



# Metal Accumulation and Ion Regulation in the Fish *Hyphessobrycon luetkenii* Living in a Site Chronically Contaminated by Copper: Insights from Translocation Experiments

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Received: 27 June 2021 / Accepted: 28 September 2021 / Published online: 18 October 2021  
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## Abstract

Fish living in the João Dias creek (southern Brazil) have to deal with trace-metal contamination in the long-term basis, as this aquatic environment has been historically impacted by copper mining activities. In order to survive in this harsh environment, the local biota had to develop adaptations related to pollution tolerance. The aim of this study was to test if biochemical mechanisms related to osmoregulation were among these adaptations, using translocation experiments. Water ionic and trace-metal compositions were measured in a nonmetal impacted site (NMIS) and in a metal impacted site (MIS) of this creek. Also, whole-body metal accumulation, ion concentration and branchial enzyme activity (Na,K-ATPase and carbonic anhydrase) were evaluated in *Hyphessobrycon luetkenii*. In both NMIS and MIS, fish were collected and immediately stored, kept caged or translocated from sites. The result shows that waterborne Cu was 3.4-fold higher at the MIS. Accordingly, animals that had contact with this site showed elevated whole-body Cu levels. Moreover, both translocated groups showed elevated Na,K-ATPase activity. Additionally, fish translocated from the NMIS to the MIS showed lower carbonic anhydrase activity. These findings indicate that *H. luetkenii* chronically or acutely exposed to naturally elevated waterborne Cu showed a rapid Cu bioaccumulation but was unable to readily excrete it. Moreover, classic Cu osmoregulatory toxicity related to Na,K-ATPase inhibition was not observed. Conversely, impacts in ammonia excretion related to carbonic anhydrase inhibition may have occurred.

Trace-metals are one of the most frequent pollutants released into natural areas (IPCC 2014; Zebral et al. 2019a). Some of these elements are essential components for life due to their redox ability, such as copper ions ( $\text{Cu}^{+2}$ ). Indeed, this metal is used to form cuproenzymes, molecules involved in several biological functions acting as enzymes, transporters and signaling transducers (Wood et al. 2011; Zhao et al. 2014). Cu is an essential element for life but can be toxic at elevated concentrations (Zebral et al. 2020). In several freshwater animals, including fish, this metal has been reported to inhibit

key enzymes such as Na/K-ATPase (Laurén and McDonald 1987) and carbonic anhydrase (Zimmer et al. 2012) leading to disfunction in ionic regulation (Grosell et al. 2002; Grosell 2011). Indeed, pollution is now considered as one of the greatest challenges that humans and other animals have to face (IPCC 2014; Zebral et al. 2019a).

The Na/K-ATPase is abundant in epithelial tissues, where this enzyme is used for the maintenance of intracellular ionic gradients used for absorption or excretion of compounds and for cellular homeostasis maintenance (McCormick et al. 2009). At the gills of freshwater fish, this enzyme can be majorly found in specialized cells responsible for ion uptake, the chloride cells (McCormick et al. 2009). Within this organ, Na/K-ATPase is responsible for the reduction of the intracellular concentration of  $\text{Na}^+$ , producing the gradient used for the entry of this ion through sodium channels (McCormick et al. 2009). On the other hand, the carbonic anhydrase is a ubiquitous enzyme responsible for catalyzing the reversible hydration of  $\text{CO}_2$  to form  $\text{H}^+$  and  $\text{HCO}_3^-$  (Zebral et al. 2019a). This reaction is important to

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many biochemical processes, such as acid–base regulation, ammonia excretion and ionic and osmotic regulation (Zebral et al. 2019a).

In order to adapt and survive in metal-contaminated environments, aquatic fauna has to develop physiological mechanisms associated with adjustments in metal uptake, storage and excretion, leading to reduced sensitivity to these elements (Depledge and Rainbow 1990; Whitehead et al. 2011; Romero et al. 2012; Uren Webster et al. 2013; Hamilton et al. 2016). Within this context, field experiments of reciprocal translocation are elegant strategies to elucidate such physiological mechanisms (Larsen et al. 2011). For example, the fish *Fundulus heteroclitus* locally adapted to different levels of chemical contamination showed divergent physiological patterns among populations. These patterns were related with the negative impacts caused by long-term exposure to contaminants such as polychlorinated biphenyls, dioxins, polycyclic aromatic hydrocarbons, trace-metals and pesticides (Whitehead et al. 2011).

