Sublethal concentrations of waterborne copper induce cellular stress and cell death in zebrafish embryos and larvae

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ABSTRACT

Copper is an essential ion that forms part of the active sites of many proteins. At the same time, an excess of this metal produces free radicals that are toxic for cells and organisms. Fish have been used extensively to study the effects of metals, including copper, present in food or the environment. It has been shown that different metals induce different adaptive responses in adult fish. However, until now, scant information has been available about the responses that are induced by waterborne copper during early life stages of fish. Here, acute toxicity tests and LC50 curves have been generated for zebrafish larvae exposed to dissolved copper sulphate at different concentrations and for different treatment times. We determined that the larvae incorporate and accumulate copper present in the medium in a concentration-dependent manner, resulting in changes in gene expression. Using a transgenic fish line that expresses enhanced green fluorescent protein (EGFP) under the hsp70 promoter, we monitored tissue-specific stress responses to waterborne copper by following expression of the reporter. Furthermore, TUNEL assays revealed which tissues are more susceptible to cell death after exposure to copper. Our results establish a framework for the analysis of whole-organism management of excess external copper in developing aquatic animals.
**INTRODUCTION**

Multicellular organisms have developed sophisticated strategies to maintain a proper balance of copper and other metals in the body. To date, substantial information has been accumulated on the effects of metals like copper, cadmium, mercury and other heavy metals in organisms (Blechinger et al., 2002b; Coronado et al., 2001; Dave and Xiu 1991; Gravenmier et al., 2005; Szebedinszky et al., 2001). Two disorders in humans are associated with either lack or excess of copper in the organism. In Menkes disease, intestinal copper transport is blocked (Ambrosini and Mercer 1999; Camakaris et al., 1999; Mercer and Llanos 2003), while in Wilson's disease there is an excess of copper in the organism due to a failure in excretion of the metal from hepatic cells (Harris 2001; Mercer and Llanos 2003). Disorders in copper metabolism can be emulated in animal models, for example, using knockout mice or mutant zebrafish recovered in genetic screening. The absence of the high affinity copper transporter, Ctr1, causes early embryonic death in mice homozygous for this mutation (Kuo et al., 2001; Lee et al., 2001). Heterozygous animals are viable, but show increased sensitivity to low copper levels in the diet and accumulate subnormal copper levels in some tissues (Kuo et al., 2001; Lee et al., 2001). The zebrafish Ctr1 protein, which has 70% identity with its human cognate, is also essential during early development, as loss of function of this gene causes larvae to die three days after fertilization (Mackenzie et al., 2004). Recently, the identification of a zebrafish hypomorphic mutation in the ortholog of the Menkes disease gene (*atp7a*) was reported (Mendelsohn et al., 2006; Madsen et al., 2008; Madsen and Gitlin, 2008). This mutant shows a notochord abnormality, a lethal defect for the developing embryo.

Zebrafish have been widely used for detection of heavy metal contamination, making this animal a convenient biological water contaminant sensor (Blechinger et al., 2002b; Chan and Cheng 2003; Dave and Xiu 1991; Li et al., 2004; Ribeyre et al., 1995; Rougier et al., 1996). However, there is little information on the effects of copper on early fish development and its potential toxicity at sub-lethal concentrations (around 5 µM of Cu) (Hernandez et al., 2006; Sandahl et al., 2007; Seok et al., 2006). In adult fish, copper uptake is largely regulated by the gills (Grosell et al., 2004; Kamunde et al., 2003; Matsuo et al., 2004; Taylor et al., 2003), but in developing embryos, prior to gill development; other mechanisms could account for this process.

The purpose of this work is to characterize the acute effects of copper on zebrafish early life stage survival, development and physiology. We aim to know whether raising zebrafish under conditions of excess copper in the water has an impact on the uptake of the metal and on gene expression and cell survival. We have taken advantage of a useful transgenic line that expresses EGFP under the control of the promoter for the heat shock protein 70 (*hsp70*) gene (Halloran et al., 2000). Hsp70 is a stress-induced protein and appears as a convenient marker for the response of cells with altered metabolism and avoidance of cell death (Mosser et al., 2000; Stankiewicz et al., 2005), and it has been used before to assay the effects of external cadmium on developing zebrafish (Blechinger et al., 2002b; Krone et al., 2005; Matz and Krone, 2007). Our results show that dissolved copper in the medium is uptaken and accumulates in zebrafish larvae in a concentration and exposure-time dependent...
manner. In sub-lethal doses, copper elicits a cellular stress response and induces cell apoptosis in tissues involved in copper uptake and metabolism. Our studies suggest that long-term exposure to low doses of copper in the aquatic environment can affect embryonic health and, ultimately, survival in fish.

