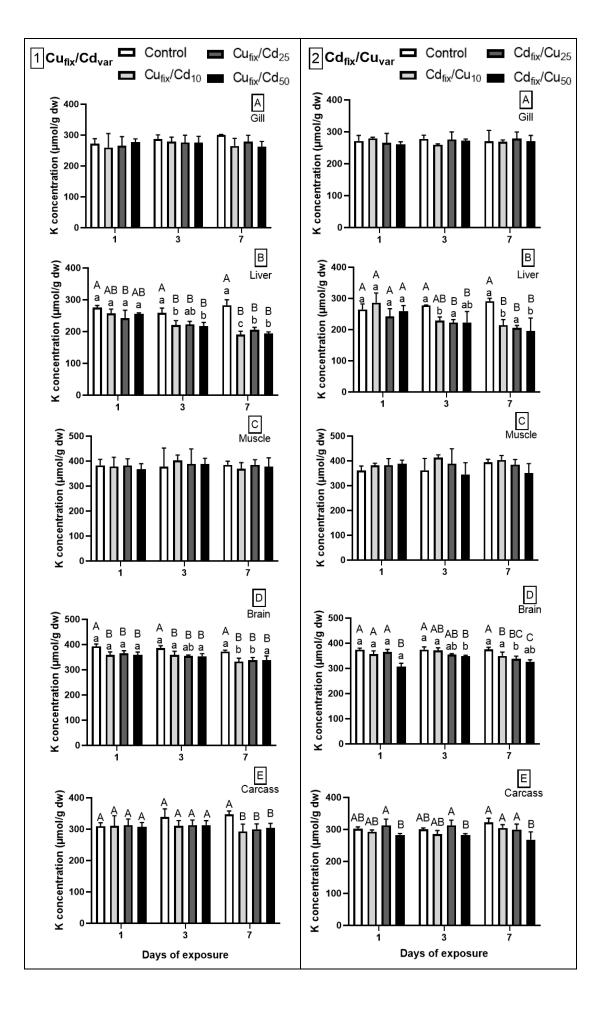


Fig. 6. Potassium (K) concentration (μ mol/g dry weight) in liver (A), brain (B) and carcass (C) of *Cyprinus carpio* exposed to Cu_{fix}/Cd_{var} (1) or Cd_{fix}/Cu_{var} (2) mixtures for 1, 3 and 7 days. Mean \pm SD, N=5. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p <0.05) among treatments within the same sampling day.

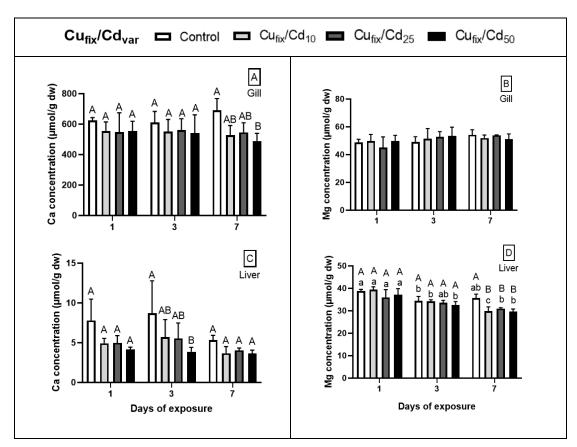


Fig. 7. Calcium and magnesium concentration (μ mol/g dry weight) in gills (A,B) and liver (C,D) of *Cyprinus carpio* exposed to Cu_{fix}/Cd_{var} mixtures for 1, 3 and 7 days. Mean \pm SD, N=5. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p < 0.05) among treatments within the same sampling day.

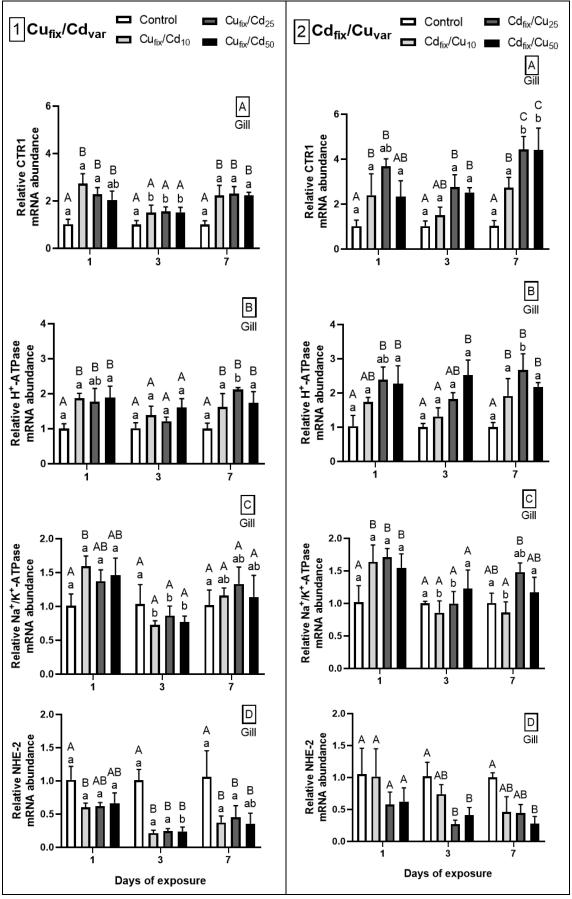


Fig. 8. Relative copper transporter 1 (A), H⁺-ATPase (B), Na⁺/K⁺-ATPase (C) and Na⁺/H⁺-exchanger (D) mRNA abundance in gills of *Cyprinus carpio* exposed to to Cu_{fix}/Cd_{var} (1) or Cd_{fix}/Cu_{var} (2) mixtures for 1, 3 and 7 days.

Mean \pm SD, N=4. Letters were only added when statistical differences occurred. Lower-case letters indicate significant differences (p < 0.05) of treatments among sampling days, capital letters indicate significant differences (p <0.05) among treatments within the same sampling day.

Conflict of Interest

Declaration of interests

☑ The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.	
☐The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:	

Giovanni Castaldo: Conceptualization, Methodology, Investigation, Formal analysis, Writing- Original draft preparation, Writing-Reviewing and Editing.: Flipkens Gunter: Conceptualization, Investigation, Formal analysis, Writing - Original draft preparation, Writing- Reviewing and Editing.: Pillet Marion: Investigation, Methodology.: Lieven Bervoets: Supervision, Funding acquisition.: Raewyn M. Town: Conceptualization, Validation.: Ronny Blust: Funding acquisition, Project administration.: Gudrun De Boeck: Conceptualization, Supervision, Funding acquisition, Project administration, Writing- Reviewing and Editing formal analysis.

Supplementary Material

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