The sub-basin of João Dias creek, a tributary of the Camaquã River located at Camaquã municipality (Caçapava do Sul city, southern Brazil), was directly impacted by Cu mining activity for more than a century (1870–1996) (Laybauer 1998) and even though the extraction has been discontinued for almost two decades, elevated levels of Cu can still be found in water and sediments from this area (Laybauer 1998; Ronchi and Lobato 2000; Bidone et al. 2001; Abril et al. 2018). To give an example, a recent study reported Cu waterborne levels corresponding to 8.5 µg/L in the region. Interestingly, the same study also showed that a non-impacted portion of this creek, located upstream the mining area, had Cu concentrations corresponding to 4.3 µg/L (Abril et al. 2018). In face of the presented facts, it is clear that the João Dias creek is an interesting environment to perform studies related to chronic exposure of aquatic organisms to naturally elevated levels of Cu. This kind of study is majorly needed, taking into consideration that most of the studies on the toxic effects of this metal were conducted under laboratory conditions and throughout acute exposure periods (Zebral et al. 2018). As a result of that, our knowledge related to Cu chronic effects on biochemical and physiological parameters is still limited, especially in wild fish populations (Whitehead et al. 2011; Uren Webster et al. 2013; Hamilton et al. 2016; Abril et al. 2018; Zebral et al. 2018; Anni et al. 2019a; Anni et al. 2019b; Zebral et al. 2019b; Zebral et al. 2021).

Despite the chronic contamination of João Dias creek, 36 fish species can still be found at the region, representing around 50% of all fish species reported for Camaquã river basin (Konrad and Paloski 2000). Among them, the lambari *Hypessobrycon luetkenii* is a small fish that can be found at Rio Grande do Sul water basins (southern Brazil), at coastal rivers of Rio de Janeiro (southeastern Brazil), and

at Paraguay and Uruguay rivers. At Rio Grande do Sul, this species can be found at the Patos Lagoon system and at the drainage basin of the following rivers: Camaquã, Uruguay, Negro, Paraguay, Tramandaí, Mampituba (Weiss 2013) and Guaíba (Konrad and Paloski 2000). *H. luetkenii* is known to form localized populations that usually swims in small groups of two to five individuals (Lima et al. 2008). These small (7 cm) animals are omnivorous freshwater fish, with a diet composed of small invertebrates, detritus and algae (Graciolli et al. 2003). Unfortunately, studies describing *H. luetkenii* life history are still scarce, but it is known that this fish species are fast growing animals, with parceled spawning of adhesive eggs. It is also interesting to comment that *H. luetkenii* is not migratory and completes its full reproductive cycle in localized environments (Giora et al. 2000; Lima et al. 2008).

The objective of this work was to search for physiological adaptations involved in osmoregulation in wild populations of the fish *H. luetkenii* living in the pollution portion of João Dias creek. Additionally, acute physiological adjustments were also evaluated, using translocation experiments. Physiological parameters evaluated were whole-body metal (Cd, Cu, Fe, Mn, Pb and Zn) and major ions (Mg, Na, K and Ca) concentrations and the activity of branchial enzymes involved in ionic regulation (Na/K-ATPase and carbonic anhydrase). To achieve our goal, fish living in a nonmetal impacted site (NMIS) and in a metal impacted site (MIS) of João Dias creek were collected for analysis. Also, fish collected at the NMIS were kept caged at the collection site or translocated and kept caged at the MIS and vice versa. Following 96 h of experiment, fish were collected for analysis. Our hypothesis was that animals chronically exposed to Cu contamination would show elevated Cu bioaccumulation and unimpacted levels of branchial enzymes and major ions, suggesting the presence of physiological adjustments at the populational level. On the other hand, it was hypothesized that animals translocated from the NMIS to the MIS would show clear signs of physiological impacts, such as inhibition in Na/K-ATPase and carbonic anhydrase and reduced levels of major ions.

## Materials and Methods

### Fish Translocation Experiment

The present work was performed in two areas of the João Dias creek located at Minas do Camaquã, southern Brazil. These areas were composed by a NMIS located at 7 km upstream the place where the mining activity occurred (30°53'47"S–53°25'28"W) and a historically MIS located within the mining area (30°52'55"S–53°27'11"W). An aerial photograph showing the collection points can be found

in Fig. 1, and photos of each site can be found in the supplementary material (Fig. 1a). At each site, 30 individuals divided among male and female fish (mean total body length = 4.32 cm) were caught using a fish trap and were divided into six groups. The first two groups were composed by animals collected at each site and immediately anesthetized with benzocaine, quickly rinsed in MilliQ water, euthanized by spine cord sectioning and stored in liquid nitrogen for the analysis of whole-body metal concentrations. Additionally, another group of animals were collected and euthanized as stated above but had their gills dissected and stored in liquid nitrogen for enzyme activity determination (Na/K-ATPase and carbonic anhydrase). The animals immediately collected at the NMIS and at the MIS were designated as control (C;  $n=5$  for Cu body burden and enzyme analysis) and polluted (P;  $n=5$  for Cu body burden and enzyme analysis), respectively.

