Research Article

Nickel Enhances Zinc-Induced Neuronal Cell Death by Priming the Endoplasmic Reticulum Stress Response

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Trace metals such as zinc (Zn), copper (Cu), and nickel (Ni) play important roles in various physiological functions such as immunity, cell division, and protein synthesis in a wide variety of species. However, excessive amounts of these trace metals cause disorders in various tissues of the central nervous system, respiratory system, and other vital organs. Our previous analysis focusing on neurotoxicity resulting from interactions between Zn and Cu revealed that Cu^{2+} markedly enhances Zn^{2+} -induced neuronal cell death by activating oxidative stress and the endoplasmic reticulum (ER) stress response. However, neurotoxicity arising from interactions between zinc and metals other than copper has not been examined. Thus, in the current study, we examined the effect of Ni²⁺ on Zn²⁺-induced neurotoxicity. Initially, we found that nontoxic concentrations (0–60 μ M) of Ni²⁺ enhance Zn²⁺-induced neurotoxicity in an immortalized hypothalamic neuronal cell line (GT1-7) in a dose-dependent manner. Next, we analyzed the mechanism enhancing neuronal cell death, focusing on the ER stress response. Our results revealed that Ni²⁺ treatment significantly primed the Zn²⁺-induced ER stress response, especially expression of the CCAAT-enhancer-binding protein homologous protein (CHOP). Finally, we examined the effect of carnosine (an endogenous peptide) on Ni²⁺/Zn²⁺-induced neurotoxicity and found that carnosine attenuated Ni²⁺/Zn²⁺-induced neuronal cell death and ER stress response. Thus, compounds that decrease the ER stress response, such as carnosine, may be beneficial for neurological diseases.

1. Introduction

In many organisms, trace metals such as zinc (Zn), copper (Cu), and nickel (Ni) play important roles in various physiological functions such as immunity, cell division, and protein synthesis. Indeed, these trace metals are well-known cofactors for hundreds of enzyme proteins [1]. Thus, deficiency of these trace metals causes immune system dysfunction, physical development retardation, dwarfism, learning disabilities, and taste and olfaction disorders in humans [2, 3]. However, excessive amounts of these trace metals cause disorders of various organs, such as the central nervous system and respiratory system. In particular, disorders involving excessive amounts of zinc in the central nervous system have been attracting keen attention. Zn binds firmly to certain metalloproteins and enzymes in their steady state but has also been shown to exist in the form of free Zn ions (Zn^{2+}) or loosely bound to proteins in a subset of excitatory neurons [4]. In pathological situations such as stroke or transient global ischemia, interrupted blood flow induces excessive release of Zn via long-lasting membrane depolarization [5]. The released Zn accumulates in neurons and induces neuronal cell death [6, 7]. These findings suggest that Zn is a key modulator of delayed neuronal cell death after ischemic injury, and moreover, Zn neurotoxicity is central to the pathogenesis of

poststroke dementia [8]. Furthermore, Zn is reportedly involved in the progression of Alzheimer's disease [9].

We previously examined molecular mechanisms underlying Zn²⁺-induced neurotoxicity using an immortalized hypothalamic neuronal cell line (GT1-7) and found that Zn²⁺ induced a marked upregulation of endoplasmic reticulum (ER) stress-related genes, especially CCAAT-enhancerbinding protein homologous protein (CHOP) and growth arrest and DNA damage-inducible gene 34 (GADD34), and loss of mitochondrial membrane potential (mitochondrial injury) [10]. These results suggested that Zn²⁺ induces neuronal cell death via the ER stress response and mitochondrial injury. We also identified compounds capable of decreasing Zn²⁺-induced neurotoxicity and neuronal cell death, including carnosine (an endogenous dipeptide), pyruvic acid (an organic acid involved in glycolysis and the tricarboxylic acid cycle), and thioredoxin-albumin fusion protein (HSA-Trx; an antioxidative protein) [10-12].

Other trace metals, such as Cu²⁺, Ni²⁺, iron (Fe²⁺, Fe³⁺), and manganese (Mn²⁺), are also present in the brain and/or cerebrospinal fluid [13, 14]. Recent studies suggest that intracellular Cu²⁺ accumulates in synaptic vesicles and is then released into the synaptic cleft during neuronal excitation, with a reported concentration of Cu²⁺ in the synaptic cleft of 2–15 μ M [15–17]. The translocated Cu²⁺ influences various receptors, including AMPA-type glutamate and GABA receptors, and contributes to the modulation of neuronal excitability [16]. Mn²⁺ is important both for neurotransmitter synthesis and as a component of superoxide dismutase 2 in mitochondria [18]. However, excessive amounts of Mn²⁺ are thought to induce neurotoxicity and cause a Parkinson'slike syndrome [19]. Ni²⁺ also reportedly causes neurotoxicity, reactive oxygen species (ROS) production, and mitochondrial dysfunction in neuronal cells [20]. Considering these reports, other trace metals may interact with Zn²⁺ in brain tissue and synaptic clefts, to act together in neuronal cells. Therefore, we recently examined the effects of various metal ions on Zn²⁺-induced neurotoxicity in GT1-7 cells and found that Cu^{2+} or Ni^{2+} enhanced Zn^{2+} -induced neurotoxicity [21]. Moreover, we found that Cu^{2+} enhanced Zn^{2+} -induced neurotoxicity by activating oxidative stress, the ER stress response, and mitochondrial injury [11, 21, 22]. However, the mechanisms by which Ni²⁺ enhanced Zn²⁺-induced neurotoxicity have not been examined.

