The Role of Zinc in the Modulation of Neuronal Proliferation and Apoptosis

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Abstract Although a requirement of zinc (Zn) for normal brain development is well documented, the extent to which Zn can modulate neuronal proliferation and apoptosis is not clear. Thus, we investigated the role of Zn in the regulation of these two critical events. A low Zn availability leads to decreased cell viability in human neuroblastoma IMR-32 cells and primary cultures of rat cortical neurons. This occurs in part as a consequence of decreased cell proliferation and increased apoptotic cell death. In IMR-32 cells, Zn deficiency led to the inhibition of cell proliferation through the arrest of the cell cycle at the G0/G1 phase. Zn deficiency induced apoptosis in both proliferating and quiescent neuronal cells via the intrinsic apoptotic pathway. Reductions in cellular Zn triggered a translocation of the pro-apoptotic protein Bad to the mitochondria, cytochrome c release, and caspase-3 activation. Apoptosis is the resultant of the inhibition of the prosurvival extracellular-signal-regulated kinase, the inhibition of nuclear factor-kappa B, and associated decreased expression of antiapoptotic proteins, and to a direct activation of caspase-3. A deficit of Zn during critical developmental periods can have persistent effects on brain function secondary to a deregulation of neuronal proliferation and apoptosis.

Keywords Zinc · Neuron · Proliferation · Apoptosis · Caspase · Nuclear factor-kappa B (NF-κB) · Extracellular-signal-regulated kinase (ERK) · Zinc deficiency · Cortical neuron · Bad

Abbreviations Cdks Cyclin-dependent kinases DTPA Diethylenetriamine pentaacetic acid EMSA Electrophoretic mobility shift assay ERK Extracellular-signal-regulated kinase FBS Fetal bovine serum MAPK Mitogen-activated kinases NF-κB Nuclear factor-kappa B Zn Zinc

Introduction

Zinc (Zn) is an essential nutrient that has multiple roles in the nervous system (Frederickson et al. 2005); deficits of this nutrient can result in numerous diverse pathologies (Rogers et al. 1990; Keen et al. 1995; Sandstead 2003). Zn exists largely bound to proteins, and numerous enzymes require Zn including DNA and RNA polymerases, several enzymes in the intermediary metabolism, and most metalloproteinases. Zn is also a structural component of a family of Zn-binding proteins known as Zn-finger proteins that act as DNA-binding transcription factors. While protein
bound Zn is important, large amounts of Zn are not protein bound. For example, about 10% of Zn in the nervous system is associated with pre-synaptic vesicles of glutaminergic neurons; this Zn, which is not protein bound, is released as free Zn into the synaptic cleft where it can modulate the activity of a variety of post-synaptic receptors (Frederickson et al. 1995).

In rodent models, severe Zn deficiency throughout pregnancy can result in multiple brain malformations (Hurley and Swenerton 1966), and deficits of this nutrient during the early postnatal period can affect the number and differentiation of neurons in the cerebellum (Dvergsten et al. 1983, 1984a, b). Mild Zn deficiency can result in alterations in learning, attention, and memory in rodents and non-human primates (reviewed in Golub et al. 1995, 2000). In humans, maternal dietary Zn intakes during pregnancy have been reported to be inversely associated with the head circumference of the infant at birth (Lagiou et al. 2005), and Zn supplementation has been associated with larger head circumferences in infants from women with low body mass index (Goldenberg et al. 1995). There is a growing body of evidence that supports the concept that in humans, suboptimal Zn nutrition during development is associated with altered neonatal/infant behavior, cognitive, and motor performance (Kirksey et al. 1994; Bentley et al. 1997; Penland et al. 1997; Gardner et al. 2005).

Zn plays an essential role in cell proliferation (MacDonald et al. 1998; Wang et al. 2002; Wong et al. 2007; Corniola et al. 2008) and apoptosis (reviewed in Truong-Tran et al. 2003; Clegg et al. 2005; Fraker 2005; Corniola et al. 2008) in different tissues and cell types. The regulation of cell proliferation by Zn can occur at different levels including the requirement of Zn for the activity of enzymes involved in DNA synthesis (i.e., deoxythymidine kinase), and the modulation of regulatory signals directly, as well as indirectly, through its effects on the hormonal regulation of cell division. Illustrative of this, the pituitary growth hormone-insulin-like growth factor-1 axis is responsive to Zn status (MacDonald 2000), although its role in Zn deficiency-associated decreases in cell proliferation is an issue of controversy (Ninh et al. 1998; Wang et al. 2002). Low cellular Zn concentrations can trigger apoptosis in numerous cell types, including fibroblasts, hepatocytes, T-cell precursors, glioma, and testicular cells. The mechanisms involved in the induction of apoptosis secondary to Zn deficiency are multiple, and differ among different cell types. Zn can act as an inhibitor of caspase-3, a cysteine protease that is central in apoptosis cleaving key proteins [e.g., poly(ADPribose) polymerase (PARP)] that continue the apoptotic cascade (reviewed in Truong-Tran et al. 2001). Zn deficiency can also induce apoptosis by disrupting growth factors signal transduction pathways mediated by receptor tyrosine kinases (Clegg et al. 2005). While somewhat controversial (Zago et al. 2005), it has also been proposed that an increase in cell oxidants triggers apoptosis (Clegg et al. 2005).