The other experimental groups were composed by animals collected and kept caged at the NMIS (CCC;  $n=5$  for Cu body burden and enzyme analysis), collected and kept caged at the MIS (CCP;  $n=5$  for Cu body burden and enzyme analysis), collected at the NMIS and translocated to MIS (CP;  $n=5$  for Cu body burden and enzyme analysis) and animals collected at the MIS and translocated to the NMIS (PC;  $n=5$  for Cu body burden and enzyme analysis). Fish translocation was performed immediately following collection and animals were transported in buckets filled with local water continuously aerated with mobile air pumps. At the final destination, animals were acclimated to the new conditions by the addition of small volumes of water from the translocation experiment to the bucket. Following an hour, animals were transferred to the containment cages. Fish from CCC, CP, CCP and PC groups were kept caged for

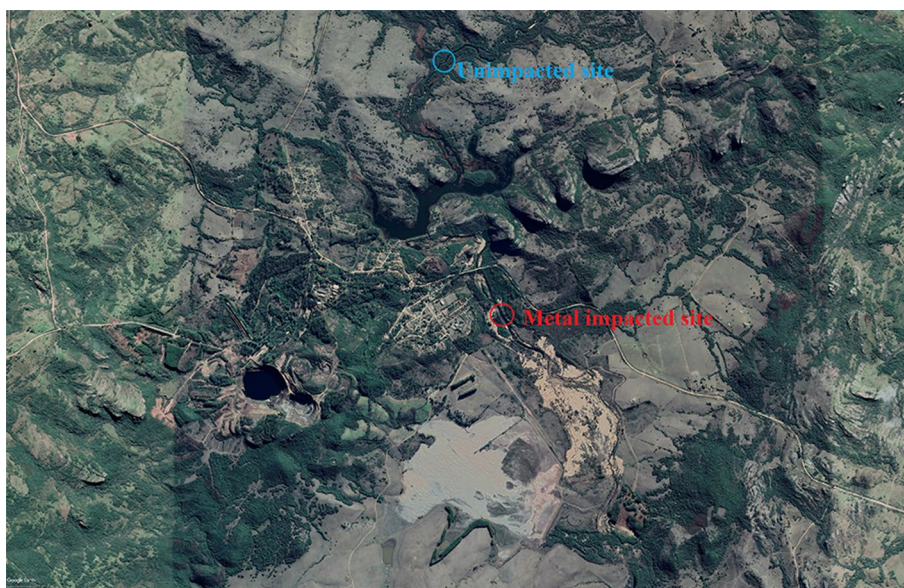
96 h. This is the most traditional experimental period used in acute ecotoxicological experiments, therefore, using it allows direct comparisons with previous and future studies. Additionally, longer experimental periods could implicate in the intensification of possible confounding factors related with caging the animals, such as different patterns of food availability and less water renew inside some of the cages resulting from mesh clogging with algae and detritus.

Following this period, animals were anesthetized with benzocaine, quickly rinsed in MilliQ water, humanely euthanized by spine cord sectioning and stored in liquid N<sub>2</sub> for determination of whole-body metal concentrations and enzyme analysis. Cages had a total volume of 5L and were surrounded by holes blocked with a fine mesh, in order to allow water flow. Cages were anchored with local stones using ropes. Considering the total volume of the cages and fish mean weight, it is possible to state that animals were stocked at <0.1 g fish/L. It is important to state that no fish mortality was observed, in any of the experimental groups.

### Environmental Parameters

Water quality parameters, including dissolved O<sub>2</sub>, pH and temperature, were daily monitored during fish collection and during translocation experiments. Also, water samples were collected, acidified (final concentration: 1%) with 65% HNO<sub>3</sub> (Suprapur) and kept at 4 °C until analysis. In these water samples, we determined the levels of total carbon concentrations (Total Organic Carbon analyzer; 5050A, Shimadzu, Japan), Mg, Na, K, Ca (flame photometer, model B262, Micronal, São Paulo, Brazil), Fe, Cd, Cu, Pb, Mn and Zn (HR-CS GF AAS; model Control-A 700; Analytik Jena, Germany). Standard curves were made by serially

**Fig. 1** Aerial photograph showing the sites where the freshwater fish *Hyphessobrycon luetkenii* was collected. The blue circle is indicating the precise location of the unimpacted site (NMIS) and the red circle is indicating the precise location of the metal impacted site (MIS). This aerial photograph was obtained using the software Google Earth Pro 7.3





dilutions of 1000 mg/L certified stock solutions (Multi-Element Standards Certipur<sup>®</sup>, Merck, Darmstadt, Germany). Methodological procedures were performed accordingly to Marques et al. (2019), additionally, LOD and LOQ are shown in supplementary material (Table A1). Metal recovery ranged from 92.6 to 102.8%. All reagents were of high-purity grade. Water used for preparing reagents and reference solutions was deionized and further purified using a Milli-Q system (Millipore Corp., Bedford, USA).

### Whole-Body Metal Concentration

Whole-body metal determination was performed according to Marques et al. (2019) with minor modifications. Fish were weighed before and after drying in an oven at 60 °C and were completely digested with HNO<sub>3</sub> (65%; Suprapur, Merck, Darmstadt, Germany). Mg, Na, K and Ca concentrations were measured using a flame photometer (São Paulo, Brazil, Micronal, model B262). Metal (Fe, Cu, Cd, Zn, Pb and Mn) concentrations were determined using a high-resolution atomic absorption spectrometer coupled with a graphite furnace (HR-CS GF AAS) (Germany, Analytik Jena, model Control-A 700). Spiked matrices and regular blank analysis were used as quality control and quality assurance for metal quantifications. Standard curves were made using standard solutions (Multi-Element Standards Certipur<sup>®</sup>, Merck, Darmstadt, Germany). Additionally, certified reference material (Fish protein DORM-3, National Research Council Canada, Ottawa, Canada) was analyzed to confirm extraction efficiency. Standard reference material was prepared as described for tissue samples and were used for the calculation of metal recovery. This evaluation was performed similarly to Sahuquillo et al. (1999) and Qu et al. (2014a, 2014b). Metal recovery ranged from 91.1 to 106.4%. All procedures were performed in triplicate.