MATERIALS AND METHODS

Zebrafish maintenance

A breeding colony of AB wild type zebrafish (Danio rerio) was maintained at 28.5°C on a 14:10 hour light: dark cycle (Westerfield, 1995). All embryos used were collected by natural spawning and staged according to Kimmel et al., (1995) and were raised at 28°C in E3 medium (5 mM NaCl, 0.17 mM KCl, 0.33 mM CaCl₂, 0.33 mM MgSO₄, and 0.1% Methylene Blue (Merk, Darmstadt, Germany) in Petri dishes as described by Haffter et al., (1996). We express the larval ages in hours post-fertilization (hpf) or days post-fertilization (dpf). Copper was added as CuSO₄ (Merk, Darmstadt, Germany) to E3 medium, larvae were exposed for the required times and copper concentrations and subsequently rinsed a minimum of three times in fresh medium before processing for further analysis.

Mortality curves and determination of LC50 values

Embryos or larvae were exposed to different concentrations of copper sulphate. For the LC50 calculation, 72 hpf larvae were exposed to 0, 1, 5, 10, 25, 50, 100, 250 or 500 µM CuSO₄ dissolved in E3 medium. The larvae were maintained in the medium and the number of dead larvae were counted and removed at 1, 2, 3, 6, 18, 24, 30, 42, 48, 54, 66, 71, 78, 90 and 96 h during incubation with the metal. For each concentration of copper, 15 larvae were incubated in Petri dishes (10 cm, with 40 ml of medium) and at each time point we recorded the number of dead larvae and calculated the mean from three separate experiments; results were expressed as the percentage of dead larvae with respect to the starting number of fish. The medium was changed once a day as normal maintenance, maintaining the corresponding copper concentration or control E3 medium. Then, we plotted the data and calculated LC50 values using a 95% confidence test, Trimmed Spearman-Karber software, version 1.5 (Hamilton et al., 1977).

Determination of copper uptake and accumulation in larvae

We determined copper accumulation in the larvae using two methods: measuring the uptake of ⁶⁴Cu and quantifying the total copper content in an atomic absorption spectrometer (ASS).

Uptake of ⁶⁴Cu in zebrafish larvae

72 hpf zebrafish larvae (10 larvae per sample, in triplicate) were exposed to 10 mM ⁶⁴CuSO₄ (specific activity of 1.6 mCi/ mg, Chilean Nuclear Energy Commission) dissolved in the medium for 0, 5, 15, 30, 60 or 120 minutes. Then the larvae were washed 5 times with ice-cold 10 mM EDTA-PBS and dissolved in scintillation liquid.


64Cu was quantified in a beta-counter (Packard Tricarb 2100TR liquid scintillation analyzer) and calibrated to a standard curve of radiolabeled CuSO₄ dissolved in metal-free water. To quantify the protein in each sample, we took triplicate samples for each incubation time determined the protein concentrations using the Bradford method.

**Accumulation of copper in larvae**

To quantify the level of copper in the larvae, 72hpf larvae were exposed to concentrations of 0, 1, 5, 10, 25, or 50 CuSO₄ for 6 hrs. Then the larvae were washed 8 time with 1 ml of metal-free water (10 larvae per sample, in triplicate) and then the larvae were digested with concentrated ultrapure nitric acid (1: 1) overnight at 60°C. Copper content were determined by atomic absorption spectrometer (AAS) equipped with a graphite furnace (SIMAA 6100, Perkin Elmer, Shelton, CT). MR-CCHEN-002 (Venus clams) and DOlt-2 (Dogfish liver) preparations were used as reference materials to validate the mineral analyses. Proteins of each sample were determined using the Bradford method in triplicate samples.

