

The Critical Roles of Zinc: Beyond Impact on Myocardial Signaling

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Zinc has been considered as a vital constituent of proteins, including enzymes. Mobile reactive zinc (Zn^{2+}) is the key form of zinc involved in signal transductions, which are mainly driven by its binding to proteins or the release of zinc from proteins, possibly via a redox switch. There has been growing evidence of zinc's critical role in cell signaling, due to its flexible coordination geometry and rapid shifts in protein conformation to perform biological reactions. The importance and complexity of Zn^{2+} activity has been presumed to parallel the degree of calcium's participation in cellular processes. Whole body and cellular Zn^{2+} levels are largely regulated by metallothioneins (MTs), Zn^{2+} importers (ZIPs), and Zn^{2+} transporters (ZnTs). Numerous proteins involved in signaling pathways, mitochondrial metabolism, and ion channels that play a pivotal role in controlling cardiac contractility are common targets of Zn^{2+} . However, these regulatory actions of Zn^{2+} are not limited to the function of the heart, but also extend to numerous other organ systems, such as the central nervous system, immune system, cardiovascular tissue, and secretory glands, such as the pancreas, prostate, and mammary glands. In this review, the regulation of cellular Zn^{2+} levels, Zn^{2+} -mediated signal transduction, impacts of Zn^{2+} on ion channels and mitochondrial metabolism, and finally, the implications of Zn^{2+} in health and disease development were outlined to help widen the current understanding of the versatile and complex roles of Zn^{2+} .

Key Words: Metallothionein, Signal Transduction, Zinc, Zinc exporter, Zinc importer

INTRODUCTION


Zinc is the 24th most abundant element in the Earth's crust, with five naturally occurring isotopes. In 1869, scientists first described zinc as an essential nutrient for the growth of living organisms, due to the lack of its storage in major organs [1]. Zinc deficiency had been identified as a causative factor in the delayed sexual development and stunted growth in humans, because of the vital role of zinc

in both male and female hormone production [2]. Normally, the human body contains 2~3 g of zinc (approx. 15.5 μ M), 0.1% of which is exchanged daily [3]. Cellular zinc is most abundant in the cytoplasm (50%), followed by the nucleus (30~40%), and occurs least in the membrane (10%) [4]. Metalloproteomic analysis has revealed that prokaryotes use approximately 83% of their zinc proteins to carry out enzymatic catalysis, whereas eukaryotes utilize practically equal ratios of zinc-related proteins in catalytic reactions (47%) and DNA transcription (44%), while the remaining are involved in transport (5%) and signaling (3%) [5]. According to Costello et al., zinc occurring in mammalian biological systems can be classified into at least three forms, depending on its molecular behavior [6]: first, the non-exchangeable and non-reactive immobile zinc, which is tightly bound predominantly to proteins and comprises about 54% of the zinc pool; second, the "mobile reactive zinc" that is loosely bound to a ligand and is an exchangeable reactive pool of zinc (44.7%); and third, the reactive pool of zinc or the free zinc (<1%, ~5 pM~1 nM) (Fig. 1). Moreover, five types of Zn^{2+} sites are apparent from the

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ABBREVIATIONS: MT, metallothionein; Zip, zinc importer; Zn, zinc; Zn^{2+} , free zinc or labile zinc; ZnT, zinc exporter.

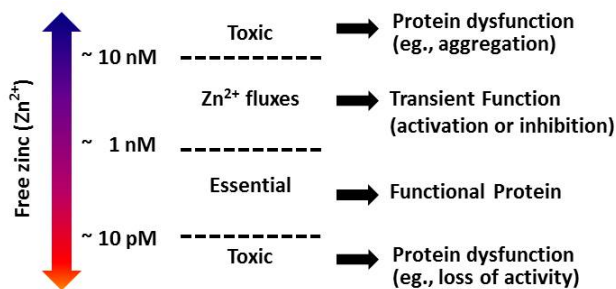


Fig. 1. Impact of Zn^{2+} concentration on protein function in the cell. The total zinc concentration in a eukaryotic cell is relatively high (100–300 μM), but the actual amount of Zn^{2+} ranges from $\sim pM$ to $\sim nM$ values, depending on the cell type. Both extremely low and extremely high levels of Zn^{2+} have adverse effects on the cell. Excess Zn^{2+} causes irreversible effects on proteins, such as aggregation, which leads to the dysfunction of many proteins. Low levels of Zn^{2+} are also detrimental to the cell, since it is an important metal cofactor and signal transducer.

examination of Zn^{2+} -containing proteins: structural, multi-nuclear (clustered), catalytic (over 300 various enzymes), transport, and interprotein [7-9]. This reinforces the possibility of varied activities of zinc in cellular processes. Here, Zn^{2+} represents an exchangeable reactive pool and a reactive pool of zinc (or free zinc). The availability of Zn^{2+} inside the cell influences protein function, more evidently via interprotein Zn^{2+} binding sites, which responds to Zn^{2+} in a concentration-dependent manner [9]. Signaling cascades are induced not by non-exchangeable zinc (tightly bound to proteins), but by mobile reactive or free zinc.