### Enzyme Activities

Na/K-ATPase activity was measured in gill homogenates following procedures described by Bianchini and Wood (2003), with modifications. Tissue samples were homogenized in 0.5 mL of ice-cold buffer solution containing 150 mM sucrose, 50 mM imidazole, 10 mM ethylenediaminetetraacetic acid (EDTA) and 11.5 mM sodium deoxycholate. Homogenates were centrifuged at 4 °C, for 30 s, at 5,000 g and the supernatant was used as enzyme source. Two reaction mixtures were assayed: salt solution A (10.5 mM MgCl<sub>2</sub>, 100 mM NaCl, 30 mM KCl and 50 mM imidazole, pH 7.5) and salt solution B (50 mM imidazole, 10.5 mM MgCl<sub>2</sub>, 130 mM NaCl and 1 mM ouabain; pH 7.5). Reaction mixture A was made with 20 µL of sample homogenates and 200 µL of salt solution A. Reaction mixture B was made with 20 µL of sample homogenates

and 200 µL of salt solution B. Enzyme assays were run in duplicate at 20 °C (room temperature) during 10 min. For reaction cessation, 0.2 mL of trichloroacetic acid (50%) was added to reaction medium. Inorganic phosphate (Pi) concentrations in the reaction solution were assessed by commercial reagent kit (Fosfato, Doles, Goiânia, Brazil), based on the colorimetric method described by Fiske and Subbarow (1925).

The enzyme activity corresponded to the difference in Pi concentration produced in the two reaction mixtures (A and B). The idea behind this method is that the activity of Na/K-ATPase is directly related with elevated levels of Pi only on reaction mixture A, because the ouabain present in reaction mixture B is a strong inhibitor of this enzyme. Therefore, reaction mixture A yields Pi background levels present in homogenates together with the Pi produced by Na/K-ATPase activity during the assay. On the other hand, reaction mixture B only yields Pi background levels already present in the homogenates. Analyses were performed together with blank samples. Enzyme activity was normalized considering the protein content in sample homogenates, measured with Bradford reagent (Bio-Rad, Richmond, CA, USA). Total protein content was estimated using standard curves and blank analyses. Finally, enzyme activity was expressed as µmoles ADP/mg protein/h. Absorbance for Na/K-ATPase and Bradford analysis was made using a microplate reader (ELx-800, Biotek, Winooski, VT, USA).

Carbonic anhydrase activity was determined in gill homogenates using the delta pH method (Henry 1991) with modifications. This method is based in pH decrease following H<sup>+</sup> release upon the catalytic hydration of CO<sub>2</sub>. Reaction mixtures contained 15 mM sucrose, 225 mM mannitol, 10 mM phosphate and 10 mM Tris-Base (pH 8.5). Sample homogenates (10 µl) were added to 2 ml of reaction solution. The enzyme substrate (260 µl of MiliQ-water saturated with CO<sub>2</sub>) was then added to the mixture to start the reaction. Reaction mixture pH was measured every 5 s for up to 30 s. Blank reactions were prepared by replacing the sample homogenate with the buffer solution (10 µl) used for sample homogenization. Sample and blank measurements were taken simultaneously. The slope of pH values with time was estimated by a linear regression model. The catalyzed reaction ratio was considered as being the regression slope obtained for each individual sample homogenate. In turn, the non-catalyzed reaction ratio was considered as being the regression slope obtained in the blank measurement. Data were normalized based on the total protein content in sample homogenates, measured with Bradford reagent (Bio-Rad, Richmond, CA, USA). Enzyme activity was expressed as enzyme units/mg protein. Carbonic anhydrase analysis was made using a microplate reader (ELx-800, Biotek, Winooski, VT, USA).

## Data Presentation and Statistical Analysis

Data are shown as mean  $\pm$  standard error. Endpoints were analyzed using Analysis of Variance (One-way ANOVA) followed by the Tukey post hoc test. Parametric assumptions were graphically verified by residuals analysis (data normality) and the by the Cochran C test (homogeneity of variances). The significance level adopted was 95% ( $\alpha=0.05$ ). Analyses were performed using the software Statistica 12.0.

## Ethics and Legal Statements

All experimental and laboratorial procedures performed in the present work were previously approved by the university ethics committee (CEUA; protocol # 23,116.001365/2015-44) and by the Brazilian Ministry of the Environment (MMA; research license # 44,769-1).