Based on the results of our previous study [21], we first examined whether Ni²⁺ enhanced Zn²⁺-induced neurotoxicity using GT1-7 cells in the current investigation. As a result, we found that Ni²⁺ (nontoxic concentrations by single treatment) enhanced Zn²⁺-induced neurotoxicity in GT1-7 cells via a mechanism involving the ER stress response. Furthermore, we examined the effect of carnosine, an endogenous peptide, on neuronal cell death induced by cotreatment with Ni²⁺ and Zn²⁺.

2. Materials and Methods

2.1. Chemicals and Reagents. An antibody against actin (SC-47778) was purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Antibodies against CHOP (#5554) and

goat anti-rabbit IgG (horseradish peroxidase- (HRP-) conjugated, #7074) were purchased from Cell Signaling Technology Japan (Tokyo, Japan). Tauroursodeoxycholic acid (TUDCA) was purchased from Tokyo Chemical Industry (Tokyo, Japan). The Zinc Assay Kit was purchased from Metallogenics Co. Ltd. (Chiba, Japan). HRP-conjugated donkey anti-mouse IgG was purchased from GE Healthcare Japan (Tokyo, Japan). Carnosine, ZnCl₂, NiCl₂, and RIPA buffer (20 mmol/L Tris-HCl (pH 7.4), 0.05 *w/v*% NP-40 substitute, 2.5 mmol/L MgCl₂, and 200 mmol/L NaCl) were purchased from Wako Pure Chemicals (Tokyo, Japan). Protease and phosphatase inhibitors (87786 and 78420), NuPAGE[®] Novex 4–12% Bis-Tris Protein Gel, iBlot[™] Transfer Stack, and SuperSignal[™] West Dura Extended Duration Substrate were from Thermo Fisher Scientific K.K. (Tokyo, Japan).

2.2. Cell Culture. GT1-7 cells, which were provided by Dr. R. Weiner, University of California San Francisco, were grown in Dulbecco's Modified Eagle's Medium/Ham's Nutrient Mixture F-12 supplemented with 10% fetal bovine serum. After trypsin digestion, cells were resuspended in serum-free medium, distributed into culture dishes, and cultured in a humidified incubator (7% CO_2) at 37°C [12].

2.3. Measurement of Cell Viability, Cytotoxicity, and Reactive Oxygen Species (ROS) Levels. Cell viability was measured as described previously [21, 23]. Briefly, dissociated GT1-7 cells were distributed into 96-well culture plates at a density of 3×10^4 cells per well in 200 μ L of culture medium. After 24 h incubation, cells were treated with or without N-acetylcysteine (NAC: 0–250 μ M), carnosine (0–4 mM), or TUDCA (0–100 mM) prior to the addition of NiCl₂ and ZnCl₂ to the medium. After 24 h exposure, cell viability was quantified using CellTiter-Glo[®] 2.0 (Promega Corporation, Madison, WI). Cytotoxicity was quantified using an LDH-GloTM Cytotoxicity Assay kit (Promega Corporation, Madison, WI) after exposure for 24 h.

GT1-7 cells were precultured in black 96-well microplates (3×10^4 cells/well). After incubation for 24 h, cells were incubated with 2 ',7 '-dichlorodihydrofluorescein diacetate (DCFHDA, 100 μ M), an indicator of ROS, in the absence (control) or presence of NiCl₂ and/or ZnCl₂ for 2 h. ROS levels were then measured using a microplate reader (Tecan, Kawasaki, Japan) (Ex: 480 nm; Em: 530 nm).