Studies with experimental animals, as well as in cell models, have shown that neurons can be particularly affected by a low availability of Zn. In rats, postnatal Zn deficiency impaired the normal development of the cerebellum, affecting the number and differentiation of Purkinje, basket, stellate, and granule cells (Sandstead 2003). In human neuroblastoma IMR-32 cells, we observed that with Zn deficiency there is a reduction in cell viability, with evidence of increased apoptotic death (Verstraeten et al. 2004; Zago et al. 2005).

In the nervous system, cell proliferation and apoptosis actively occur during the perinatal period, and are critical for normal neurodevelopment. Alterations in these tightly regulated events can affect the function of the nervous system latter in life. Given the close interrelationships between cell proliferation and death in the development of the nervous system, and to better understand the role of Zn in this process, in this study we investigated if a low Zn availability can affect neuronal proliferation and survival. As experimental models, we investigated the effects of Zn deficiency on neuronal proliferation in human neuroblastoma cells (IMR-32), and on apoptosis in IMR-32 cells, and primary cultures of differentiated rat cortical neurons. We show that with a deficit of Zn, there is an inhibition of neuronal cell proliferation that is secondary to an arrest at the G0/G1 phase of the cell cycle. Interestingly, in proliferating as well as in quiescent neurons, Zn deficiency triggers apoptotic neuronal death. The apoptosis occurs via the intrinsic pathway which can be a consequence of extracellular-signal-regulated kinase (ERK) inhibition, caspase-3 activation, and the down-regulation of nuclear factor-kappa B (NF-κB)-dependent anti-apoptotic genes.

Materials and Methods

Materials

The IMR-32 neuroblastoma cells were obtained from the American Type Culture Collection (Rockville, MA, USA). Cell culture media and reagents were purchased from Invitrogen Life Technologies (Carlabad, CA, USA). The CellTiter-Glo Luminescent Cell Viability Assay, the oligonucleotides containing the consensus sequence for NF-κB (5'-AGTTGAGGGGACTTTCCAGGC-3') and OCT-1, and the reagents for the electrophoretic mobility shift assay (EMSA) assay were obtained from Promega (Madison, WI, USA). Polyclonal antibodies for Akt, phosphorylated Akt (Ser 473), Bcl-2, c-myc, c-IAP1, and XIAP, and phosphorylated Bad (Ser 112 or Ser 136) were
obtained from Cell Signaling Technology (Beverly, MA, USA). Antibodies for p53, ERK1/2, phosphorylated ERK, p21, cyclin D1 and E, PARP, Bad, cytochrome c, Bcl-xL, and β-tubulin were from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Polyvinylidene difluoride (PVDF) membranes were obtained from Bio-RAD (Hercules, CA, USA). Merocyanine 540 and propidium iodide were obtained from Molecular Probes (Eugene, OR, USA). The ECL Plus western detection reagents were from GE Healthcare (Piscataway, NJ, USA). The APO-BrdUTM kit and BD CyclestestTM Plus DNA reagent kit were obtained from Pharmingen-Becton-Dickinson Co. (San Diego, CA, USA). The ApoAlert-Caspase-3 Assay was purchased from Clontech (Mount View, CA, USA). All other reagents were from the highest quality available and were purchased from Sigma (St. Louis, MO, USA).

Cell Cultures and Incubation

Zn-deficient fetal bovine serum (FBS) was prepared by chelation with diethylenetriamine pentaacetic acid (DTPA) as previously described (Oteiza et al. 2000; Mackenzie et al. 2002b). The chelated FBS was subsequently diluted with complex medium or DMEM to a final concentration of 3 mg protein/ml to match the protein concentration of the control non-chelated medium [10% (v/v) FBS]. The concentration of Zn in the deficient medium was 1.5 μM. Aliquots of the Zn-deficient media were supplemented with ZnSO₄ to a final Zn concentration of 15 μM.