**Exposure of the hsp70:egfp transgenic zebrafish line to waterborne copper**

In order to obtain information on the stress response induced by acute exposure of zebrafish to copper, we incubated transgenic Tg(hsp70:egfp) zebrafish in different concentrations of copper sulphate. 72 hpf larvae were exposed for 2 h to 100, 200, and 400 µM CuSO₄ dissolved in E3 growth medium (n=65 to 70 larvae each concentration). We analyzed EGFP expression under UV light to detect the activation of the hsp70 promoter compared to control animals. The larvae were maintained in Petri dishes and E3 medium was renewed daily. After incubation, the animals were incubated in E3 copper-free medium for 24 h and the accumulation of EGFP was monitored over the next five days. The larvae were observed under UV fluorescence in a Leica MZ12.5 dissecting scope and recorded with an Optronics 60800 camera. We recorded the visible expression of EGFP in five tissues during the assays: gills, olfactory pit, liver, spinal cord and brain.

**Immunohistochemistry**

To further analyze the organ-specific expression of EGFP induced by copper exposure in the Tg(hsp70:egfp) larvae, we carried out immunostains against the reporter protein. 72 hpf transgenic larvae were exposed for 2 h to 300 µM CuSO₄ dissolved in E3 Medium, washed and then maintained in E3 medium for 72 hours. Then the embryos were fixed in fresh 4% paraformaldehyde in phosphate buffered saline (PBS) overnight at 4°C. The fixed larvae were dehydrated through an ethanol series (50%, 70%, 80%, 90%, 100% I and 100% II; Merk, Darmstadt, Germany), cleared in xylene (Merk, Darmstadt, Germany) and mounted in Paraplast Plus® (Tyco healthcare group LP, Gosport, UK). All incubations were for 20 minutes. The sections (6 mm) were mounted in silane-coated slides. Mounted sections were dewaxed in xylene and rehydrated through an ethanol series. Immunohistochemistry was performed in slides incubated in a humidity chamber. Washing and dilution of immunoreagents was performed with PBS with 0.1% Tween-20 (PBST) throughout, and three PBST washes (5 min each) were performed between each antibody incubation. To avoid non-specific binding of antibody, we incubated with blocking solution (20% lamb serum, 0.1%, DMSO and 0.1% Tween-20; Sigma-Aldrich Co.) at room temperature for 1 h. EGFP
detection was performed using rabbit anti-GFP polyclonal antibody (Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA, SC-8334, 1:500), incubated overnight at 4°C. An alkaline phosphatase (AP)-conjugated goat anti-rabbit IgG (H+L) was used as the secondary antibody (KPL, Inc, Gaitherburg, MD, USA, Cat. N°375-1506, 1:500), incubated for 1 h at room temperature. The stain was developed by incubation for 15 min with 0.4 mg/ml NBT (Nitro Blue Tetrazolium Chloride, Roche Diagnostic, Manheim, Germany), 0.19 mg/ml BCIP (5-bromo-4-chloro-3-indolylphosphate, toluidine salt, Roche, Diagnostic, Manheim, Germany) in 100 mM Tris buffer (pH 9.5) and 50 mM MgSO₄.

After immunostaining, all sections were dehydrated in ascending concentrations of ethanol, cleared with xylene, and coverslipped with Cytoseala mounting media (Stephen Scientific, NJ, USA).