## Results

### Environmental Parameters

Water pH (MIS:  $5.02 \pm 0.02$ ; NMIS:  $5.08 \pm 0.06$ ) and dissolved oxygen (NMIS:  $7.07 \pm 0.10$  mg O<sub>2</sub>/L; MIS:  $6.95 \pm 0.11$  mg O<sub>2</sub>/L) showed no significant differences when the two sites were compared. Additionally, no significant differences were observed when major cations concentrations (Ca, K, Na and Mg) were compared (Table 1). However, a significant difference between sites in water temperature (MIS:  $26.8 \pm 0.37$  °C; NMIS:  $24.6 \pm 0.39$  °C) was observed, as well as in the Cu concentration. As expected, the MIS showed higher concentrations (~twofold) in comparison to the NMIS (Table 1). Additionally, no other significant differences among trace-metals (Cd, Fe, Mn, Pb and Zn) were found when the MIS and the NMIS were compared (Table 1).

### Whole-Body Metal and Major Cation Content

As expected, whole-body Cu content in P, CCP, CP and PC fish was elevated in comparison to C and CCC fish (Fig. 2). Conversely, no differences were observed for *H.luetkenii* Cd, Pb (Fig. 2), Fe, Mn and Zn (Fig. 3) whole-body concentrations among all experimental groups. In accordance, no differences were observed for whole-body content of Na, K, Ca and Mg among the experimental groups evaluated (Fig. 4).

### Na/K-ATPase and Carbonic Anhydrase Activities

No significant differences were observed in Na, K-ATPase branchial activity among non-translocated fish (C, CCC, P

**Table 1** Carbon (TOC), major cations (Na, K and Ca) and trace-metals (Mg, Cd, Cu, Fe, Mn, Pb and Zn) concentrations in water samples from the metal impacted site (MIP) and the nonmetal impacted site (NMIS) at the João Dias creek (Caçapava do Sul city, Rio Grande do Sul state, southern Brazil)

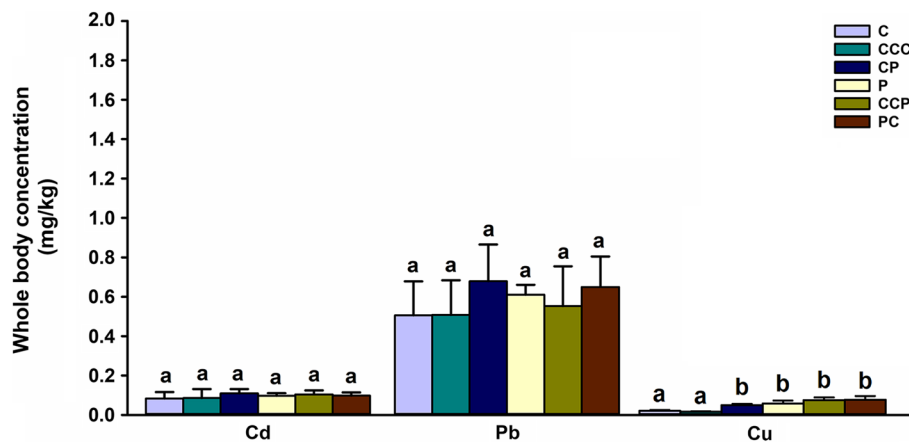
Parameter	MIS	NMIS
TOC (mg/L)	$4.02 \pm 0.69^a$	$9.09 \pm 0.51^b$
Na (mg/L)	$3.50 \pm 0.72^a$	$4.83 \pm 0.31^a$
K (mg/L)	$0.76 \pm 0.09^a$	$0.78 \pm 0.14^a$
Ca (mg/L)	$0.63 \pm 0.09^a$	$0.46 \pm 0.07^a$
Mg (mg/L)	$0.78 \pm 0.01^a$	$0.73 \pm 0.04^a$
Cd ( $\mu$ g/L)	$5.28 \pm 0.63^a$	$6.25 \pm 0.38^a$
Cu ( $\mu$ g/L)	$8.50 \pm 0.76^a$	$4.32 \pm 0.68^b$
Fe (mg/L)	$2.73 \pm 0.18^a$	$2.45 \pm 0.36^a$
Mn ( $\mu$ g/L)	$28.00 \pm 4.80^a$	$28.38 \pm 3.31^a$
Pb ( $\mu$ g/L)	$18.42 \pm 1.91^a$	$20.66 \pm 1.57^a$
Zn ( $\mu$ g/L)	$0.94 \pm 0.13^a$	$1.17 \pm 0.03^a$

Data are expressed as mean  $\pm$  standard error ( $n=10$ ). Different letters indicate significant difference among conditions (T-test;  $P < 0.05$ ; TOC = Total Organic Carbon)

and CCP). Conversely, this enzyme activity was significantly elevated in translocated animals (CP and PC) in comparison to non-translocated fish (Fig. 5). Despite that, no differences were observed between CP and PC fish (Fig. 5). In the case of carbonic anhydrase activity, the only group that showed significant differences was the CP fish, as an enzymatic inhibition was observed in these animals in comparison to all other groups (Fig. 6).