IMR-32 cells were cultured at 37°C in complex medium [55% (v/v) DMEM high glucose, 30% (v/v) Ham F-12, 5% (v/v) α-MEM] supplemented with 10% (v/v) FBS and antibiotics-antimycotic (50 U/ml penicillin, 50 μg/ml streptomycin, and 0.125 μg/ml amphotericin B). Cells were grown in control medium [complex medium containing 10% (v/v) non-chelated FBS] until 90% confluence. Depending on the experimental conditions, synchronized or unsynchronized IMR-32 cells were incubated in control non-chelated medium or in chelated media containing 1.5 or 15 μM Zn. Cells were harvested at different time points (6–48 h) depending on each experiment. We previously reported that the incubation of IMR-32 cells in media containing low Zn concentrations (1.5 and 5 μM Zn) results in a reduction of cellular labile Zn pools within 3 h (Mackenzie et al. 2002a).

For the isolation of cortical neurons, timed-pregnant Sprague–Dawley rats were obtained from Charles Rivers Laboratories (Wilmington, MA, USA). All animal procedures were in agreement with standards for the care of laboratory animals as outlined in the NIH Guide for the Care and Use of Laboratory Animals. Primary cortical cultures were prepared from E19 embryos according to Banker and Goslin (1998). In brief, embryos were rapidly removed under sterile conditions, and after removal of the skull and meninges, the cerebral hemispheres were dissected out and placed in Hanks’ balanced salt solution prewarmed to 37°C. Cortical tissue was dissociated by enzymatic digestion (0.025% w/v trypsin). Trypsinization was stopped by the addition of Hanks’ solution containing 20% (v/v) FBS, and the tissue was mechanically dissociated in Hanks’ balanced salt solution containing 0.004% (w/v) DNAse with Pasteur pipettes of different diameters. The resulting cell suspension was centrifuged at 200 × g for 5 min. The cells were resuspended in a known volume of Neurobasal culture medium supplemented with 2% (v/v) serum-free additive B27, and plated on poly-L-lysine-coated dishes. Cell cultures were incubated at 37°C in an atmosphere of 95% (v/v) air, 5% (v/v) CO₂ with 90–95% (v/v) humidity. After 24 h in culture, the media were replaced by DMEM high glucose supplemented with 10% (v/v) FBS. After 72 h, cell cultures were supplemented with 40 μM cytosine-β-D-arabinoside for 24 h to inhibit non-neuronal cell proliferation. After 7–9 days in culture, the media were replaced by control non-chelated medium, or chelated medium containing 1.5 or 15 μM Zn, and cells were further incubated for 6–24 h depending on the experiment.

Cell viability for IMR-32 and cortical neurons was measured evaluating the cell ATP content (CellTiter-Glo Luminiscent Cell Viability) according to manufacturer’s protocol.

Evaluation of Cell Cycle Progression

For synchronization, IMR-32 cells were serum-starved for 48 h in complex medium containing 0.1% (v/v) FBS. Re-entry into G₁ phase of the cell cycle was initiated by replacement of the starvation medium with complete or chelated media containing 1.5 or 15 μM Zn. After 0, 12, 16, and 24 h, cells were collected, washed and the analysis of nuclear DNA was performed using the CyclestestTM Plus DNA Reagent Kit. Propidium iodide-stained nuclei were analyzed by flow cytometry on a Becton-Dickinson cytomter and the percentage of cells in the G₀-G₁, S, and G₂–M phases of the cell cycle were determined by using the Mod Fit LT cell cycle analysis software (Verity Software, Topsham, ME, USA).

Western Blot

Cells were washed in PBS and whole cell extracts were prepared by lysis in 50 mM Hepes buffer (pH 7.4), containing 150 mM NaCl, 0.5 mM EDTA, 2% (v/v) Igepal, phosphatase inhibitors (1 mM sodium pervanadate, 100 μM NaF), and protease inhibitors (1 mM PMSF, 5 μg/ml leupeptin, 1 mg/l pepstatin, and 10 μg/ml aprostatin). Lysates were exposed to one cycle of freezing and thawing,
incubated for 30 min on ice and centrifuged at 15000×g for 30 min. Nuclear fractions were isolated as previously described (Mackenzie et al. 2002b). The protein content in the samples was measured according to Bradford (1976).

Aliquots of total or nuclear fractions containing 25–50 μg of protein were separated by SDS-PAGE [10–12.5% (w/v) acrylamide] (Laemmli 1970) blotted onto PVDF membranes and subsequently probed with the different primary antibodies followed by a secondary horseradish peroxidase-conjugated antibody. The bands were visualized by enhanced chemiluminescence (ECL plus, Amersham Biosciences, Piscataway, NJ, USA) and measured using a Phosphoimager 840 (GE Healthcare).