**Whole-mount TUNEL staining**

To detect apoptotic cells induced by copper we incubated 72 hpf larvae with 300 μM copper for 2 h. The embryos were then rinsed with E3 copper-free medium and maintained for 24 or 72 h and fixed. The larvae were assayed for whole-mount terminal deoxynucleotidyl transferase-mediated dUTP nick-end labeling (TUNEL) using an in situ cell death detection kit (Roche, Diagnostic, Manheim, Germany, Cat. N°11684809910). The TUNEL assay was performed according to a previously described method (Kozlowski et al., 2005) with some modifications. After the treatment larvae were fixed overnight at 4°C in 4% of paraformaldehyde in PBS. The samples were washed three times for 15 min in PBST (0.1% Tween-20 in PBS) and dehydrated in methanol 100% at -20°C for 24 h. To rehydrate the samples following methanol incubation, each was rehydrated for 5 min with a methanol/PBST gradient until reaching 100% PBST. Then, larvae were digested with Proteinase K (GIBCO, Carlsbad, CA, concentration 10 mg/ml) for 30 min at room temperature. Digestion was stopped by washing in PBST twice for 5 min and re-fixed in 4% paraformaldehyde for 20 min at room temperature, followed by three washes with PBST for 5 min each and one wash in terminal deoxynucleotidyl transferase (TdT) buffer for 10 min. Then the larvae were pre-incubated with TdT enzyme with labeled nucleotides (DIG-dUTP) at 4°C for 60 min, and then incubated successively at 37°C for 1 h. After the enzymatic labeling, the larvae were maintained at 4°C in blocking solution for 2-3 h and overnight at 4°C with anti-DIG (1:2000 in blocking solution). After incubation, the embryos were washed in PBST for 20 min four times and pre-incubated with AP buffer (three times for 5 min). The samples were stained using a solution containing NBT/BCIP in AP buffer. The reaction was stopped by transferring the larvae to PBST and washed 3 times for 5 min. The embryos where observed in a stereomicroscope (Leica MZ12.5) and photographed with Optronic camera.

**Data analyses**

Variables were tested in triplicates, and experiments were repeated at least twice. One-way ANOVA was used to test differences in mean values, and Turkey's post-hoc test was used for comparisons (InStat program from GraphPad Prism). Differences were considered significant if p < 0.05.

**RESULTS**
Effect of copper on larval survival

We determined the lethality curves for copper sulphate dissolved in embryo medium (E3) by incubating 72 hpf larvae in different concentrations of the metal and measuring mortality up to 96 hours after initiation of the treatment (7 days post-fertilization, dpf). Mortality was recorded at different times (1, 2, 3, 6, 18, 24, 30, 42, 48, 54, 66, 71, 78, 90 and 96 hours after initiation of treatment) by counting larvae that showed evident necrosis, absence of heart beat or failure to move with mechanical stimulation. Fish were counted and the percentage of dead larvae with respect to the initial number was calculated. In control samples of fish (no copper added to the medium), a mortality of approximately 10% was observed after 72 hours of incubation (6 dpf), which increased to 20% at 96 hours postincubation (7 dpf) (Figure 1A). Larvae incubated in the range of 1 to 10 µM of copper show mortality rates similar to those of the control larvae. At 25 µM CuSO₄, we observe 100% mortality at 90 hours post-incubation (hpi). The fish treated with 50 and 100 of CuSO₄ show 100% mortality at 54 hpi. Concentrations of 250 and 500 µM dissolved copper present 100% mortality at 24 and 18 hpi, respectively (Figure 1A). Thus, this result showed a direct correlation between copper concentration in the water and lethality.
From the cumulative mortality curves we calculated the lethal concentration 50 (LC50) values for each incubation time. After plotting the LC50 values versus time we found a logarithmic decay curve (Figure 1B), the expected fit for LC50 values versus time obtained from lethality curves or mathematical modeling (Tsai and Liao 2006).

Figure 1. Waterborne copper-induced lethality of zebrafish larvae depends on concentration of the metal. (A) Copper lethality curves were determined at different copper concentrations in the media. Larvae of 72 hpf were incubated in presence of 0 (○), 1 (∆), 5 (○), 10 (□), 25 (□), 50 (●), 100 (▲), 250 (●) and 500 (■) µM of CuSO₄ and maintained at 28 °C for 96 h exposed to the metal. The number of dead larvae were counted and removed at the indicated times: % mortality refers to the accumulated percentage of dead larvae with respect to the initial number of larvae. (B) LC50 values were calculated from the lethality curves using the Trimmed Spearman-Karber method. LC50 values diminish, as the time of incubation is longer. Fitting the logarithmic tendency curve to the actual data shows a strong correlation (R² = 0.9221).
Copper incorporation and accumulation in zebrafish larvae

In order to determine whether dissolved copper present in the medium is uptaken and accumulated into developing zebrafish, we incubated 72 hpf larvae with 10 µM $^{64}$CuSO$_4$ for different lengths of time and incorporated radioactivity was measured. When compared to untreated larvae, we found that radiolabeled copper is uptaken from the medium into the $^{64}$CuSO$_4$ treated larvae and this incorporation increases with incubation time (Figure 2A). While copper uptake was not significant between 5 and 30 min compared to control fish, after 60 and 120 minutes of incubation, copper uptake was significantly higher compared to control larvae (one-way ANOVA, p<0.01).