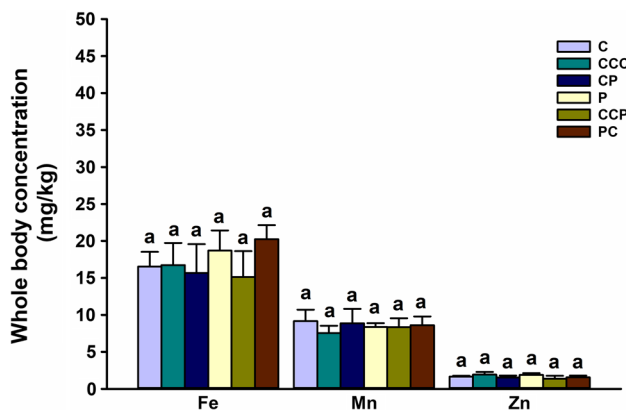
## Discussion

In agreement with the elevated levels of Cu found in the water samples, the whole-body content of this metal was significantly higher in P fish in comparison to C animals, indicating that chronic exposure to Cu under natural conditions resulted in the bioaccumulation of this metal. Interestingly, the same results have been reported by similar studies (Pyle et al. 2005; Couture and Pyle 2008; Uren Webster et al. 2013). Additionally, a significant increase in whole-body Cu burden was also observed in CP fish, indicating that *H.luetkenii*, like other freshwater fishes (Grosell and Wood 2002; Grosell 2011; Eyckmans et al. 2012; Zebra et al. 2019c), can readily accumulate Cu following an acute exposure (96 h) to increased levels of this metal. Indeed, Cu tissue uptake and accumulation by fish is known to be a fast (Grosell and Wood 2002; Grosell 2011; Eyckmans et al. 2012). Despite the aforementioned results, animals translocated from the MIS to the NMIS did not have any reduction in whole-body Cu concentration. This was an unexpected result, considering that one could have hypothesized that translocation



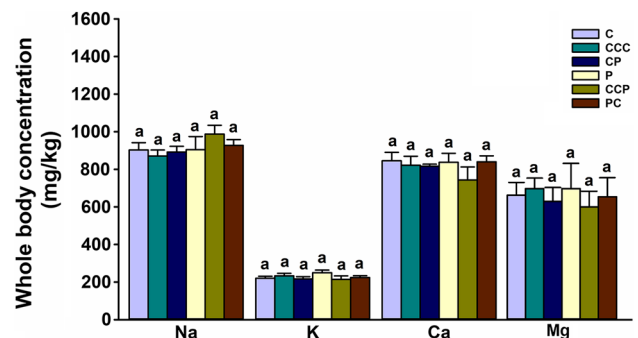
**Fig. 2** Whole-body Cd, Pb and Cu concentrations in the freshwater fish *H.luetkenii* collected at a nonmetal impacted site (C), kept caged at this site (CCC) or translocated to the metal impacted site (CP) for 96 h. Also, fish were collected at the metal impacted site (P), kept caged in this site (CCP) or translocated to the nonmetal impacted

site (PC fish) for 96 h. Data are expressed as mean  $\pm$  standard error ( $n=5$ ). Different letters indicate significant differences among fish groups for the same parameter analyzed (ANOVA followed by Tukey test;  $P < 0.05$ )



**Fig. 3** Whole-body Fe, Mn and Zn concentrations in the freshwater fish *H.luetkenii* collected at a nonmetal impacted site (C), kept caged at this site (CCC) or translocated to the metal impacted site (CP) for 96 h. Also, fish were collected at the metal impacted site (P), kept caged in this site (CCP) or translocated to the nonmetal impacted site (PC fish) for 96 h. Data are expressed as mean  $\pm$  standard error ( $n=5$ ). Different letters indicate significant differences among fish groups for the same parameter analyzed (ANOVA followed by Tukey test;  $P < 0.05$ )

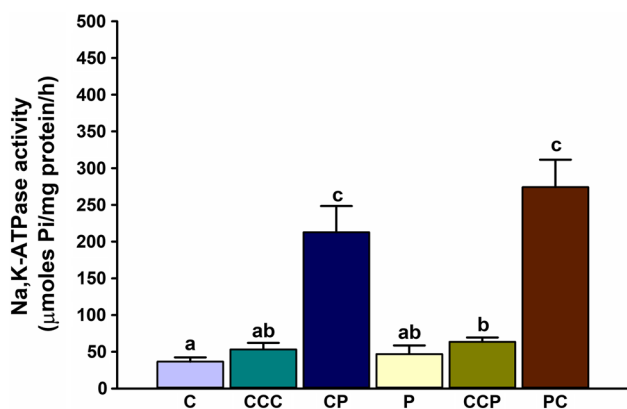
from a contaminated environment to a non-contaminated area would result in a clearance process. One possible explanation to this apparent contradiction is that 96 h was not enough time for an efficient metal depuration. Further supporting this idea, Subathra and Karuppasamy (2008) showed that the fish *Mystus vittatus* took 21 and 39 days to depurate the Cu bioaccumulated in gills and kidney, respectively, following exposure to 5.98 mg/L for 28 days. It was also possible to observe that none of the other trace-metals assessed in the water or in *H.luetkenii* whole-body were elevated in the MIS in comparison to the NMIS. This is a strong evidence