Electrophoretic Mobility Shift Assay

Nuclear fractions were isolated as previously described (Mackenzie et al. 2006a). For the EMSA, the oligonucleotides containing the consensus sequences for NF-κB or OCT-1 were end labeled with \(^{32}\)P-ATP using T4 polynucleotide kinase and purified using Chroma Spin-10 columns. Samples were incubated with the corresponding labeled oligonucleotide (20000–30000 cpm) for 20 min at room temperature in 1× binding buffer [5×: 50 mM Tris–HCl buffer, pH 7.5, containing 20% (v/v) glycerol, 5 mM MgCl₂, 2.5 mM EDTA, 2.5 mM DTT, 250 mM NaCl, and 0.25 mg/ml poly(dI-dC)]. The products were separated by electrophoresis in a 6% (w/v) non-denaturing polyacrylamide gel using 0.5× TBE (45 mM Tris/borate, 1 mM EDTA) as the running buffer. The gels were dried and radioactivity was measured using a Phosphoimager 840.

Evaluation of Apoptosis

Assay of Caspase-3 Activity

Caspase-3 activity was determined in cell lysates after incubating IMR-32 cells for 12, 24, 36, and 48 h, and cortical neurons for 12, 18, and 24 h, in control or chelated media containing 1.5 or 15 μM Zn. Caspase-3 activity was measured using the ApoAlert-Caspase-3 Assay following manufacturer’s protocol.

Caspase-3 activity was also determined measuring the proteolytic processing of full-length PARP [poly(ADP-ribose) polymerase] by Western blot analysis as described above.

TUNEL Assay

Apoptotic nuclei were identified in IMR-32 and cortical neurons after 48 and 24 h of incubation, respectively, in control or chelated media containing 1.5 or 15 μM Zn by evaluating DNA breaks with the TUNEL assay, either by confocal fluorescence microscopy (laser scanning confocal microscope Nikon D-ECLIPSE C1si) or by flow cytometry. For analysis by flow cytometry, control and treated cells were collected in their culture medium, pelleted by centrifugation at 300×g for 5 min and rinsed with PBS. Cells were fixed with 1% (w/v) paraformaldehyde in PBS and after 15 min on ice, washed, and resuspended in PBS. Seventy percent (v/v) cold ethanol was added to the suspension and placed overnight at −20°C. After centrifugation, the cells were washed and resuspended in 50 μl of the staining solution containing the terminal deoxynucleotidyl transferase enzyme and FITC-dUTP and incubated for 2 h at 37°C. At the end of the incubation time, the cells were washed, resuspended in 0.5 ml of the propidium iodide/RNase solution, and analyzed on a Becton-Dickinson flow cytometer.

Translocation of Phosphatidyl Serine to the Outer Layer of the Plasma Membrane

The translocation of phosphatidyl serine was evaluated measuring the cell binding of merocyanine (Verstraeten et al. 2004). IMR-32 (1 × 10⁵ cells) cells suspended in 0.3 ml of 50 mM Hepes buffer (pH 7.4) containing 125 mM KCl were added with 10 μM merocyanine, and incubated at room temperature for 10 min in the darkness. Merocyanine binding to phosphatidylserine was evaluated as the increase in the fluorescence emission at 578 nm (excitation, 555 nm) using an SFM-25 Kontron fluorometer (Kontron Instruments s.p.A, Milan, Italy). To evaluate DNA content, cells were disrupted by a 30-min incubation in presence of 0.1% (v/v) Igepal, and incubated for further 20 min with 50 μM propidium iodide. The fluorescence at 590 nm (excitation, 538 nm) was measured, and results are expressed as the merocyanine/propidium iodide fluorescence ratio.

Cytochrome c Release

To characterize the release of cytochrome c, 18 h after incubating IMR-32 cells in the presence of control or chelated media containing 1.5 or 15 μM Zn, cells were harvested in their culture medium, pelleted by centrifugation at 300×g for 5 min, rinsed with PBS, and resuspended in 250 μl of 20 mM Hepes buffer (pH 7.5) containing 10 mM KCl, 1.5 mM MgCl₂, 1 mM EDTA, 1 mM EGTA, 1 mM dithiothreitol, 250 mM sucrose, and a mixture of protease inhibitors. Cells were homogenized with a glass Pyrex homogenizer and centrifuged at 10000×g for 10 min to remove nuclei and unbroken cells. The resulting supernate was subjected to a 10000×g centrifugation for 20 min at 4°C and the pellet, corresponding to the mitochondrial fraction, was solubilized in 50 μl of 10 mM Tris buffer (pH 8), 0.05% (v/v) Igepal, and 5 mM CaCl₂. The resulting supernatant was transferred to a new tube and centrifuged at 10000×g for 10 min. The pellet was then solubilized in 50 μl of 10 mM Tris buffer (pH 8), and the resulting supernatant was assayed for cytochrome c content.
supernate was centrifuged at 100,000×g for 1 h at 4°C to obtain the cytosolic fraction. Cytochrome c and Bad content in mitochondrial and cytosolic fractions were measured by Western blot analysis as described above.