Additionally, we determined if copper present in the medium accumulated in the larvae. The copper uptake measured by AAS show that larvae incubated with 1 to 25 µM of CuSO$_4$ do not show an increase in copper content (Figure 2B). However at 50 µM (approximately 3.15 mg/L) of copper in the medium we detected accumulation of the metal in the larvae, increasing from 2.87 to 7.36 nmol of Cu/mg protein, compared with control larvae (one-way ANOVA, p<0.05).
Exposure to copper induces hsp70 gene promoter activation in transgenic zebrafish larvae

Cells respond to metabolic stress by inducing the expression of a number of genes including those of the heat shock response such as Heat Shock Proteins (HSPs). We took advantage of the availability of an \textit{hsp70:egfp} transgenic zebrafish line, \textit{Tg(hsp70:egfp)} (Halloran et al., 2000) to assay the effect of a copper-induced stress response in developing larvae. This strain of fish expresses enhanced green fluorescent...
protein (EGFP) in an almost identical pattern to that of the endogenous hsp70 gene either during normal development, after heat shock (Halloran et al., 2000), or after cadmium exposure (Blechinger et al., 2002b).

We exposed transgenic 72 hpf larvae to a 2 h pulse of high copper-loaded media (Figure 3A). Control larvae show a basal expression in the lens as previously reported (Halloran et al., 2000; Blechinger et al., 2002a) (Figure 3A), while heat-shocked larvae show the characteristic pattern of EGFP label throughout the body (Supplementary Figure 1). Exposure to low concentrations of copper (below 100µM CuSO₄) did not appreciably modify the levels of detectable EGFP compared to control fish. Transient exposure to 100µM CuSO₄, induces expression of EGFP in gills and olfactory pits, and weak expression in brain, liver and spinal cord (Figure 3B and 3E); this expression increases at 200 ^iM of CuSO₄ and fluorescence appears in the olfactory epithelium and spinal cord (Figure 3C and 3F). Treatment with a higher concentration of copper (400µM CuSO₄) induces strong expression in spinal cord, liver and brain (Figure 3D and 3G). We documented the expression of EGFP in the different larval tissues for each concentration at different exposure times by counting the larvae in each sample that showed label in these tissues and expressing this value as a percentage of the total number of larvae (Figure 3 E-G). These results show that induction of the hsp70 promoter by waterborne copper elicits tissue specific effects depending on the amount of metal present, and that different tissues may show varying susceptibility to copper toxicity or that copper load is distributed differently under different conditions.
Figure 3. The Hsp70 promoter is induced by copper in a dose-dependent manner. Transgenic Tg(hsp70:egfp) zebrafish larvae of 72 hpf were exposed transiently to a 2 h pulse with 0 µM (A), 100 µM (B), 200 µM (C) and 400 µM (D) of waterborne copper (as CuSO$_4$) in the medium. Larvae were transferred to copper free medium, kept for a further 24 hours, and observed under a fluorescence stereomicroscope for recording EGFP expression. Note the rise in fluorescence as concentrations of copper increased. As different organs showed fluorescence under the different conditions used and there was variability within a batch of embryos, we recorded the tissues where fluorescence was visible. The graphs in panels E-F show the percentage of larvae per batch that had visible expression of EGFP at the indicated times in each analyzed tissue. Note that the EGFP expression profile is specific to each time point and at each concentration of copper used. Panels A-D show lateral views, anterior to left. 70 larvae were used per each concentration for panels E, F and G.
To analyze the expression of EGFP in higher detail, 72 hpf transgenic larvae were exposed to 300 µM copper for 2 h and processed for EGFP immunodetection at 72 h after copper exposure (Figure 4). Observations of living control larvae under fluorescent illumination shows expression of EGFP exclusively in the lens (Figure 4A). In contrast, exposed larvae show expression in olfactory pit, gills, liver, spinal cord and brain (Figure 4B). Anti-GFP immunohistochemistry in sectioned slices of copper exposed fish shows expression of EGFP in the retina and also in the telencephalon (r and t, respectively, Figure 4C). Interestingly, posterior transverse sections evidence the expression of EGFP in the spinal cord (sc), pronephric glomeruli (pg), pronephric ducts (pd), pneumatic duct (pnd), stomach (s), liver (L) and intestine (i) (Figure 4D and 4E).