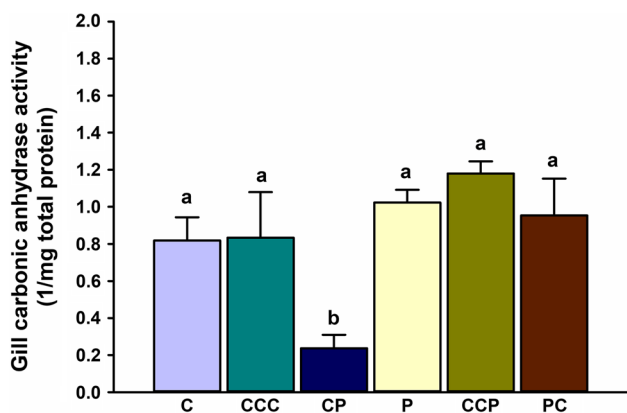
**Fig. 4** Whole-body major cations (Na, K, Ca and Mg) content in the freshwater fish *H.luetkenii* collected at the nonmetal impacted site (C), kept caged at this site (CCC) or translocated to the metal impacted site (CP) for 96 h. Also, fish were collected at the metal impacted site (P), kept caged in this site (CCP) or translocated to the nonmetal impacted site (PC fish) for 96 h. Data are expressed as mean  $\pm$  standard error ( $n=5$ ). Different letters indicate significant differences among fish groups for the same parameter analyzed

showing that the water bodies impacted by the mining activity at Minas do Camaquã District are contaminated with Cu specifically, and not with trace-metals in general.

Moving forward, it is well known that the main mechanism involved in Cu toxicity in freshwater fish is associated with inhibition of brachial Na/K-ATPase activity, resulting in ionic and osmoregulatory disturbances (Grosell and Wood 2002; Grosell 2011). With that in mind, we expected that the elevated levels of whole-body Cu observed in P, CP, PC and CCP would be accompanied by reductions in the tissue content of major cations (Na, K and Ca) due to Na/K-ATPase disruption. However, no significant differences were observed. For the case of P fish, this lack of disturbances



**Fig. 5** Gill Na,K-ATPase activity in the freshwater fish *H.luetkenii* collected at the nonmetal impacted site (C), kept caged at this site (CCC) or translocated to the metal impacted site (CP) for 96 h. Also, fish were collected at the metal impacted site (P), kept caged in this site (CCP) or translocated to the nonmetal impacted site (PC fish) for 96 h. Data are expressed as mean  $\pm$  standard error ( $n=5$ ). Different letters indicate significant differences among fish groups for the same parameter analyzed



**Fig. 6** Gill carbonic anhydrase activity in the freshwater fish *H.luetkenii* collected at the nonmetal impacted site (C), kept caged at this site (CCC) or translocated to the metal impacted site (CP) for 96 h. Also, fish were collected at the metal impacted site (P), kept caged in this site (CCP) or translocated to the nonmetal impacted site (PC fish) for 96 h. Data are expressed as mean  $\pm$  standard error ( $n=5$ ). Different letters indicate significant differences among fish groups for the same parameter analyzed

was also paralleled by unaltered branchial Na/K-ATPase and carbonic anhydrase activities. Altogether, these findings indicate that *H.luetkenii* wild individuals chronically exposed to Cu were able to deal with the excess of this metal in its tissues, showing no disturbances in ionic content. This result may indicate that the *H.luetkenii* population living in the MIS has developed local adaptations to deal with Cu contamination. These adaptations could be related with the production of Na/K-ATPase and carbonic anhydrase

isoforms that are less susceptible to Cu. Interestingly, as it can be observed in Fig. 1 (supplementary material), the populations of *H.luetkenii* living at the MIS and at the NMIS are separated by a dam catchment. This may result in reproductive isolation among them. Additionally, *H.luetkenii* are fast growing animals (Lima et al. 2008). Together, these two facts support the idea that animals living at the MIS have developed local adaptations at the populational level. Although extremely interesting, this point is still very hypothetical and further studies comparing genetic variations among populations living at the NMIS and in the MIS are still needed. There is no doubt that this is a prolific line of work and our future studies will be further assessing this matter.

For the case of Na/K-ATPase in translocated animals, an unexpected result was observed as both CP and PC fish had augmented levels of this enzyme in the gills. Despite that, one can hypothesize that the physiological basis for each of these outcomes are likely to be different. For example, Abril et al. (2018) showed in a companion paper that PC fish had a reduction in brachial Cu content, explaining the elevation in branchial Na/K-ATPase activity seen in the present study, as a reduction in the inhibition effect induced by this metal would also be expected. On the other hand, the induction of this enzyme in the gills of CP fish may indicate the activation of compensatory mechanisms to avoid osmoregulatory disturbances associated with higher Cu accumulation (Grosell 2011), such as the diffusive loss of Na due to oxidative damage to the gills (Craig et al. 2007; Wang et al. 2015; Ransberry et al. 2016). In this case, the suggested compensatory mechanisms could be related with the intensification of the ionic gradient produced by Na/K-ATPase in the gills, the force that drives Na absorption in this tissue (McCormick et al. 2009), counteracting any osmotic disturbances induced by Cu toxicity. It is interesting to comment that this was an unexpected result, considering that in most studies this enzyme was unaffected (Zimmer et al. 2012; Moysen et al. 2016; Canli et al. 2016) or was inhibited (Grosell and Wood 2002; Grosell 2011; Chowdhury et al. 2016) by Cu. It is important to remember that experimental animals were exposed to naturally elevated Cu levels together with a complex mixture of other environmental cues. In this context, the slightly elevated water temperature observed at the MIS could also help to understand this unexpected elevation in Na/K-ATPase of CP fish, as this parameter is known to impact this enzyme activity (Schwarzbaum et al. 1992; Yang et al. 2018; Monroe et al. 2019; Vargas-Chacoff et al. 2020). Therefore, the results of the present study may indicate that under a real-case scenario, the physiological mechanisms related to Cu-induced osmoregulatory toxicity may actually be more complex than previously expected. Certainly, this is an exciting field to be further assessed in future studies.