Zn^{2+} can act as an efficient Lewis acid and is often integrated to the amino acid side chains of aspartic acid, glutamic acid, cysteine, and histidine. Among them, cysteine has an important role in the cellular mobility of zinc in proteins. Thus, its flexible coordination geometry with proteins allows a rapid shift of conformations in order to perform certain biological reactions [10-13]. Zinc, while an essential requirement for good health, can be harmful when reaching excessive and toxic levels, or when its homeostasis is perturbed by the suppressed absorption of other metal ions, such as copper and iron [9,14]. The impact of Zn^{2+} concentrations on protein function in the cell is summarized in Fig. 1. The importance and complexity of Zn^{2+} activity has been presumed to parallel the degree of calcium's participation in cellular processes [8,15]. This study aims to summarize the versatile roles of Zn^{2+} , especially on cellular signaling, to better understand its physiological and pathophysiological roles in the heart and other organ systems. In this review, we will briefly summarize: (1) the regulation of cellular Zn^{2+} , (2) Zn^{2+} -mediated cellular functions, (3) Zn^{2+} activity on ion channels, (4) Zn^{2+} and mitochondrial metabolism, and finally, (5) the implications of Zn^{2+} on disease and health.

REGULATION OF CELLULAR ZINC LEVELS IN BLOOD AND IN THE CELL

Zinc homeostasis is maintained through mechanisms that regulate the gene expressions involved in zinc homeostatic pathways; Zn^{2+} transporters (Official name: Solute

Carrier Family 30A, SLC30A, ZnTs), Zn^{2+} importers (Official name: Solute carrier Family 39A, SLC39A, ZIPs), and the intracellular zinc-binding protein, metallothionein (MT) [8,12,16-18]. An outline of cellular Zn^{2+} movement by MTs, ZnTs, and ZIPs is depicted in Fig. 2. Dietary zinc is typically absorbed in the intestine and MT synthesis in mucosal cells is triggered by both fasting and high luminal zinc concentrations [19,20]. Generally, Zn^{2+} is impermeable across the plasma membrane. Hence, after the uptake of Zn^{2+} via ZIPs in the small intestine, the distribution of Zn^{2+} is carried out through the plasma membrane, where it is bound by proteins, mostly albumin ($\sim 77\%$), which is the major protein component of blood plasma and is responsible for the circulatory transport of a range of small molecules. The remaining Zn^{2+} is bound to globulin ($\sim 20\%$) [21,22].

Uptake and extrusion of Zn^{2+}

Intracellular Zn^{2+} must be tightly regulated to prevent the adverse effects of zinc deficiency or toxicity, but since it cannot pass through biological membranes unassisted, two families of zinc-transport proteins aid in this process. A total of 10 ZnT and 14 ZIP gene loci in humans allows for the fine-tuning of Zn^{2+} distribution among tissues and intracellular organelles, such as the endoplasmic reticulum (ER) or Golgi apparatus [15,23-26]. ZnT and ZIP proteins are energy independent, and their production is regulated by Zn^{2+} levels [27]. The fundamental role of Zn^{2+} flux from intracellular stores can be anticipated from the ubiquity of ZIP7 among different organisms and cell types [28], since Zn^{2+} release from stores is stimulated without the addition of exogenous zinc [15]. Most of the ZIP family is predicted to have eight transmembrane domains with extracellular and intravesicular amino- and carboxyl-termini, while most ZnT transporters have six. The metal binding domain of ZnTs is represented by a common histidine-rich loop [29], and this transporter promotes zinc efflux from cells or into intracellular vesicles to reduce intracellular Zn^{2+} availability. However, ZIPs do the opposite by enabling the uptake of extracellular Zn^{2+} , and possibly releasing vesicular Zn^{2+} into the cytoplasm. These transporter families exhibit tissue-specific expression and heterogeneous sensitivity to dietary zinc deficiency and excess, as well as to other physiologic stimuli via hormones and cytokines [30].

Zn^{2+} binding by MT

MT is a small protein (~ 14 kDa) and comprises a family of cysteine (cys)-rich proteins [31]. The human body synthesizes MT in large quantities, mainly in the liver, kidney, intestine, and pancreas. MTs are localized in the membrane of the Golgi apparatus. The thiol group of its cysteine residues ($\sim 30\%$ of its amino acid residues) allows MTs to bind both physiological and xenobiotic heavy metals. The homeostasis of zinc in a system is achieved by the coordination of MT with metal-free thionein, whose production is induced by increased zinc availability. The specific knockout of MT genes in mice with multiple MT-1 genes does not exhibit major developmental defects, suggesting a minimal role in normal growth and development; however, during conditions of major infection and inflammation, MTs have been known to provide survival advantages [32].

MT gene expression is triggered by various stimuli, and the MT promoter region has specific sequences for metal response elements (MRE), glucocorticoid response ele-