**Figure 4.** Tissue-specific stress response after treatment with high concentrations of waterborne copper: 72 hpf *Tg(hsp70:egfp)* transgenic larvae were exposed for 2 h to 300 µM CuSO₄ and...
observed under UV light 72 hours after copper exposure. Control animals show background EGFP fluorescence only in the lens (A). In treated larvae, expression of EGFP was evident in olfactory epithelium (oe), gills (g), central nervous system (cns), and liver (B). EGFP detection was also carried out by immunohistochemistry in sections by using an antibody against EGFP and a secondary antibody conjugated to alkaline phosphatase (AP). Sections show EGFP expression in retina (r), telencephalon (t), liver (L), stomach (s), pronephric ducts (pd), pneumatic duct (pnd), pronephric glomeruli (pg), spinal cord (sc) and intestine (i) (C to E). Sections were counterstained with eosine after developing the alkaline phosphatase activity; the corresponding transverse sections are indicated in B. In A and B pictures are lateral views, anterior to left. In C, D and E are transverse sections, dorsal is up; y, yolk.

**Cell death in zebrafish larvae exposed to copper**

Having established the expression pattern of the *hsp70* gene in larvae exposed to copper, we investigated the spatial distribution of apoptotic cells in copper-treated larvae after acute exposure treatments by using the TUNEL assay. Larvae exposed for 2 hours to 300µM CuSO$_4$ and processed by TUNEL 72 h after exposure show an increase of apoptotic staining in the central nervous system (cns), gills (ga) and also strong labeling in the pronephros (prn) (Figure 5B and 5D) compared to control larvae at the same stages (Figure 5A and 5C).

**Figure 5.** Apoptosis is induced by waterborne copper exposure. Wild type AB strain larvae of 72 hpf were exposed continuously for 2 h to 300 µM of CuSO$_4$ and after a further 72 h in E3 medium (6 dpf) whole-mount TUNEL staining was performed to identify apoptotic cells. Exposed larvae (B, D) show staining in the central nervous system (cns), gills (g) and strong labeling in the pronephros (prn) in comparison to sibling controls (A, B).

**DISCUSSION**
A variety of studies have shown that fish, as well as other aquatic organisms, are particularly sensitive to pollutants and heavy metals in the early stages of their life cycle (Herkovits et al., 1998; Hernandez et al., 2006; Linbo et al., 2006; Witeska et al., 1995). However, some studies have shown that fish eggs are more resistant than young fish to the toxic effects of copper sulfate. In fact, experiments using zebrafish as an animal model show that adult zebrafish are more sensitive to copper than embryos (Palmer et al., 1998). These authors calculated a 4-day median lethal concentration 50 (LC50) for adults (five to nine months old) of 1.90 mM and a 12-day LC50 for embryos of 2.2 mM. Nevertheless, copper toxicity in fishes varies among species and depends on the physical and chemical characteristics of the water (Kamunde et al., 2003).

In our study, acute toxicity tests of waterborne copper were performed in zebrafish larvae (72 hpf) and the LC50 values for dissolved copper sulphate were calculated. Our lethality curves indicate that there is an increase in the mortality rate as the copper concentration is increased, as well as when the incubation with waterborne copper is longer. We found that at concentrations lower than 10 µM of copper (approximately 0.63 mg/L of copper) the larvae present similar mortality rates to controls. However at concentrations higher than 25 µM (1.6 mg/L) the mortality was significant increased. Calculations of the LC50 value for each exposure time show a logarithmic dependence between the LC50 and time of exposure. Furthermore, the LC50 value calculated for larvae exposed to copper continuously for 4 days was 13.82 µM, showing that larvae are more resistant to waterborne copper than adults, where this value was calculated at 1.9 µM (Palmer et al., 1998).