For the case of carbonic anhydrase, a significant reduction in this enzyme activity was observed in the gills of CP fish. In fact, Cu is known to be potent inhibitor of this enzyme (Grosell 2011; Zebraal et al. 2019a). The cytosolic form of carbonic anhydrase is responsible for the intracellular supply of H<sup>+</sup> to H-ATPase and Na/H-exchanger through hydration of intracellular CO<sub>2</sub>, facilitating Na uptake (Weihrauch et al. 2009; Wright and Wood 2009). Therefore, the inhibition of branchial carbonic anhydrase activity observed in CP fish could have led to disruption in Na<sup>+</sup> balance, but the whole-body content of this cation was unaltered in these animals. This apparent contradiction can be explained by the fact that, as discussed above, CP animals showed an elevation in Na/K-ATPase branchial activity, counteracting any possible ionoregulatory disturbances. On the other hand, the apical membrane-bound form of carbonic anhydrase is responsible, together with H-ATPase and Na/H-exchanger, to induce the acidification of the apical gill boundary layer, allowing the unprotonated form of ammonia to be excreted by facilitated diffusion (Weihrauch et al. 2009; Wright and Wood 2009), therefore, it is unwise to neglect a hypothetical impact in the excretion of nitrogenous compounds in CP animals. As a matter of fact, many studies have already demonstrated that Cu-induced toxicity and mortality in fish may be attributed to an inhibition in ammonia excretion through the gills (Grosell et al. 2003, Grosell et al., 2004a; Blanchard and Grosell 2006; Zimmer et al. 2012; Lim et al. 2015; Sinha et al. 2016), although the mechanism related to this toxic effect is not yet clear (Zimmer et al. 2012). In this regard, there is a strong hypothesis attributing this Cu-dependent disruption in ammonia clearance to an inhibition in the activity of carbonic anhydrase in fish gills (Grosell, 2011). The results obtained in the present study strengthens this hypothesis. Actually, as far as we know, this is the first evidence of a Cu-dependent inhibition in fish carbonic anhydrase activity in a field study. It is worth noting that any of the biological parameters assessed in the present study were different among C and CCC fish. Also, no significant differences were observed between P and CCP fish. These findings indicate that the significant effects observed in CP and PC fish can be attributed to animals' translocation and not to the fact that fish were kept caged during the experimental period (96 h).

## Conclusions

In conclusion, *H.luetkenii* chronically or acutely exposed to naturally elevated levels of Cu rapidly accumulated this metal, but was unable to readily excrete it when transferred to an uncontaminated environment. Moreover, classical Cu toxic effects related to ionic and osmotic disturbances, such as inhibition in Na/K-ATPase activity and reduced levels of major ions, were not observed. Despite that, a reduction

in carbonic anhydrase activity was seen, indicating that the excretion of nitrogenous compounds may have been compromised. Finally, it is concluded that populations of *H.luetkenii* living in an environment chronically contaminated by Cu-developed biochemical mechanisms to sustain osmoregulation even in the face of elevated accumulation of this metal.

**Supplementary Information** The online version contains supplementary material available at <https://doi.org/10.1007/s00244-021-00895-3>.

**Acknowledgements** Thanks to Dr. Sandra Isabel Moreno Abril for her help in the field experiment. The present study was financially supported by grants from the Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq, INCT—Toxicologia Aquática # 573949/2008-5), CAPES (# 230/2010), and IDRC (Ottawa, ON, Canada, # 104519-027). A.B (Proc. # 307647/2016-1) is a CNPq fellow was supported by IDRC.

**Authors' Contribution** All authors contributed equally.

**Funding** Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq, INCT—Toxicologia Aquática # 573949/2008-5), CAPES (# 230/2010), and IDRC (Ottawa, ON, Canada, # 104519-027). A.B (Proc. # 307647/2016-1).

## Declarations

**Conflicts of interest** There were no conflict of interests.

**Ethics Approval** All experimental and laboratorial procedures performed in the present work were previously approved by the university ethics committee (CEUA; protocol # 23116.001365/2015-44) and by the Brazilian Ministry of the Environment (MMA; research license # 44769-1).

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