2.4. Real-Time RT-PCR. Dissociated GT1-7 cells were distributed into 6-well culture plates at a density of 7.5×10^5 cells per well in 1.5 mL of culture medium. After 24 h incubation, cells were treated with or without carnosine (0–4 mM) prior to the addition of NiCl₂ and ZnCl₂ to the medium. After 4 h exposure, total RNA was extracted using an RNeasy kit (QIAGEN, Hilden, Germany) according to the manufacturer's protocol. Samples were reverse-transcribed using a PrimeScript[®] First Strand cDNA Synthesis Kit (Takara Bio, Otsu, Japan). Synthesized cDNA was used in real-time RT-PCR experiments with Thunderbird[®] SYBR qPCR Mix (Toyobo, Osaka, Japan) and analyzed with a CFX96[™] Real-Time System and CFX Manager[™] software (Bio-Rad, Hercules, CA). Specificity was confirmed by electrophoretic

Name	Forward	Reverse
Gapdh	AACTTTGGCATTGTGGAAGG	ACACATTGGGGGTAGGAACA
Chop	CCACCACACCTGAAAGCAGAA	AGGTGAAAGGCAGGGACTCA
Gadd34	TCCCTCATGGGGAGACTGAA	AGCTGTGCGTTCCATTTCCT
Grp78	TTCAGCCAATTATCAGCAAACTCT	TTTTCTGATGTATCCTCTTCACCAGT
Edem	CTACCTGCGAAGAGGCCG	GTTCATGAGCTGCCCACTGA
Atf4	GGGTTCTGTCTTCCACTCCA	AAGCAGCAGAGTCAGGCTTTC
Grp94	AAGAATGAAGGAAAAACAGGACAAAA	CAAATGGAGAAGATTCCGCC
Pdi	GGATTGCACTGCCAACACAA	AGCTGGTCCTGCTTGTTTCT
Hif1a	CAAGATCTCGGCGAAGCAA	GGTGAGCCTCATAACAGAAGCTTT
Nrf2	TGGAGAACATTGTCGAGCTG	CTGAGCCGCCTTTTCAGTAG
Ho1	GAACCCAGTCTATGCCCCAC	GGCGTGCAAGGGATGATTTC
Gstm1	GCTCATCATGCTCTGTTACAACC	GCCCAGGAACTCAGAGTAGAG

TABLE 1: Sequences of primers.

analysis of reaction products and template- or reverse transcriptase-free controls. To normalize the amount of total RNA present in each reaction, glyceraldehyde-3-phosphate dehydrogenase (*Gapdh*) mRNA was used as an internal standard. Primers were designed using Primer-BLAST. Primer sequences are listed in Table 1.

2.5. Western Blotting Analysis. Zn-induced expression of CHOP and actin was assessed by Western blotting analysis. GT1-7 cells grown in 6-well culture plates $(7.5 \times 10^5 \text{ cells})$ per well) were lysed with RIPA buffer containing protease and phosphatase inhibitors. Protein concentrations were measured using a Bradford Protein Assay Kit (Takara Bio). Samples were applied to a NuPAGE® Novex 4-12% Bis-Tris Protein Gel and electrophoresed at a constant voltage of 180 V, and then, proteins were transferred to an iBlot[™] Transfer Stack (PVDF membranes) using the iBlot® 7-Minute Blotting System (Thermo Fisher Scientific K.K.). Membranes were blocked with 5% nonfat dry milk at room temperature for 1 h and then incubated with rabbit anti-CHOP antibody (1:1000 dilution) or mouse anti-actin antibody (1:1000 dilution) in 5% BSA, 1x Tris-buffered saline (TBS), and 0.1% Tween-20 overnight. The following day, membranes were incubated with goat anti-rabbit IgG (1:2000 dilution) or donkey anti-mouse IgG (1:4000 dilution) HRP-conjugated secondary antibodies in 1x TBS containing 0.1% Tween-20 for 1 h, and finally, bands were visualized using SuperSignal[™] West Dura Extended Duration Substrate. Band intensities were quantitated using ImageJ software (version 1.39u), and the band intensity of each protein was determined and normalized with respect to actin intensity.

2.6. Statistical Analysis. All data are expressed as mean \pm S. E.M. Significant differences among groups were examined using a one-way of analysis of variance (ANOVA) followed by Tukey's multiple comparison. SPSS 24 software was used for all statistical analyses. A probability value of P < 0.05 was considered to indicate statistical significance.

3. Results

3.1. Ni^{2+} Enhanced Zn^{2+} -Induced Neuronal Cell Death. We previously examined the effect of various metal ions on Zn²⁺-induced neurotoxicity in GT1-7 cells and revealed that sublethal concentrations of Cu²⁺ markedly enhanced Zn²⁺induced neurotoxicity [21]. We also discovered that Ni²⁺ enhances Zn²⁺-induced neurotoxicity, but its mechanism was not determined. In this study, we therefore examined the effect of Ni²⁺ on Zn²⁺-induced neurotoxicity in GT1-7 cells. As shown in Figure 1(a), Zn²⁺ induced neurotoxicity in GT1-7 cells in a dose-dependent manner. The viability of cells exposed to 20, 30, or 40 μ M of Zn²⁺ was 98.4% ± 0.6%, 79.8% ± 2.0%, and 48.0% ± 2.7% (mean ± S.E.M., n = 4) of the control, respectively. In contrast, the indicated concentrations of Ni²⁺ (0–40 μ M) in Figure 1(b) did not reduce the viability of GT1-7 cells.