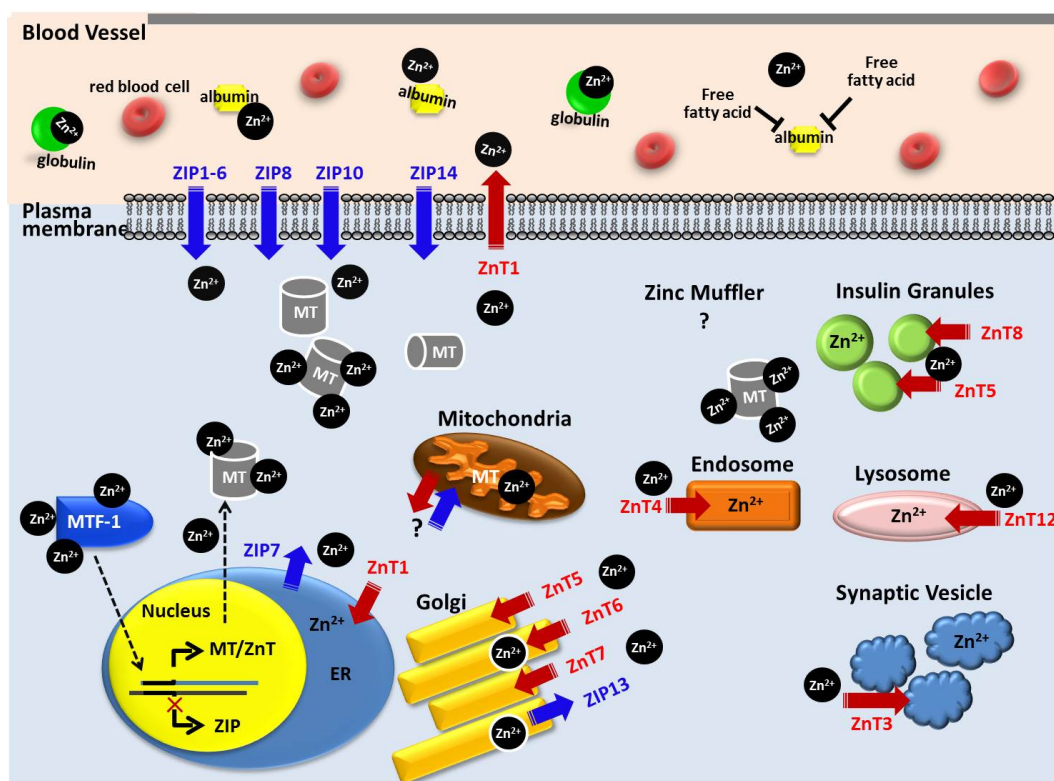


Fig. 2. Subcellular localization of Zn^{2+} transporters. Zinc absorbed in the intestine as an ion goes into the blood stream. Serum albumin is a major Zn^{2+} binding protein and serves as a macro Zn^{2+} transporter to target tissues; thus, changes in the concentration of albumin or an increase of fatty acids, which interferes with Zn^{2+} binding to albumin, impairs the control of blood Zn^{2+} levels. The availability of intracellular Zn^{2+} is governed by the action of MTs (MTs; binding or trapping of Zn^{2+}), zinc transporters (ZnTs; Zn^{2+} efflux), Zn^{2+} importers (ZIPs; Zn^{2+} influx) residing in the plasma membrane, redox potential, pH, and other unknown modulators. ZIPs and ZnTs residing in the organelle membrane have an inverse direction compared to those located in the plasma membrane. The activation of metal-responsive-element-binding transcription factor-1 (MTF-1) increases the expression of MTs, and thereby reduces Zn^{2+} levels upon binding to Zn^{2+} . Specific forms of MTs, ZnTs, and ZIPs are dependent on cell-type, which will give varying responses, even at the same zinc levels. Extracellular Zn^{2+} entering the cell could be immediately buffered within the still undefined 'zinc muffer' and translocated into an intracellular Zn^{2+} store, such as the endoplasmic reticulum (ER) and Golgi apparatus by ZnT. Inversely, Zn^{2+} released into the cytosol from Zn^{2+} stores by ZIPs is also buffered by a cytosolic Zn^{2+} muffer. Arrows show the predicted direction of Zn^{2+} mobilization.

ments, GC-rich boxes, basal level elements, and thyroid response elements [32]. Transcriptional activation also occurs at higher zinc levels, as observed in the gene regulatory response mediated by MRE-binding transcription factor 1 (MTF1) to copies of the MRE consensus sequence in the promoter region [33]. MT also serves as a free radical scavenger [31]. Mammalian MT-1/MT-2 isoforms are involved in zinc homeostasis and protection against heavy metal toxicity and oxidative stress. MT-3 is expressed mainly in neurons, but also in glia [34], while MT-4 is mostly present in differentiating stratified squamous epithelial cells [31].

Sequestration and release of Zn^{2+}

As extracellular Zn^{2+} enters the cell, it is immediately buffered within a 'zinc muffer' via a still unidentified Zn^{2+} transporter, and is then translocated into an intracellular zinc store, such as the ER or Golgi [35]. The Zn^{2+} is then released into the cytosol from the ER through the action of ZIP7, creating a zinc wave [36]. Mitochondria are another possible site of intracellular zinc sequestration. Zn^{2+} can be released from zinc-bound MT, physiological Zn^{2+} ligands

[37], cysteine, histidine, glutathione (GSH), aspartate, glutamate, and glycine, in an acidic condition such as in ischemia [38,39], from ER and Golgi through ZnTs, mitochondria with unknown mechanisms [40], or acidic organelles [41]. The partial cationic character of Zn^{2+} enables it to attract anionic prooxidants, such as peroxynitrite ($ONOO^-$). For example, zinc is released from endothelial nitric oxide synthase (eNOS) upon exposure to peroxynitrite, which results in dimer disruption and enzyme uncoupling [42]. Maret and colleagues demonstrated that the presence of GSH enabled a higher transfer rate, and caused the release of more of the available Zn^{2+} from MT by glutathione disulfide (GSSG), and, thereafter, can facilitate Zn^{2+} removal and addition to inhibitory sites on enzymes involved in glycolysis and signal transduction [43]. A pH-dependent release of Zn^{2+} by cytosolic proteins is evident with cysteine, wherein the liberation of Zn^{2+} rapidly accelerates when the pH is decreased from 6.6 to 6.1 [37]; however, MTs are not likely to release Zn^{2+} at the same pH trend [39] and Zn^{2+} release from GSH-Zinc is less sensitive to an acidic pH (pH 6.1) [37]. At physiological pH, cytosolic proteins and ligands prevent large increases in intracellular Zn^{2+} by rapidly buffer-

ing the Zn^{2+} released from the mitochondria. When the mitochondrial matrix pH drops from 8 to 7, GSH starts to release Zn^{2+} , which then leaks into the cytosol. At a pH of 7, ligands, such as MTs and cysteine, intercept the released Zn^{2+} ; only after the pH drops below 6.6 and cysteine loses its affinity for Zn^{2+} does the elevation of intracellular Zn^{2+} levels become evident [37].