Statistical Analysis

One-way analysis of variance (ANOVA) with subsequent post-hoc comparisons by Scheffe were performed using Statview 5.0 (SAS Institute, Cary, NC, USA). A P-value <0.05 was considered statistically significant. Values are given as means ± SEM.

Results

A Decrease in Zn Availability Decreases IMR-32 Neuroblastoma Cell Viability

To investigate the requirement of Zn for neuronal proliferation, the human neuroblastoma cell line IMR-32 was used as a model. These cells can be differentiated in vitro to a phenotype of cortical neurons, thus they can complement observations made using primary cortical neurons.

The effects of low extracellular Zn concentrations on IMR-32 cell viability were initially investigated (Fig. 1). IMR-32 cells were incubated for 12–48 h in control medium or in chelated medium containing 1.5 or 15 μM Zn, and cell viability was evaluated measuring the intracellular content of ATP. Cell viability was similar among the groups after 12 h of incubation, but at 24 and 48 h, cell viability was significantly lower (P < 0.05) for the cells incubated in the Zn deficient (1.5 μM Zn) media compared to controls (15 and 35%, respectively). The finding that cell viability was similar for the groups incubated in chelated medium supplemented with 15 μM Zn and in control medium at all the time points investigated, suggesting that the low cell viability found in the 1.5 Zn group can be attributed to the decrease in Zn availability. Under similar experimental conditions, we previously described that the incubation of cells in media containing 1.5 μM Zn leads to a rapid decrease in, labile, as well as total, Zn concentrations (Mackenzie et al. 2002a, 2006a).

A Decrease in Zn Availability Inhibits IMR-32 Neuroblastoma Cell Proliferation

The observed decrease in IMR-32 cell viability in cells exposed to medium containing 1.5 μM Zn could have occurred secondary to a requirement of Zn for cell proliferation and/or to the occurrence of apoptotic cell death. To investigate this, the progression of the cell cycle was investigated. Cell cycle in IMR-32 cells occurs within 24 h, and the entry into the S phase occurs between 12 and 16 h (30 and 35% of cells in S phase, respectively) (data not shown). The effects of varying extracellular Zn concentrations on the distribution of cells in S, G0–G1, and G2–M phases were characterized after 0, 9, 12, 16, and 24 h, respectively. At time 0, in synchronized cells, approximately 75% of the cells were in G0–G1 phase of the cell cycle. At 12 h, there were no significant differences in the proportion of cells in G1, S, and G2–M phases among the groups. At 16 h (Fig. 2a), cells incubated in the 1.5 μM Zn media exhibited a significant (P < 0.05) growth arrest in G1 phase (74 ± 1, 61 ± 1, and 59 ± 2% cells in G0–G1 phase for the 1.5, 15 μM Zn, and control cells, respectively) (Fig. 2a).

Key proteins involved in the control of the G1/S checkpoint of the cell cycle were subsequently investigated. In synchronized IMR-32 cells, the incubation in control medium for 9 h was associated with a marked reduction in p21; a protein that inhibits G1–S transition, p21 levels remained low for the subsequent 15 h. In the cells incubated in medium containing 1.5 μM Zn, p21 levels remained relatively high throughout the 24 h culture period. Consistent with the above, the levels of p53, a protein that positively regulates p21, were higher in IMR-32 cells incubated in 1.5 μM Zn medium compared to controls at all time points (6–24 h). In concordance with the arrest of the cell cycle in G1 phase, Zn deficiency resulted in a reduction in cyclin D1 and E protein levels after 6 h of incubation, the levels remained low throughout the 24 h study period (Fig. 2b). The mitogen-activated kinase (MAPK) ERK has an important role in the regulation of cell proliferation. The incubation of IMR-32 cells in 1.5 μM Zn medium was
associated with a marked decrease in ERK phosphorylation that occurred throughout the 24 h period (Fig. 2b).

A Decrease in Zn Availability Induces Apoptotic Death in IMR-32 Neuroblastoma Cells and Primary Cortical Neurons

The involvement of apoptotic cell death in the decreased cell viability associated with Zn deficiency was assessed by evaluating different steps in the apoptotic cascade, the activation of the effector caspase-3 (caspase-3 activity and degradation of its substrate, PARP), DNA fragmentation, and the translocation of phosphatidyl serine to the outer cell membrane leaflet.