The effect of Ni²⁺ on Zn²⁺-induced neurotoxicity in GT1-7 cells is shown in Figure 1(c). At a constant Zn²⁺ concentration of 25 μ M, Ni²⁺ enhanced Zn²⁺-induced neurotoxicity in GT1-7 cells in a dose-dependent manner within the tested Ni²⁺ concentration range (0–60 μ M), whereas Ni²⁺ treatment alone did not cause any neurotoxicity. The viability of cells exposed to 0, 20, 40, or 60 μ M of Ni²⁺ in the presence of 25 μ M Zn²⁺ was 95.9% ± 0.8%, 72.0% ± 1.5%, 39.7% ± 1.6%, and 7.1% ± 3.2% (mean ± S.E.M., *n* = 4) of the control, respectively. We then measured LDH release from GT1-7 cells to monitor cytotoxicity. As shown in Figure 1(e), Ni²⁺ enhanced Zn²⁺-induced LDH release from GT1-7 cells in a dose-dependent manner within the tested Ni²⁺ concentration range (0–60 μ M). Based on the results shown in Figure 1, Ni²⁺ exacerbated Zn²⁺-induced neuronal cell death.

3.2. Activation of the ER Stress Response by Cotreatment with NiCl₂ and ZnCl₂. We previously showed that Zn²⁺ and Cu²⁺ increase the expression of ER stress-related genes in GT1-7 cells and that Cu²⁺ enhances Zn²⁺ induced neurotoxicity by activating the ER stress response [21]. Thus, we monitored whether Ni²⁺ primes Zn²⁺-induced ER stressrelated gene expression using real-time RT-PCR. ZnCl₂



FIGURE 1: GT1-7 cells (96-well culture plates at a density of 3×10^4 cells per well) were incubated with the indicated concentrations (μ M) of ZnCl₂ (a) or NiCl₂ (b) for 24 h. GT1-7 cells (96-well culture plates at a density of 3×10^4 cells per well) were incubated with the indicated concentrations (μ M) of NiCl₂ in the absence (control) or presence of ZnCl₂ (25 μ M) for 24 h (c). Cell viability was determined using CellTiter-Glo[®] 2.0. Values represent mean ± S.E.M. *P* values are described in the figure when *P* < 0.05 (black: vs. control, red: vs. ZnCl₂ alone).



FIGURE 2: GT1-7 cells (6-well culture plates at a density of 7.5×10^5 cells per well) were incubated with NiCl₂ (Ni, 40 μ M) in the absence (control) or presence of ZnCl₂ (Zn, 25 μ M) for 4 h. Total RNA was extracted from GT1-7 cells and subjected to real-time RT-PCR using primer sets specific for each gene. Values were normalized to *Gapdh* and expressed relative to the control. Values represent mean ± S.E.M. *P* values are described in the figure when *P* < 0.05 (black: vs. control, red: vs. ZnCl₂ alone).

(25 μ M) treatment induced the expression of *Chop* and activating transcription factor 4 (*Atf4*) mRNA (Figure 2). Moreover, Ni²⁺ treatment primed the expression of *Chop*, *Gadd34*, and *Atf4*; in particular, the relative expression of *Chop* was most significantly increased by cotreatment with Ni²⁺ and Zn²⁺. The relative expression of *Chop* after cotreatment with Ni²⁺ and Zn²⁺ was 28.6 ± 0.3 – fold (mean ± S.E.M., *n* = 3), which was significantly increased compared with Zn²⁺

alone $(2.3 \pm 0.1 - \text{fold})$. In contrast, other ER stress-related genes including glucose-regulated protein 78 (*Grp78*), ER degradation-enhancing α -mannosidase-like protein (*Edem*), glucose-regulated protein 94 (*GrpP94*), and protein disulfide isomerase (*Pdi*) were not increased in this experimental condition. Ni²⁺ treatment alone did not increase expression of these genes (Figure 2). Next, we used Western blotting analysis to quantify the amount of CHOP protein.