ZINC AND ITS REGULATION OF CELLULAR FUNCTIONS

The role of Zn^{2+} in signal transduction pathways has recently become a subject of great interest, since numerous signaling pathways are influenced in the presence or loss of Zn^{2+} [12]. Binding of Zn^{2+} has been observed to prompt protein structural rearrangement, which eventually affects protein function [9]. Fukada et al. suggested that signaling evoked by intracellular Zn^{2+} may be classified by time windows, as transcription-independent early Zn^{2+} signaling happened within several minutes and transcription-dependent late Zn^{2+} signaling happened several hours after Zn^{2+} exposure, based on the actions of immune cells [12]. However, it remains unclear whether this Zn^{2+} -mediated signaling is sequential from early to late Zn^{2+} signaling, or whether both early and late Zn^{2+} signaling are separate or combinatory.

Zn²⁺ in signal transduction

In cardiomyocytes or other cells, the intracellular zinc concentrations are 100-fold less than those of intracellular Ca^{2+} [4,44]. The relatively low abundance of Zn^{2+} in cardiomyocytes suggests its availability as second messengers for gene expression [45]. The inhibition of cyclic AMP (cAMP) signaling by Zn^{2+} has been attributed to both an inhibition of adenylyl cyclase (AC), particularly the AC5 and AC6 isoforms [46,47] that are present in the cardiomyocytes [48], and of the function of the Gs alpha subunit (*G α s*). Zn^{2+} induces aggregates of pure tubulin, implying the activation of G proteins by tubulin dimers [49]. This possible regulation of cAMP signaling by Zn^{2+} will provide complex outcomes of cAMP and cAMP-protein kinase A signaling, which are initiated by various kinds of chemicals and hormones. Therefore, cAMP-mediated signaling may be influenced or amplified at different levels of cellular Zn^{2+} .

Protein tyrosine phosphatase 1B (PTP1B) inhibits phosphoinositide 3-kinase, a key enzyme in the insulin and leptin signaling pathways, making it a negative regulator of both signaling pathways [50]. Zn^{2+} binds to the catalytic pocket of the phosphointermediate form of PTP1B at a nanomolar range [51], and works as an inhibitor of protein tyrosine phosphatases (PTPs) on account of an active site HC(X5)R signature motif and the nucleophilic attack of the phosphorus atom of the phosphate group by the sulfur ion of the cysteine thiol in the first step of the dephosphorylation reaction and/or formation of inactive dimers between thiolates of PTPs and zinc [16,51]. ER-bound PTP1B is proposed to be inhibited by ZIP7-mediated Zn^{2+} release from the ER, because the co-localization of ZIP7 and PTP1B does not require a global increase in cytosolic Zn^{2+} in order to inhibit the enzyme. The inhibitory effect of Zn^{2+} on PTPs implicates zinc-induced signaling in the increase in tyrosine phosphorylation of insulin receptor (IR), insulin-like growth factor 1 receptor (IGF-1R), epidermal growth factor re-

ceptor (EGFR), or unrecognized receptor tyrosine kinases, depending on the cell type [52]. Moreover, both insulin and adiponectin directly work together with Zn^{2+} , affecting their oligomerization, and hence their activities [21]. Therefore, an elevation in disturbances in intracellular Zn^{2+} status observed in metabolic disorders, such as obesity and diabetes, may directly impact leptin signaling by PTP1B inhibition [53]. In insulin-resistant rodent models, key regulators of the insulin signaling pathway, extracellular-signal-regulated kinases 1 and 2 (ERK1/2) and protein kinase B (PKB), were activated by Zn^{2+} , and insulin-mimetic and anti-diabetic properties were also observed [52].

Zn²⁺ in redox homeostasis

The cell's response to a perturbation in its surrounding environment heavily relies on the regulation of protein activity. Reactive disulfide bonds act as molecular switches, and proteins with closely spaced cysteines or cysteine-coordinating Zn^{2+} centers, including MT, may reveal that thiol-disulfide exchanges play a much more widespread role in protein regulation at the post-translational level than had been previously anticipated [54]. Intracellular Zn^{2+} levels can be increased by thiol-reactive oxidants. In cardiomyocytes, reactive oxygen species (ROS) are major contributors to a rapid increase in intracellular Zn^{2+} levels, due to the mobilization of Zn^{2+} from intracellular stores. Moreover, during oxidative stress, including ultraviolet A irradiation, Zn^{2+} is released from MTs, either by nitrosylation or thiol ligand oxidation [55,56]. Intracellular Zn^{2+} may also contribute to oxidant-induced alterations of excitation-contraction coupling [55], changes in signal transduction, and alterations of mitochondrial function due to modification of protein structure or functions by Zn^{2+} release, or its binding to proteins [29]. These findings explain why antioxidant intervention, resulting in the prevention of both Zn^{2+} and Ca^{2+} leaks, is helpful against diabetic heart disease [45,57]. Zn^{2+} plays a central role in the redox signaling network, in that when it binds to a sulfur ligand, it generates a redox-active environment, which allows conformational changes of the proteins it is attached to, activating several signaling molecules. For example, the activation of protein kinase C (PKC) could be initiated by the release of Zn^{2+} from its zinc finger regulatory domain by lipid second messengers or oxidation [58]. Another example is the Kv4.1 channel, a voltage-gated potassium channel that is a key regulator of membrane excitability in neuronal and cardiac tissues [59,60]. Under physiological conditions, the native Kv4 channel complex activity is regulated by thiol-disulfide exchange reactions at the Zn^{2+} site upon exposure to reactive nitrosative or oxygen species. Intracellular nitric oxide (NO) rapidly inhibits the Kv4.1 channel in a Zn^{2+} -dependent manner through a disulfide bridge across the T1-T1 interface (Cys¹¹⁰ to Cys¹³²), which may be responsible for the nitrosative/oxidative modulation of rapidly inactivating (A-type) K^+ currents in the heart and smooth muscle, as well as in neurons [59]. Other Kv2 and Kv3 channels share the similar T1-T1 Zn^{2+} site, and thus they also can be regulated by zinc-dependent redox modulation.