For IMR-32 cells, their incubation in low Zn medium led to a time-dependent activation of caspase-3 (Fig. 3a, b). The increased activation of caspase-3, measured as enzyme activity, was significant after incubating the cells for 24 h in the 1.5 μM Zn medium compared to the control cells. Caspase-3 activity remained elevated for the subsequent 24 h (threefold higher than in controls) (Fig. 3a). When cells were simultaneously incubated for 24 h in 1.5 μM Zn medium supplemented with the antioxidant α-lipoic acid (0.5 mM), the activity of caspase-3 remained high (100 ± 8, 176 ± 6, and 184 ± 20, for cells incubated in the following media: control, 1.5 μM Zn, and 1.5 μM Zn added with α-lipoic acid, respectively). The activation of caspase-3 in the 1.5 μM Zn group was also observed when
Fig. 3  Zn deficiency induces apoptotic cell death in IMR-32 cells. Cells were incubated in control non-chelated medium (C) or chelated media containing 1.5 (1.5 Zn) or 15 µM (15 Zn) Zn for the different periods of time. a Caspase-3 activity after 12, 24, 36, and 48 h of incubation, in the different experimental conditions. b Western blot for full-length and cleaved-PARP in IMR-32 cells incubated in the corresponding media. One representative image out of three independent experiments is shown. c Western blots for Bad and cytochrome c (cyt c) in cytosolic or mitochondrial fractions after 18 h incubation. Representative images out of two to three independent experiments are shown. DNA nicks were evaluated with the TUNEL staining after 48 h incubation, and the percentage of apoptotic cells was calculated. Results are shown as means ± SEM of four independent experiments. e After 18, 48, and 72 h incubation in the corresponding media, the amount of phosphatidylserine in the outer layer of the cell membrane was evaluated fluorometrically measuring merocyanine 540 (MC) binding, and is expressed relative to the amount of DNA measured as propidium iodide (PI) fluorescence. All results are shown as means ± SEM of two to four independent experiments. * Significantly different compared to C and 15 µM Zn groups (P < 0.01, one-way ANOVA test)
the cleavage of PARP, from the full-length protein (116 kDa) to the cleaved form (85 kDa), was assessed by Western blot (Fig. 3b). In the control and the 15 μM Zn groups, caspase-3 activity was stable over the culture period; values were similar in the two groups (Fig. 3a, b).

The key event in the intrinsic apoptotic pathway is the release of cytochrome c, a component of the mitochondrial respiratory chain, to the cytosol where it activates Apaf-1 leading to the subsequent activation of caspase-9 and caspase-3. In support of the activation of this pathway in Zn deficiency, cytochrome c levels were high in the cytosol of IMR-32 cells incubated in 1.5 μM Zn medium for 18 h (Fig. 3c). Accordingly, the mitochondrial content of the pro-apoptotic protein Bad, involved in the mitochondrial release of cytochrome c, was high in mitochondrial fractions, isolated from the Zn-deficient cells (Fig. 3c). Evaluation of DNA fragmentation, as assessed by TUNEL staining of DNA nicks and subsequent quantitation by cell cytometry, showed 45% of apoptotic cells after 48 h incubation in the 1.5 μM Zn medium (Fig. 3d). The increase in phosphatidylserine translocation was statistically significant after 48 and 72 h incubation in the 1.5 μM Zn medium, compared to cells incubated in control and 15 μM Zn media (Fig. 3e). DNA fragmentation and phosphatidyl serine externalization levels were similar for the control and 15 μM Zn groups (Fig. 3d, e). Similar to the observations with the IMR-32 cells, a decrease in the availability of Zn induced apoptotic cell death in primary cultures of rat cortical neurons (Fig. 4a). Cortical neurons incubated in 1.5 μM Zn medium were characterized by low cell viability after 24 h incubation that was prevented by supplementing the chelated medium with 15 μM Zn. Zn deficiency induced apoptotic cell death in rat cortical neurons, as evidenced by the activation of caspase-3 after 18 h incubation (Fig. 4b); and by high levels of DNA fragmentation after 24 h, as evaluated by confocal microscopy (Fig. 4c). The simultaneous incubation with 0.5 mM z-lipoic acid did not prevent the activation of caspase-3 that occurs when neuronal Zn decreases (Fig. 4b).

Neuronal Signaling Pathways Involved in the Triggering of Apoptosis When the Availability of Zn Decreases

The kinases ERK and Akt participate in the inhibition of the intrinsic apoptotic pathway, by phosphorylating Bad, which prevents its translocation to the mitochondria. We next investigated if the ERK/Akt/Bad pathways were involved in the induction of apoptosis when neuronal Zn decreases. After incubating IMR-32 and cortical neurons in Zn-deficient medium for 9 h, phosphorylated ERK1/2 levels were low, compared to levels in cells incubated in control or 15 μM Zn media (Fig. 5a). At the same time, the levels of phosphorylated Akt were high in cortical neurons and IMR-32 cells incubated in the Zn-deficient medium compared to levels in controls (Fig. 5a). While Akt promotes cell survival through its ability to phosphorylate Bad at Ser136, ERK phosphorylates Bad at Ser112. In IMR-32 cells and rat cortical neurons incubated for 9 h in 1.5 μM Zn medium, the levels of Bad phosphorylated at Ser136 were high, whereas those of Bad phosphorylated at Ser112 were similar to those observed in control cells (Fig. 5b). These results suggest that Akt activation in the Zn-deficient cells cannot counteract the pro-apoptotic consequences of ERK inhibition.