FIGURE 3: GT1-7 cells (6-well culture plates at a density of 7.5×10^5 cells per well) were incubated with NiCl₂ (Ni, 40 μ M) in the absence (control) or presence of ZnCl₂ (Zn, 25 μ M) for 7 h. Whole-cell extracts were analyzed by immunoblotting with an antibody against CHOP or actin (a). Band intensity was determined using ImageJ software (b). GT1-7 cells (96-well culture plates at a density of 3×10^4 cells per well) were pretreated with the indicated concentrations (μ M) of TUDCA just before Ni²⁺/Zn²⁺ treatment. Next, GT1-7 cells were incubated in the absence (control) or presence of NiCl₂ (20 or 40 μ M) and ZnCl₂ (25 μ M) for 24 h (c, d). GT1-7 cells (96-well culture plates at a density of 3×10^4 cells per well) were treated with the indicated concentrations of TUDCA for 24 h (e). Cell viability was determined using CellTiter-Glo[®] 2.0 (c–e). Values represent mean ± S.E.M. *P* values are described in the figure when *P* < 0.05: (black: control, red: vs. ZnCl₂ alone (b)) or (black: vs. control, red: vs. Zn(25)/Ni(20 or 40) (c, d)).

As shown in Figure 3(a), we found significantly increased amounts of CHOP protein after cotreatment with Ni^{2+} and Zn^{2+} (78.2 ± 1.3 – fold), compared with Zn^{2+} alone (10.3 ± 0.2 – fold). Ni²⁺ treatment alone did not increase the expression of CHOP protein (Figure 3(b)). Furthermore,

we examined whether the ER stress inhibitor, TUDCA, attenuates Ni^{2+}/Zn^{2+} -induced neurotoxicity. As shown in Figures 3(c) and 3(d), TUDCA significantly attenuated Ni^{2+}/Zn^{2+} -induced neurotoxicity in GT1-7 cells in a dose-dependent manner. For example, the viability of cells

exposed to Ni²⁺/Zn²⁺ (20 μ M/25 μ M) or Ni²⁺/Zn²⁺ plus TUDCA (100 μ M) was 58.5% ± 0.7% or 80.3% ± 1.0% (mean ± S.E.M., *n* = 4), respectively. Treatment with TUDCA alone did not affect the viability of GT1-7 cells (Figure 3(e)). These results suggest that Ni²⁺ enhanced Zn²⁺-induced neurotoxicity by priming the ER stress response.

3.3. Activation of Oxidative Stress by Cotreatment with NiCl, and ZnCl₂. We next examined the involvement of oxidative stress on Ni²⁺/Zn²⁺-induced neurotoxicity or the ER stress response. As shown in Supplementary Figure S1A, cotreatment with Ni²⁺ and Zn²⁺ induced ROS production. In contrast, Ni²⁺ or Zn²⁺ treatment did not always induce ROS production. Moreover, Ni²⁺/Zn²⁺ treatment induced the expression of oxidative stress-related genes, as indicated by increases in hypoxia-inducible factor (Hif)1 α , nuclear factor-erythroid 2-related factor 2 (Nrf2), heme oxygenase 1 (Ho1), and glutathione S-transferase m1 (Gstm1) mRNA levels. In contrast, Ni^{2+} or Zn^{2+} treatment did not increase expression of these genes (Supplementary Figure S1B). Thus, we examined the effect of N-acetylcysteine, an antioxidative compound, on Ni²⁺/Zn²⁺-induced neurotoxicity. As shown in Supplementary Figure S1C, N-acetylcysteine significantly attenuated Ni²⁺/Zn²⁺-induced neurotoxicity in GT1-7 cells in a dose-dependent manner. The viability of cells exposed to Ni²⁺/Zn²⁺ (40 μ M/25 μ M) or Ni²⁺/Zn²⁺ plus N-acetylcysteine (250 μ M) was 20.8% ± 6.5% or 71.5% ± 1.7% (mean ± S.E.M., n = 4), respectively (Supplementary Figure S1C). Treatment with only N-acetylcysteine did not affect the viability of GT1-7 cells (Supplementary Figure S1D). Furthermore, N-acetylcysteine (250 μ M) reduced the amount of CHOP protein induced by Ni^{2+}/Zn^{2+} (40 μ M/25 μ M) treatment (Supplementary Figure S1E and S1F). These results suggested that Ni²⁺/Zn²⁺-induced ER stress responses were mediated by upregulating ROS production.

3.4. Effect of Carnosine on Cu^{2+}/Zn^{2+} -Induced Neurotoxicity and the ER Stress Response. Carnosine (β -alanyl-L-histidine) is a small dipeptide with numerous activities, including antioxidant effects, proton buffering capacity, and inhibitory effects on protein carbonylation [24, 25]. Although we found previously that carnosine protected against Zn²⁺-induced neurotoxicity [10], we have not studied the effect of carnosine on neuronal cell death induced by cotreatment with Zn^{2+} and other metals. Thus, we examined the effect of carnosine on Ni²⁺/Zn²⁺-induced neurotoxicity and the ER stress response in GT1-7 cells. The concentration of the chosen carnosine was according to a previous report [10]. As shown in Figures 4(a) and 4(b), cotreatment of GT1-7 cells with NiCl₂ (20 or 40 μ M) and ZnCl₂ (25 μ M) induced neurotoxicity with cell viabilities of $48.5\% \pm 0.7\%$ and $25.5\% \pm 0.9\%$ (mean \pm S.E.M., n = 4) of the control, respectively. In contrast, carnosine significantly attenuated Ni²⁺/Zn²⁺-induced neurotoxicity in GT1-7 cells in a dose-dependent manner. The viability of cells exposed to Ni²⁺/Zn²⁺ (20 μ M/25 μ M) plus carnosine (1, 2, and 4 mM) was $55.2\% \pm 0.7\%$, $61.6\% \pm$ 2.3%, and 70.0% \pm 3.3% (mean \pm S.E.M., n = 4), respectively, (Figure 4(a)). The viability of cells exposed to Ni^{2+}/ Zn^{2+} (40 μ M/25 μ M) plus carnosine (1, 2, and 4 mM) was