Zn²⁺ in the modulation of ion channels

Numerous proteins are targets of Zn^{2+} binding, and this direct binding of Zn^{2+} to proteins is able to directly alter the function of proteins, including ion channels. Zn^{2+} is able

to induce changes of the ionic current and significantly impede the activation kinetics of most K^+ channels through its direct binding [60]. In addition, Zn^{2+} is among the most potent inhibitors of voltage-gated proton (H^+) and calcium channels [61,62]. The abundance of potential synaptic targets and the localization of Zn^{2+} in synaptic vesicles all seem to point to a role of Zn^{2+} in synaptic transmission [63]. The entry of Zn^{2+} in cortical neurons is mediated by L- and N-type voltage-gated Ca^{2+} channels, which may be enhanced by an acidic extracellular pH [64], whereas Zn^{2+} entry into heart cells is via dihydropyridine-sensitive Ca^{2+} channels and is reliant on electrical stimulation [65]. In the dorsal root ganglion (DRG) neurons, the rise in nanomolar concentrations of intracellular Zn^{2+} can activate transient receptor potential ankyrin 1 (TRPA1), which is a sensory receptor for environmental irritants and both oxidative and thiol-reactive compounds, possibly through interaction with the intracellular cysteine and histidine residues of TRPA1, and thus potentiates the perception of discomfort [66,67].

Zn^{2+} in mitochondrial metabolism

Mitochondrial enzymes are also influenced by Zn^{2+} or oxidative stress-related Zn^{2+} release, and its dysregulation can take a toll on mitochondrial metabolism [40,68-70]. Although there is limited information available regarding Zn^{2+} -dependent mitochondrial functions, changes in mitochondrial metabolism could be introduced by the direct binding of Zn^{2+} to mitochondrial proteins or the release of Zn^{2+} from its bound proteins, due to oxidative stress or redox switches, which result in an alteration of its enzymatic activity and function [71]. In the mitochondrial matrix, GSH has a more prominent action in Zn^{2+} chelation than it does in the cytosol. The application of a sulfhydryl reactive agent, N-ethylmaleimide, leads to a major intracellular Zn^{2+} release, indicating a large amount of zinc storage in Zn^{2+} -cysteine complexes [39,72]. While GTP binds to MT [73], ATP also binds to mammalian MT-2 [74]. Thus, it has been proposed that released Zn^{2+} from MT may be involved in the inhibition of ADP-stimulated respiration, or the formation of an ATP- Zn^{2+} complex, which is a preferred cofactor of pyridoxal kinase and flavokinase, which will be able to influence energy metabolism [74].

Zinc can inhibit mitochondrial electron transport by binding Zn^{2+} in complexes [75,76] and stimulating a burst of mitochondrial ROS production [77]. Several zinc-regulated enzymes have been identified. Fructose 1, 6-diphosphatase, glyceraldehyde 3-phosphate dehydrogenase, aldehyde dehydrogenase, and lipoamide dehydrogenase (LADH) are inhibited by nanomolar concentrations of Zn^{2+} [71,78]. Inside the mitochondrial matrix, an elevated Zn^{2+} concentration interrupts the Krebs cycle and mitochondrial energy metabolism by inhibiting the production of a reduced form of nicotinamide adenine dinucleotide (NADH) production. Besides, the Zn^{2+} pool in prostate mitochondria ($0.1 \sim 1 \mu g$ zinc/mg protein) is essential for inhibiting mitochondrial aconitase in the Krebs cycle, possibly through MT-mediated Zn^{2+} transfer [79], which results in the accumulation of citrate for secretion into prostatic fluid [80,81]. In neurons, excessive Zn^{2+} could induce mitochondrial swelling and ROS release. However, whether or not this induced swelling also occurs in the liver remains controversial [82,83].

In mitochondria-mediated apoptosis, Zn^{2+} increases the Bax-associated mitochondrial pore forming process that releases cytochrome C, activating the caspase cascade that

leads to apoptosis [84]. Zn^{2+} also increases the gene expression and cellular production of Bax and the Bax/Bcl-2 ratio, which facilitates the mitochondrial Bax-associated apoptogenic effect [85]. Zn^{2+} chelation at the onset of neuronal ischemia attenuates mitochondrial cytochrome C release and downstream caspase-3 activation [86]. In contrast to brain ischemia [38], Zn^{2+} is considered as an important modulator involved in cardioprotection upon reperfusion, since several suggested cardioprotectants share zinc activation [10].