We previously demonstrated that Zn deficiency can decrease the nuclear translocation of transcription factor NF-κB into the nucleus, inhibiting the NF-κB-regulated expression of select genes (Mackenzie et al. 2002b). As expected, NF-κB-DNA binding was lower in nuclear fractions from IMR-32 and rat cortical neurons incubated in 1.5 μM Zn medium for 36 and 24 h, respectively, compared to controls (Fig. 6a). Several proteins (c-IAP-1, XIAP, Bcl-xl, and Bcl-2) involved in the protection of cells from apoptosis are regulated by NF-κB. c-IAP1 acts inhibiting caspases, XIAP exerts its anti-apoptotic action through the interaction with several partners (caspase-3, caspase-9, DIABLO/Smac) and Bcl-xL and Bcl-2 inhibit apoptosis by preventing the opening of the mitochondrial pore. Thus, we next investigated the effects of varying extracellular Zn concentrations on the expression of NF-κB-regulated anti-apoptotic proteins. In IMR-32 and rat cortical neurons incubated in Zn-deficient medium for 36 and 24 h, respectively, a decreased content of c-IAP1, XIAP, Bcl-xL, and Bcl-2 was observed in cortical neurons and of XIAP and Bcl-2 in IMR-32 cells (Fig. 6b). These findings support a role of NF-κB inhibition in the triggering of apoptosis when neuronal Zn availability decreases.

Discussion

In this study, we demonstrate Zn modulates neuronal proliferation and survival. A deficit of Zn can result in an inhibition of neuronal cell proliferation secondary to an arrest at the G0/G1 phase of the cell cycle. In proliferating as well as in quiescent neurons, Zn deficiency can trigger apoptotic neuronal death. Apoptosis occurs through the intrinsic pathway which can be a consequence of ERK inhibition, caspase-3 activation, and the down-regulation of NF-κB-dependent anti-apoptotic genes.

Under the conditions employed in this study, Zn deficiency caused an arrest in IMR-32 cell cycle progression at the G0/G1 phase. A similar finding was observed in HepG2 human hepatoblastoma cells (Wong et al. 2007). Accordingly, male rats fed a Zn-deficient diet have a decreased...
number of proliferating stem cells in the subgranular zone and granular cell layer of the dentate gyrus compared to controls (Corniola et al. 2008). A key aspect in the cell cycle regulation is the MAPK pathway, Raf/MEK/ERK1/2 that can be activated by mitogenic signals, cytokines, and growth factors (Steelman et al. 2004). The ERK signaling pathway is involved in the integration of environmental signals (nutrients, mitogens, and cytostatic factors) that lead to DNA replication during the S phase, and in relaying the information to the cell cycle control system. ERK activation throughout G1 is required for the cell entry into S phase (Yamamoto et al. 2006). Zn deficiency caused a rapid inhibition of ERK1/2 phosphorylation in IMR-32 cells supporting a role of this pathway in the associated cell
Cycle arrest. Accordingly, the ERK inhibitor (PD 98059) causes a significant decrease in IMR-32 cell viability within 24 h (Zago et al. 2005), supporting a role of ERK in IMR-32 cell proliferation. A down-regulation of cyclins D and E was also observed in Zn-deficient cells. Major components of the regulation of G1 to phase S transition are the cyclin-dependent kinases (Cdks) and their regulatory cyclins (Sherr 2000). During the G1 phase, cyclin D assembles with its catalytic partners Cdk4 or Cdk6 which in turn phosphorylate members of the retinoblastoma family of proteins. Cyclin E-Cdk2 activates later in G1 phase. Because ERK upregulates cyclins D1 and E, the findings of decreased cyclins D1 and E levels in the Zn-deficient IMR-32 cells are consistent with the observed ERK inhibition and Go/G1 arrest. Zn deficiency also increased p53 protein levels which can occur as a consequence of Zn deficiency-induced activation of the MAPK (JNK/p38)-AP-1 signaling pathway (Zago et al. 2005). As previously proposed (Corniola et al. 2008), Zn deficiency-induced p53 upregulation can underlie an increased expression of downstream target genes which could contribute to the triggering of cell cycle arrest (Cdk inhibitor p21, and others) and apoptosis (Corniola et al. 2008; Miyashita and Reed 1995; Miyashita et al. 1994).

Other molecular events may contribute to the decreased cell proliferation in the Zn-deficient neuronal cells. It has been proposed that the regulation of the cell cycle by Zn could be secondary to its requirement for enzyme systems that influence cell division and proliferation (Clegg et al. 1989; MacDonald 2000; Beyersmann and Haase 2001), and to the impaired tubulin polymerization that is associated with Zn deficiency (Oteiza et al. 1990). An altered neuronal microtubule assembly (Mackenzie et al. 2002b) could impair the formation of the mitotic spindle affecting mitosis.