 $32.0\% \pm 4.9\%$, $39.9\% \pm 2.2\%$, and $51.1\% \pm 1.1\%$ (mean ± S.E.M., n = 4), respectively, (Figure 4(b)). Treatment with carnosine alone did not affect the viability of GT1-7 cells (Figure 4(c)). Next, we examined the effect of carnosine on Ni²⁺/Zn²⁺-induced ER stress responses. As shown in Figure 5, cotreatment of GT1-7 cells with Ni² and Zn²⁺ increased the expression of Chop, Gadd34, and Atf4. In contrast, carnosine treatment significantly decreased the expression of these genes in a dose-dependent manner (Figure 5). Furthermore, the amount of CHOP protein observed after cotreatment with Ni^{2+} and Zn^{2+} was reduced by carnosine treatment (Figures 6(a) and 6(b)). These results suggest that carnosine significantly attenuated Ni² and Zn²⁺-induced neuronal cell death by decreasing the ER stress response. Moreover, carnosine may decrease not just neuronal cell death induced by Zn²⁺ alone but also neuronal cell death induced by Zn^{2+} and other metals.

4. Discussion

Zn is the second most abundant trace metal and has essential roles in many physiological functions such as the immune system, cell cycle, DNA replication, and protein synthesis. Moreover, Zn is a well-known cofactor for over 300 enzymes and metalloproteins [1]. Deprivation of Zn in humans causes atrophy, learning disorders, delayed physical development, taste and olfaction disorders, and immune system diseases [2, 3]. Thus, Zn supplementation reportedly exerts therapeutic effects against various diseases such as cirrhosis, ulcerative colitis, and asthma [26–28]. In the brain, Zn^{2+} accumulates in the synaptic vesicles of excitatory synapses and is released during neuronal excitation [29, 30]. Although Zn has essential roles in the brain and neuroprotective roles in normal neuronal function, excessively high concentrations of Zn are neurotoxic [6–8]. Although the amount of Zn^{2+} released from synaptic vesicles still needs confirmation, several studies estimate the concentration of Zn^{2+} in the synaptic cleft to be 1–100 μ M [31–34]. Therefore, Zn²⁺ may act on neuronal cells at the concentration used in this study (25 μ M). Ni, a component of enzymes such as urease and hydrogenase, is a heavy metal widely used in industrial applications [35]. A previous study showed that nickel concentration in the brain is estimated to be 0.34–1.11 μ mol/kg tissue [36]. Compared with this report, we consider that the nickel concentration used in our study is slightly higher. In contrast, environmental and occupational exposure to Ni has been reported to result in an increased nickel concentration within the body [37]. Cempel and Janicka reported that oral administration of Ni increases the amount of Ni in the rat brain by approximately 7-fold [38]. Moreover, intranasal instillation of Ni reportedly reached the brain of rats via olfactory neurons [39]. Therefore, we believe that the enhanced activity of Zn²⁺ induced by Ni²⁺ in this study may also occur in the brain.

The ER stress response is important in the pathogenesis of several diseases, such as cancer, acute lung injury, and neurological disorders [40, 41]. ER stress induces the unfolded protein response, which can be distinguished by three ER stress sensors: protein kinase R-like ER kinase (PERK),



FIGURE 4: GT1-7 cells (96-well culture plates at a density of 3×10^4 cells per well) were pretreated with the indicated concentrations (mM) of carnosine just before Ni²⁺/Zn²⁺ treatment. Next, GT1-7 cells were incubated in the absence (control) or presence of NiCl₂ (20 or 40 μ M) and ZnCl₂ (25 μ M) for 24 h (a, b). GT1-7 cells (96-well culture plates at a density of 3×10^4 cells per well) were treated with the indicated concentrations of carnosine for 24 h (c). Cell viability was determined using CellTiter-Glo[®] 2.0. Values represent mean ± S.E.M. *P* values are described in the figure when *P* < 0.05 (black: vs. control, red: vs. Zn(25)/Ni(20 or 40)).