ZINC AND ITS ROLE IN HEALTH AND DISEASE

In conjunction with the versatile roles of Zn^{2+} , the differential expression of MT, ZnT, or ZIP participates in various kinds of pathological states, including inflammation [12,32], neurodegenerative diseases, cancer, myocardial hypertrophy, and environmental metal exposure, since intracellular Zn^{2+} levels are largely influenced by these Zn^{2+} regulatory proteins [31]. The brief representation of Zn^{2+} regulation and possible outcomes is summarized in Fig. 3. Generally, dysregulated zinc metabolism in tissues with an intricate Zn^{2+} requirement is associated with metabolic diseases, such as cancer, diabetes, and infertility [9,12,14,87-90]. Aside from inadequate dietary zinc intake, zinc deficiency may result from impaired absorption, resorption, or increased excretion of Zn^{2+} . Conditions such as chronic diarrhea, widespread burns, or wounds also accelerate endogenous zinc losses [91]. Alcoholics also frequently develop zinc deficiency, since alcohol causes the body to lose critical Zn^{2+} stores through the kidneys.

Zn^{2+} in the nervous system

In the mammalian brain, vesicular zinc is co-localized with glutamate in a subset of glutamatergic nerve terminals throughout the forebrain. Zn^{2+} also shows anti-depression effects [92], which are associated with the action of G protein-coupled receptor 39 (GPR39), which is recognized as a Zn^{2+} receptor, and is a potent stimulator of GPR39 activity via the $G\alpha(q)$, $G\alpha(12/13)$, and $G\alpha(s)$ pathways [93,94]. Impairments of the central nervous system (CNS), such as defects in learning and memory, may be brought about in the fetus by maternal zinc deficiency, which cannot be reversed by subsequent supplementation of zinc to the diet after weaning [95]. This CNS defect may be associated with a reduction in the number of proliferating stem cells and neuronal precursor cells due to inadequate amount of available zinc [96,97]. In various neuronal diseases, such as stroke, epilepsy, mechanical trauma, and Alzheimer's disease, the precise mechanism of cytotoxicity is less known, but emerging evidence suggests that excess Zn^{2+} may kill neurons through the inhibition of ATP synthesis [98]. In contrast to the previously mentioned proapoptotic role of Zn^{2+} [84], Zn^{2+} could participate in the inhibition of proapoptotic caspase-3 activation [71] and in the reduction of amyloid production in experimental neuronal cells [99], since these effects are lost in response to low levels of Zn^{2+} . In zinc deficiency, neural cell apoptosis was associated with an increase in phosphorylation of p53 and its translocation into mitochondria [97].

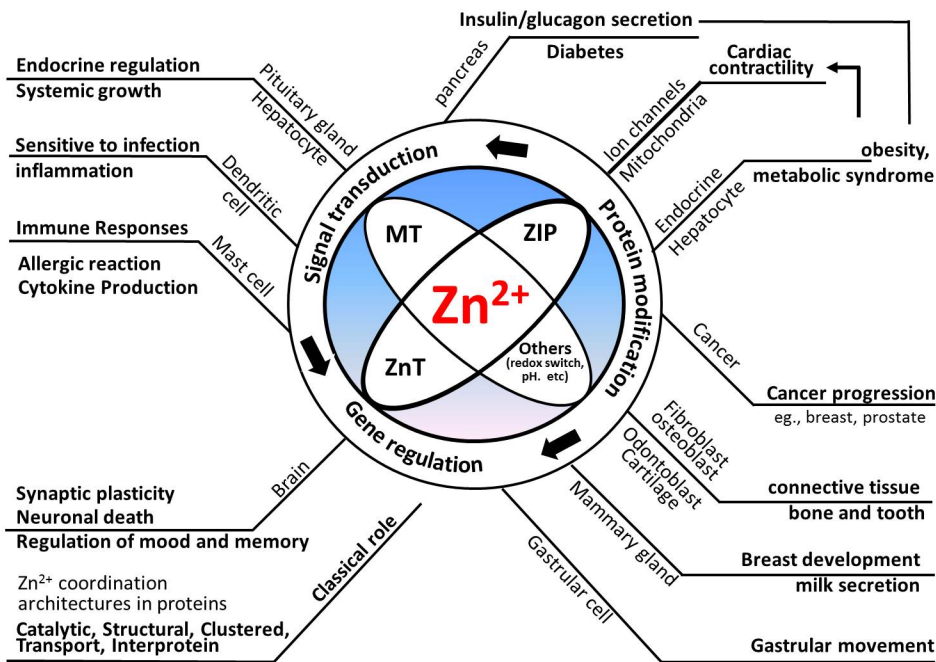


Fig. 3. Roles of Zn^{2+} in a biological system. Various tissues and organs are influenced by Zn^{2+} . The intracellular Zn^{2+} levels are primarily regulated by MTs, ZnTs, and ZIPs. The action of Zn^{2+} can be initiated within a few minutes by protein modification involved in enzyme activities and signal transduction, or in late events, which are mediated by gene regulation. Therefore, the impairment of Zn^{2+} homeostasis, such as deficiency or excess, can lead to cellular malfunction, and eventually, cell damage.

Zn^{2+} in the cardiovascular system

The extracellular and intracellular levels of Zn^{2+} may be a determinant to cardiovascular health [29], and zinc deficiency related to diseases and aging may be a factor in the development of heart failure. Moreover, a potential role of zinc in diabetic cardiomyopathy has been suggested, due to its involvement in insulin signaling and diabetes development [100,101]. In regards to the cardioprotective roles of Zn^{2+} , four beneficial roles of Zn^{2+} were suggested: (1) protective effects from acute redox stress in cardiomyocytes; (2) prevention of inflammatory processes triggered during myocardial damage; (3) wound healing; and finally (4) maintenance of cardiac stem cells essential for cardiac healing [29].