To determine whether waterborne copper is incorporated in larvae, we measured copper uptake and loading by using radiolabeled copper and AAS, respectively. Our data indicate that copper is incorporated into larvae in a time-dependent manner and also that the metal accumulates in larvae depending on its concentration in the medium. The primary uptake pathway of trace metals in fish is the food, under normal dietary and waterborne conditions (Kamunde et al., 2003; Kjoss et al., 2005; Craig et al., 2009), even though some reports suggest that copper and iron can also be taken up by the gills (Clearwater et al., 2002; Grosell and Wood, 2002; Kamude et al., 2003). In rainbow trout diet contributes more than 90% of the body burden of this metal. However, gill-uptake contributes importantly to the body Cu load when the dietary source of copper is deficient (Kamunde et al., 2003). As the larvae in our tests are not feeding, the mechanism of entry is limited to the body surface, especially exposed tissues not covered by the epidermis (gills, olfactory epithelium, mechanosensory organs). Copper could be entering cells via the same mechanisms that mediate entry in adult tissues, Na+ dependent transporters, copper transporter-1 (Ctrl) and the divalent metal transporter 1 (Dmtl). We have previously shown that Ctrl is expressed in zebrafish during early embryogenesis, providing a possible transport mechanism at this stage (Mackenzie et al., 2004).

Based in our data of mortality assays and LC50 values we designed experiments to examine whether copper is capable of inducing the expression of the hsp70 gene, a chaperone whose expression is activated by heat shock. This protein has been reported to be induced by a variety of heavy metals including Zn, Cd, Hg and Cu (Murata et al., 1999; Williams et al., 1990) and this activation is mediated by the heat shock transcription factor HSF-1 (Mosser et al., 1988; Murata et al., 1999). Recent work has shown that copper can activate the human hsp70 promoter in mosaic transgenic fish (Seok et al., 2006), and previous reports show that heat shock and cadmium induce the expression of EGFP in Tg(hsp70:egfp) fish expressing EGFP driven by the hsp70 gene promoter (Blechinger et al., 2002a; Blechinger et al., 2002b; Blechinger et al., 2002b).
Using these transgenic fish, we found that waterborne copper induces EGFP expression in a concentration-dependent manner. The histological analysis in sections shows that the expression induced by copper is localized mainly to the head and, especially, the brain. Moreover, waterborne copper induces EGFP expression in organs such as the spinal cord, pronephric glomeruli, pronephric ducts, pneumatic duct, liver, stomach and intestine. The liver, intestine and renal tissues have been shown to accumulate copper in toadfish and rainbow trout after waterborne copper exposure (Grosell et al., 2004; Kamunde et al., 2003), suggesting that this selective accumulation of copper in these tissues could be a protection mechanism against internal copper toxicity. Studies using primary cultures of rainbow trout hepatocytes exposed to CuSO4 (0-200 mM) resulted in both a dose-dependent elevation of hsp70 expression and cell death by apoptosis (Feng et al., 2003). Moreover, in vitro studies show that copper exposure is also able to induce the expression of metallothioneins (MTs) (Cheuk et al., 2008), small cysteine-rich proteins with high affinity for heavy metals, that could complex the copper ions and protect the cells from toxicity. Our results were similar to those previously reported for cadmium-induced EGFP expression (Blechinger et al., 2002a; 2007). In our experiments we found additional expression sites such as the brain and spinal cord in fish exposed to high copper concentrations, and also that appearance of the marker in some tissues depended strongly on the incubation time. While EGFP was induced in the brain only at the highest concentration tested, it could reveal that metal toxicity at this concentration overwhelms the physiological capacity of the larva to prevent its dissemination. Nonetheless, expression in the brain appeared to decrease over time, possibly indicating progressive recovery of the liver or other detoxifying systems. The gills and olfactory epithelium showed EGFP expression in the larvae exposed to 100 µM copper from the earliest time point analyzed, 24 hpf. Interestingly MT is expressed in these tissues during normal embryogenesis of zebrafish (Chen et al., 2004), suggesting that these tissues could be “primed” for protection against accumulation of copper or other metals. Prolonged incubation induced the expression of EGFP in liver (where MT is not expressed in developing fish, Chen et al., 2004). Our results also differ from those previously reported, in which mosaic transgenic fish carrying EGFP under the control of the exogenous human hsp70 promoter were exposed to copper (Seok et al., 2006). These authors show mosaic EGFP expression in gills, skin epithelium, auditory epithelium and myotubes. The differences can be explained by the fact that these authors used fish transiently expressing EGFP from the human hsp70 promoter and obtained mosaic expression of EGFP. This is in contrast to the stable transgenic Tg(hsp70:egfp) line used in our experiments that expresses EGFP in the entire embryo. In addition, these authors reported a calculated LC50 of 1.2 µM of copper in contrast to the value of 13.82 µM calculated in our experiments, a difference of an order of magnitude (Seok et al., 2006). There are several possible explanations for these differences such as the strain of fish used (AB strain in our study versus a non-typified strain), ion composition of the media (E3 in our experiments versus Ringer's solution) and water hardness (Long et al., 2004; Van Genderen et al., 2007).