inositol-requiring enzyme-1 (IRE1), and ATF6 [41]. Various signal transduction events are triggered by the activation of these sensors. For example, the α subunit of eukaryotic translation initiation factor 2 (eIF2) is phosphorylated by PERK, which affects the translation of ATF4 (a member of the ATF subfamily of the basic leucine zipper transcription factor superfamily). ATF4 activates the transcription of CHOP and GADD34. In contrast, IRE1 is phosphorylated and interacts with the adaptor protein tumor necrosis factor receptorassociated factor-2, which promotes JNK phosphorylation [40, 41]. As shown in Figure 2, Ni²⁺ treatment activated Zn²⁺-induced increases in the expression of *Chop*, *Gadd34*, and *Atf4*. Hence, it is likely that the PERK/eIF2 α /ATF4 pathway is activated by cotreatment with Ni²⁺ and Zn²⁺.

However, we have been unable to determine the upstream mechanism of ER stress response induction. Thus, the mechanism by which Ni^{2+} treatment enhances Zn^{2+} -induced neuronal cell death needs clarification. As shown in Supplementary Figure S1, we suggest that ROS is one

upstream mechanism of the ER stress response. As one line of evidence, bisphenol A (an endocrine-disrupting chemical) induced cellular apoptosis by activating the ROS-triggered PERK/eIF2 α /CHOP pathway [42]. Moreover, hydrogen peroxide reportedly induced apoptosis by activating ER stress responses in SH-SY5Y cells [43]. Furthermore, luteolin-induced ER stress response (p-PERK/p-eIF2 α /ATF4/CHOP/caspase-12 pathway) was reversed in glioblastoma cells by treatment with the antioxidant N-acetylcysteine [44]. In contrast, nickel chloride exposure induced behavioral disorders, altered neuronal microarchitecture, and induced neuronal cell death by upregulating intracellular ROS [20, 45]. Considering our results and these reports, we predict that Ni²⁺ treatment enhances Zn²⁺-induced ER stress responses by upregulating ROS production.

As shown in Figure 5, we investigated the effect of carnosine on Ni^{2+}/Zn^{2+} -induced neurotoxicity using hypothalamic neuronal GT1-7 cells. To our knowledge, this is the first evidence that carnosine decreases neuronal cell death



FIGURE 5: GT1-7 cells (6-well culture plates at a density of 7.5×10^5 cells per well) were pretreated with the indicated concentrations (mM) of carnosine just before Ni²⁺/Zn²⁺ treatment. Next, GT1-7 cells were incubated in the absence (control) or presence of NiCl₂ (Ni, 40 μ M) and ZnCl₂ (Zn, 25 μ M) for 4 h. Total RNA was extracted and subjected to real-time RT-PCR using primer sets specific for each gene. Values were normalized to *Gapdh* and expressed relative to the control. Values represent mean ± S.E.M. *P* values are described in the figure when *P* < 0.05 (black: vs. control, red: vs. Zn(25)/Ni(40)).



FIGURE 6: GT1-7 cells (6-well culture plates at a density of 7.5×10^5 cells per well) were pretreated with the indicated concentrations of carnosine just before Ni²⁺/Zn²⁺ treatment. Next, GT1-7 cells were incubated in the absence (control) or presence of NiCl₂ (Ni, 40 μ M) and ZnCl₂ (Zn, 25 μ M) for 7 h. Whole-cell extracts were analyzed by immunoblotting with an antibody against CHOP or actin (a). The band intensity of CHOP was determined using ImageJ software (b). Values represent mean ± S.E.M. *P* values are described in the figure when *P* < 0.05 (black: vs. control, red: vs. Zn(25)/Ni(40)).

induced by the cotreatment of Zn^{2+} and other metal ions. Because Zn^{2+} and other metal ions often coexist *in vivo*, we believe that the results of this study are very important. As described above, carnosine is an endogenous dipeptide with various protective activities, such as antioxidant effects and metal ion chelation [24, 25]. Carnosine is abundant in the skeletal muscle, cerebral cortex, kidney, spleen, and plasma [25]. We previously reported that carnosine reduces Zn^{2+} induced neuronal cell death by decreasing the ER stress response [10]. Moreover, we revealed that carnosine exerts its neuroprotective effect without metal ion chelation [10]. Notably, the antioxidant activity of carnosine has been described in many previous studies. For example, carnosine treatment activated antioxidative enzymes (Cu/Zn superoxide dismutase and glutathione peroxidase) in an experimental subarachnoid hemorrhage model [46]. Other groups showed that carnosine directly scavenges hydroxyl radicals [47, 48]. Furthermore, we reported that carnosine prevented lipopolysaccharide-induced ER stress response by suppressing LPS-dependent ROS production [23]. Considering these results, carnosine may counteract Ni and Zninduced ER stress responses and neuronal cell death by exerting antioxidative effects. Therefore, we believe that examining the effect of other antioxidants in these models would be highly worthwhile.