Zn deficiency induced apoptotic cell death in IMR-32 cells and in cortical neurons as evidenced by caspase-3.
activation, phosphatidyserine externalization, and DNA fragmentation. Apoptotic cell death is consistently associated with Zn deficiency in animal as well as cell models (Fraker and Telford 1997; Jankowski-Hennig et al. 2000; Duffy et al. 2001; Truong-Tran et al. 2001; Clegg et al. 2005; Fraker 2005; Corniola et al. 2008). Different hypothesis have been reported for the induction of apoptosis when cellular Zn decreases. These include the action of Zn as a direct caspase regulator, alterations in receptor tyrosine kinase signaling, mitochondrial dysfunction, and increased oxidative stress (Truong-Tran et al. 2001; Clegg et al. 2005).

We previously observed that the increase in cell oxidants and the associated activation of JNK/p38/AP-1 is not
a critical trigger for the neuronal cell death that occurs following the induction of Zn deficiency in IMR-32 cells (Zago et al. 2005). Although treatment with α-lipoic acid completely prevents the increase in oxidants in IMR-32 cells (Mackenzie et al. 2006b), it did not prevent caspase-3 activation in IMR-32 cells and cortical neurons. This provides further evidence that oxidative stress does not play a major role on Zn deficiency-induced neuronal apoptosis.

In this study, we observed a hypophosphorylation of ERK1/2 and an increased phosphorylation of Akt in association with Zn deficiency in IMR-32 and rat cortical neurons. Concomitantly, the levels of Bad phosphorylated at Ser-136 were high and no changes in the levels of Bad phosphorylated at Ser-112 were observed in association with Zn deficiency in either cell type, compared to controls. The Bcl-2 family of proteins includes anti-apoptotic and pro-apoptotic proteins, Bcl-2, and Bcl-X1 mediate cell survival while Bad and Bax induce cell death. Bad resides in the cytosol, as an inactive complex with the chaperone 14-3-3 proteins, when it is phosphorylated in at least three serine residues (Ser-112, 136, and 155) (Zha et al. 1996). These sites are phosphorylated by different kinases. The phosphorylation of Bad at Ser-112 is mediated by ERK (Zhu et al. 2002) and protein kinase A (Yang et al. 2005). Ser-136 is phosphorylated by Akt (Blume-Jensen et al. 1998), and Ser-155 by protein kinase A (Virdee et al. 2000). Recent evidence shows that phosphorylation at the three different sites are required to prevent Bad translocation to the mitochondria (Wang et al. 2007). Thus, while Akt can be activated secondarily to Zn deficiency, the inhibition of ERK can drive Bad translocation, and the subsequent mitochondrial release of cytochrome c and caspase-3 activation. The activation of caspase-3 observed in IMR-32 cells and cortical neurons can also be due to the direct inhibitory action of Zn on the enzyme. In this regard, caspase-3 is rapidly and directly activated by Zn chelation in HeLa cells without a requirement for an upstream event (Chimienti et al. 2001).

The inhibition of the pro-survival transcription factor NF-κB could also contribute to Zn deficiency-induced neuronal apoptosis. The inhibition of NF-κB with the proteasome inhibitor lactacystin causes apoptosis in IMR-32 cells (Mackenzie et al. 2002a, 2006a). We previously described that Zn deficiency alters NF-κB modulation and inhibits NF-κB-dependent gene expression in IMR-32 cells (Mackenzie et al. 2002a, 2006a). Consistently, in IMR-32 cells and rat cortical neurons incubated in Zn-deficient media, we observed a low NF-κB-DNA binding in nuclear fractions and a decreased expression of the anti-apoptotic proteins c-IAP1, XIAP, Bcl-xL, and Bcl-2.

In summary, Zn modulates a number of pro-survival (ERK, Akt, and NF-κB) and pro-apoptotic (JNK and p53) pathways, and the final consequence of a deficit of Zn will be the result of the balance between the activation/inhibition of all of them. In neurons (Fig. 7), a low Zn availability results in decreased neuronal proliferation, due to an arrest of the cell cycle in G0/G1 phase, and the triggering of neuronal apoptosis. The inhibition of ERK is an important player in both processes. Caspase-3 activation, as a direct consequence of neuronal Zn decrease, is a central event in the triggering of apoptosis. However, alterations in cell signals (NF-κB, p53) that regulate the expression of pro-survival or pro-apoptotic genes can also contribute to Zn deficiency-induced apoptosis. Given the relevance of neuronal proliferation and apoptosis in the developmental shaping of the brain, a suboptimal Zn nutrition during embryogenesis and early postnatal life can be a significant risk to the conceptus.

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