It is interesting to note that exposure of fish to copper at stages earlier than 72 hpf did not induce expression of EGFP in the organs involved in copper metabolism, such as liver, intestine and the renal system (data not shown). The onset of circulation begins at 24 hpf, while at 48 hpf a simple circulatory system has formed. Vascularization of the zebrafish pronephros and the onset of glomerular filtration occur between 40 and 48 hpf (Drummond et al., 1998). In zebrafish, as in all teleosts, nutrition in early larval stages depends on the yolk until the digestive organs develop and the larvae start to
At 96 hpf, the digestive organs are formed and liver cells start secreting bile at 72-96 hpf (Pack et al., 1996). As no hsp70 response was observed until 96-125 hpf in these tissues, we suggest that the mechanisms that regulates the distribution of copper in the body appears concomitantly with vascularization and organogenesis.

High intracellular copper has been described as toxic to the cells via generation of reactive oxygen species (ROS) (Pourahmad and O'Brien, 2000). A major cytotoxic role for ROS includes the activation of the apoptosis pathway and some studies clearly implicate a role for ROS in copper-induced apoptosis in vitro (Ma et al., 1998; Pourahmad and O'Brien, 2000; Zhai et al., 2000). Heat shock proteins are key players in the cellular stress response process and crucial for defending cells from copper toxicity (Ma et al., 1998). Our TUNEL assays in larvae acutely exposed to 300 µM (18.9 mg/L) copper show that waterborne copper induces apoptosis in tissues including the pronephros, brain and gills, organs that also showed expression of EGFP in transgenic Tg(hsp70:egfp) larvae after similar copper treatments. Stress in these tissues followed by cell death is a likely consequence of the exposure to copper. However, we did not detect TUNEL staining in the liver, an organ that showed a strong stress response by visualization of the reporter protein. In contrast, the pronephros showed the strongest TUNEL staining, even though EGFP expression in this organ was only detected by immunostaining. These observations suggest that in some tissues, such as liver tissue, the stress response induced by copper allows cell survival, probably by strong induction of MT expression, while other organs, such as the pronephros, fail to show a stress response and manifest high levels of cell death.

The data above indicate that copper is toxic for zebrafish larvae because the metal is incorporated and accumulates in diverse tissues, as is suggested by the expression of EGFP in Tg(hsp70:egfp) larvae and TUNEL assays after copper treatments. Our results provide evidence that copper can be lethal to developing fish when it is present in the water, because of the deleterious effects on several key organs.

In this work we observed that when transgenic zebrafish larvae are exposed to external copper, the hsp70:egfp transgene was expressed in different tissues in a dose-dependent manner. Thus, this resource represents an excellent system to evaluate how the larva copes with external copper and how it is able to restrict the distribution of accumulating metal to different tissues. However, it is not capable of detecting the subtle effects induced by low doses of waterborne copper, limiting its potential use as a live biosensor tool. Finally, our findings show that the most sensitive organs to stress induced by waterborne copper are the central nervous system and the liver, even though the most affected in terms of cell death are the gills and pronephros. Cell death is likely to be elicited by the induction of ROS, providing a possible immediate cause for the death of exposed larvae.

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NOTES

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