The compounds 6-hydroxydopamine (6-OHDA, a neurotoxin) and rotenone (an electron transport system inhibitor), which are used to model Parkinson's disease, induce neuronal cell death and oxidative stress [49]. Several recent studies have investigated potential interactions between these compounds and trace metals on neurotoxicity. For example, Cruces-Sande et al. demonstrated that copper increases the capacity of 6-OHDA to generate oxidative stress, which may contribute to the destruction of dopaminergic neurons [50]. Another group reported that the cotreatment of iron and rotenone induced a redox imbalance (increased malondialdehyde and decreased glutathione) in the substantia nigra of rats [51]. Zn reportedly induces abnormal phosphorylation of tau by activating various phosphatases, and its phosphorylation is suggested to be involved in the development of Alzheimer's disease [9]. Therefore, we believe that it is important to examine the neurotoxicity induced both by interactions between metals and metals and by interactions between metals and causative substances of diseases, such as 6-OHDA or rotenone.

5. Conclusion

Synergistic neurotoxicity by Ni^{2+} and Zn^{2+} may cause neurological diseases such as vascular dementia, Alzheimer's disease, and Parkinson's disease. In this study, we demonstrated that Ni^{2+} enhances Zn^{2+} -induced neurotoxicity by priming the ER stress response. Furthermore, carnosine was found to protect neuronal cells from Ni^{2+} and Zn^{2+} -dependent synergistic neurotoxicity by decreasing the ER stress response. In conclusion, compounds that decrease the ER stress response, such as carnosine, may be suitable for treating neurological diseases.

Abbreviations

ANOVA:	Analysis of variance
ATF:	Activating transcription factor
CHOP:	CCAAT-enhancer-binding protein
	homologous protein
Cu:	Copper
D-MEM/Ham's-F12:	Dulbecco's Modified Eagle's Med-
	ium/Ham's Nutrient Mixture F-12
EDEM:	ER degradation-enhancing α -manno-
	sidase-like protein

eIF:	Eukaryotic translation initiation factor
ER:	Endoplasmic reticulum
Fe:	Iron
GADD34:	Growth arrest and DNA damage-
	inducible gene 34
Gapdh:	Glyceraldehyde-3-phosphate
-	dehydrogenase
GRP:	Glucose-regulated protein
HRP:	Horseradish peroxidase
IRE1:	Inositol-requiring enzyme-1
LPS:	Lipopolysaccharide
Mn:	Manganese
Ni:	Nickel
6-OHDA:	6-Hydroxydopamine
PERK:	Protein kinase R-like ER kinase
PDI:	Protein disulfide isomerase
ROS:	Reactive oxygen species
Zn:	Zinc.

Data Availability

The data used to support the findings of this study are available from the corresponding author upon request.

Conflicts of Interest

The authors declare that they have no competing interests.

Authors' Contributions

K.T. and M. Kawahara were responsible for the study conception and design. K.T., M. Kasai., M.S., A.S., M. Kubota, and M. Kawahara were responsible for the acquisition, analysis, and/or interpretation of the data. K.T., M. Kasai., M.S., A.S., M. Kubota, and M. Kawahara were responsible for the drafting/revision of the work for intellectual content and context. K.T., M. Kasai., M.S., A.S., M. Kubota, and M. Kawahara gave final approval and have overall responsibility for the published work.

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Supplementary Materials

Supplementary Figure S1: GT1-7 cells were incubated in the absence (control) or presence of NiCl₂ (Ni, 40 μ M) and ZnCl₂ (Zn, 25 μ M) for 4 h. Total RNA was extracted and subjected to real-time RT-PCR using primer sets specific for each gene. Values were normalized to Gapdh and expressed relative to the control (A). GT1-7 cells were pretreated with the indicated concentrations (μ M) of N-acetylcysteine just before Ni²⁺/Zn²⁺ treatment. Next, GT1-7 cells were incu-

bated in the absence (control) or presence of NiCl₂ (40 μ M) and ZnCl₂ (25 μ M) for 24 h (B) or 7 h (D, E). GT1-7 cells were treated with the indicated concentrations of N-acetylcysteine for 24 h (C). Cell viability was determined using CellTiter-Glo[®] 2.0 (B, C). Whole-cell extracts were analyzed by immunoblotting with an antibody against CHOP or actin (D). The band intensity of CHOP was determined using ImageJ software (E). Values represent mean ± S.E.M. *P < 0.05; ** or ##P < 0.01 (* vs. control, #

vs. ZnCl₂ alone, Tukey's test) (A) or (* vs. control, [#] vs. Zn(25)/Ni(20 or 40); Dunnett's test (B) or Tukey's test (E)). (Supplementary Materials)

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