In the heart, a large portion of intracellular Zn^{2+} accumulates in specific cellular structures, such as in or near the terminal sarcoplasmic reticulum (SR) [65]. Calsequestrin is a Zn^{2+} -binding protein residing in the SR and can bind up to 200 mol of Zn^{2+} per mol [55]. Besides changes in the phosphorylation status of the L-type calcium channel (LTCC), thiol oxidation, and nitrosylation, the extracellular and intracellular levels of Zn^{2+} influence transmembrane Ca^{2+} movements via the LTCC. Moreover, intracellular Zn^{2+} levels could inhibit AC at even low concentrations (<1 nM), which strikingly reduces β -adrenergic stimulation [47]. Moreover, besides being cell permeant, external Zn^{2+} can prevent Ca^{2+} entry by inhibiting LTCC at concentrations as low as tens of nanomolars, which in turn suppresses the Ca^{2+} -induced Ca^{2+} release from the SR [47,102]. Zn^{2+} depresses the contractile force logarithmically with time, which seems to be related with dysregulation in calcium handling in electrically stimulated rat atrial cells [103] or smooth muscle cells [104]. In pre-diabetic rats, the improvement of diastolic relaxation by Zn^{2+} is attributed to a depressed LTCC inward current, phosphorylation of the rya-

nodine receptor and phospholamban, resulting in a reduced cellular Ca^{2+} load [89]. In guinea pigs, the administration of low concentrations of Zn^{2+} caused mainly sinus bradycardia, while higher concentrations of Zn^{2+} (30 μ M) resulted in various types of arrhythmia that lead to cardiac asystole [105]. In humans, the cardiac ejection fraction increases with Zn^{2+} content [106], and Zn^{2+} supplementation to a cardioplegic solution reverses the loss of systolic function [27]. However, the importance of Zn^{2+} in cardiac muscle function is still incompletely understood.

In patients with ischemic cardiomyopathy, a higher copper level and a lower zinc level were observed [107]. Diabetic patients with congestive heart failure also exhibited a high concentration of urinary Zn^{2+} excretion [107]. Normally, the majority of plasma Zn^{2+} is associated with human serum albumin (HSA), but this interaction is allosterically disrupted by free fatty acids (FFAs), which eventually results in hypozincemia due to altered Zn^{2+} uptake in the cell [21]. Zinc deficiency is associated with enhanced PTP activity, which increases insulin receptor signaling and insulin resistance, which is a risk factor for cardiovascular disease development [108]. Theoretically, all conditions that lead to lower energy consumption by the heart could lead to at least temporary elevations of FFAs. Therefore, the possibility of a direct correlation between lower plasma Zn^{2+} levels and higher levels of FFAs cannot be overlooked, as it is evident in acute myocardial infarction [109] and chronic heart failure [110]. Zinc deficiency may also have deleterious effects in patients with heart failure, due to decreased intake secondary to decreased absorption, increased urinary loss through diuretics, or high levels of lipid peroxides as a result of oxidative stress [108]. However, paradoxically, heart failure medications, such as thiazide diuretics, angiotensin-converting enzyme inhibitors, angiotensin receptor blockers, and the diuretic furosemide can upset zinc metabolism, and subsequently lead to zinc deficiency [108,111].

PKC is a central molecule in many signaling pathways and is particularly prominent in vascular smooth muscle and endothelial cells. As previously mentioned [58], alteration of PKC activity directed by the movement of Zn^{2+} modulates the action of multiple vasoactive factors that promote atherosclerotic events. Zn^{2+} may play a protective role in the maintenance of the integrity of endothelial cells [112] or the prevention of ultraviolet-induced skin damage [113]. Chronic zinc deprivation accelerates vascular smooth muscle cell proliferation [114], and in combination with calcification, may aggravate atherosclerosis progression [29].

Zn²⁺ in secretory tissues

Pancreatic β cells contain the highest amount of zinc among all cells within the human body. Zn^{2+} is vital for the precise handling of insulin in β -cells, from storage to the exertion of its effects. It has been suggested that the opening of ATP-sensitive K^+ channels (K_{ATP}) in the pancreas is regulated by the activation of Zn^{2+} on both sides of the membrane, with potentially discrete sites of action located on the sulfonylurea receptor (SUR) subunit [115]. It has been recognized that zinc deficiency, or the aberrant expression of ZnT (e.g., ZnT8) or ZIP (e.g., ZIP4) proteins in the pancreas, is linked to the occurrence of pancreatic cancer and chronic pancreatitis [80]. Secreted Zn^{2+} exhibits autocrine or paracrine signaling in β -cells or neighboring cells. During glucose deprivation, Zn^{2+} serves as an intercellular signaling ion, and provides an "off-switch" for glucagon release from the pancreatic α -cells. Zn^{2+} also modulates the exocrine production of digestive enzymes and their role in nutrient absorption. Zinc's entry into the digestive tract is facilitated by enzyme-containing zymogen granules secreted from the acinar cells of the pancreas. The activity of zymogen is Zn^{2+} -dependent, and ZnT2 aids in zinc transport, which eventually leads to digestive enzyme secretion [80].

Unlike other cells, prostate cells contain almost 35% of their total zinc in the cytoplasm, rather than sequester Zn^{2+} into vesicles and organelles. High zinc levels are emitted into the seminal liquid, assuming a critical role in sperm discharge and motility [116]. Antimicrobial properties have also been associated with Zn^{2+} in prostatic fluid and the gland itself. In cancer biology, evidence points to zinc's involvement in tumor suppression in several cancers [117]. Reduced levels of zinc in the prostate were also linked to prostate diseases, including benign prostatic hyperplasia and prostate carcinoma [80].

While zinc's role in mammary gland development [80] and remodeling is indisputable, the effects of its deficiency remain poorly understood. A single amino acid substitution in human ZnT2 reduces milk zinc concentration by about 75% and results in severe dermatitis and alopecia in the nursing infant [118]. Tissue expansion in the mammary gland is also mediated by Zn^{2+} -dependent matrix metalloproteinases (MMPs). Again, the availability of cellular Zn^{2+} dictates the enzyme activity; therefore, a decrease in activity could result in impaired remodeling and subsequent breast disease.

Zn²⁺ in the immune system

Zinc deficiency is manifested by an immunocompromised status due to thymic atrophy, lymphopenia, or compromised cell- and antibody-mediated responses, which causes

increased infection rates with prolonged durations [32,119]. Zinc deficiency stimulates the production of cytokines, such as interleukin (IL)-1 β , IL-2, IL-6, and tumor necrosis factor (TNF)- α , and the generation of these cytokines shows inverse responses to zinc supplementation [91]. Therefore, mild zinc deficiency with low-grade inflammation has been found in patients with atherosclerosis and diabetes mellitus [91]. In the allergic response, Zn^{2+} and its transporters are indispensable for Fc epsilon receptor I (Fc ϵ RI)-mediated mast cell activation, since blocking or supplementation of zinc largely affects this allergic reaction through the involvement of degranulation and cytokine production [12]. In monocytes, Zn^{2+} affects differentiation by increasing cAMP and its guanosine analog (cGMP) through the inhibition of ACs and cyclic nucleotide phosphodiesterases [46,120].

SUGGESTIONS OF NEW APPROACHES AND PERSPECTIVES ON ZN^{2+} MODULATION

Previous studies have focused on the general maintenance of Zn^{2+} homeostasis, but recent studies suggest the versatility of Zn^{2+} as a signaling molecule. As briefly outlined in Fig. 3, numerous functions of tissues, including the brain, immune cells, cardiovascular tissues, pancreas, and prostate and mammary glands, are influenced by the actions of Zn^{2+} . In addition, cancer progression could be associated with the dysregulation of Zn^{2+} homeostasis and its disturbed cellular signaling. However, the action of Zn^{2+} is not simple and is heterogeneous in different tissues, as confirmed by the tissue-specific expression of MT, ZnT, and ZIP proteins, and their distribution and possible roles in physiological conditions and disease progression. Cellular redox statuses are also important in understating Zn^{2+} behavior in a given system, because protein-bound Zn^{2+} , as well as free Zn^{2+} , can be released or bound to certain proteins, thus altering their structure and functions, which ultimately alters cellular responses. Considering the numerous roles of Zn^{2+} mentioned in this review, Zn^{2+} and its subsequent signaling events are not simple, and without the regulation of Zn^{2+} , catastrophic responses may occur within cells. This raises the question of how cells can tightly regulate the action of Zn^{2+} , and when it is initiated and terminated. Cellular compartments of Zn^{2+} , regional crosstalk from Zn^{2+} release or translocation from inside or outside of the cell, and the removal of Zn^{2+} by MT or other Zn^{2+} -binding proteins may provide a partial explanation as to how Zn^{2+} is able to partake in several cellular processes. These issues seem to be important, and, thus, more intensive investigation should be undertaken.

As previously mentioned, intracellular Zn^{2+} levels sensed by MTF-1 can lead to the induction of Zn^{2+} -regulating proteins. However, when cells sustain zinc deficiency, detection mechanisms and the ensuing cellular compensatory processes have not yet been proposed. Therefore, Zn^{2+} sensing mechanisms in the condition of both excess and deficient levels will be necessary to extend our understanding of the role of Zn^{2+} in physiology. Except for the known role of MTF-1, there are still gaps in our knowledge of how Zn^{2+} exerts its action. It should be noted that Zn^{2+} 's role as a signaling molecule and transcriptional regulator may not be solely attributable to Zn^{2+} , since other essential metals are equally important to maintain cellular homeostasis. However, this is beyond the scope of our current review,

and the contributions of other ions are not discussed in this manuscript. Inhibition of PTP1B by Zn²⁺ exposure has been widely recognized. In signal transduction pathways, the phosphorylation status of proteins is regulated by well-known mediators, namely kinases and phosphatases, that influence the functions of proteins. However, direct changes of their activities invoked by the introduction of Zn²⁺ or the chelation of Zn²⁺ in proteins are also largely unknown. Therefore, it is also necessary to expand our knowledge of Zn²⁺'s role as a molecular switch or connector.

Cellular Zn²⁺ levels can be measured by the aid of Zn²⁺-specific fluorescent dyes, including cell-impermeable Zinquin-A (free acid), cell permeable Zinquin-E (ethyl ester) [121], TSQ [122], FuraZin-1 AM, RhodZin, and FluoZin. Technically, the movement of Zn²⁺ is largely determined by Zn²⁺-specific fluorescent probes. Although these probes are useful in zinc biology, the specificity and sensitivity of Zn²⁺-detection probes are considered a technical hurdle. Therefore, a rapid and direct method of measurement of Zn²⁺ should be further developed. A proteomic analysis of Zn²⁺-containing proteins provides valuable insight into understanding the biological role of Zn²⁺ [9]. The development of strategies to revert the dysregulation status of cellular Zn²⁺ will be a promising approach of future direction. In addition, an integrated and systemic understanding of the newly identified roles of Zn²⁺ will be necessary.

In conclusion, a wider perspective of the roles of Zn²⁺, beyond being a mere building block of proteins, will be valuable for a comprehensive understanding of the physiological and pathophysiological functions of Zn²⁺ in the cardiovascular system and other tissues. Further research on the critical roles of Zn²⁺ and its regulation will aid in the maintenance and improvement of human health.

CONFLICTS OF INTEREST

The authors declare no conflicts of interest.

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