

REVIEW

Zinc and its role in age-related inflammation and immune dysfunction

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Zinc is an essential micronutrient required for many cellular processes, especially for the normal development and function of the immune system. Zinc homeostasis and signaling are critical in immune activation, and an imbalance in zinc homeostasis is associated with the development of chronic diseases. Zinc deficiency causes significant impairment in both adaptive and innate immune responses, and promotes systemic inflammation. The elderly are a population particularly susceptible to zinc deficiency. National surveys indicate that a significant portion of the aged population has inadequate zinc intake, and a decline in zinc status is observed with age. There are remarkable similarities between the hallmarks of zinc deficiency and immunological dysfunction in aged individuals. Both zinc deficiency and the aging process are characterized by impaired immune responses and systemic low grade chronic inflammation. It has been hypothesized that age-related zinc deficiency may be an important factor contributing to immune dysfunction and chronic inflammation during the aging process. In this review, we discuss the effects of zinc status on aging, potential molecular and epigenetic mechanisms contributing to age-related decline in zinc status, and the role of zinc in age-related immune dysfunction and chronic inflammation.

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1 Introduction

Aging is a complex process associated with physiological changes in numerous organ systems. In particular, aging of the immune system, or immunosenescence, results in a progressive dysregulation of immune responses, affecting multiple immune cell types involved in both adaptive and innate immunity. Immunosenescence is characterized by involution of the thymus, defective lymphocyte maturation and function, and a gradual decline in both cellular and

humoral immune responses, resulting in increased susceptibility to infectious diseases and compromised vaccination efficacy in the elderly [1]. At the same time, aging is associated with “inflamm-aging”, a low-grade systemic chronic inflammation characterized by constitutively elevated levels of proinflammatory cytokines [2, 3]. Inflammation is a physiological process mediated by the innate immune system to protect against pathogen infections, environmental insults, and to promote wound healing. However, aberrant or excess inflammation is detrimental, as overproduction and prolonged exposure to inflammatory mediators (including inflammatory cytokines, prostaglandins, and reactive oxygen and nitrogen species) can result in changes in cell proliferation, apoptosis, angiogenesis, and genetic instability including alterations to DNA repair, increased DNA damage, and aberrant DNA methylation [3, 4]. Accumulating evidence indicates that chronic inflammation is commonly observed in the elderly, and increases in inflammatory mediators in the blood are significant predictors of morbidity and mortality in aged individuals [5]. Chronic inflammation has been correlated to

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Abbreviations: **AE**, acrodermatitis enteropathica; **DC**, dendritic cells; **DNMT**, DNA methyltransferase; **EAR**, estimated average requirement; **LPS**, lipopolysaccharide; **MT**, metallothionein; **MTF-1**, metal-regulatory transcription factor 1; **NF- κ B**, nuclear factor-kappaB; **SLC**, solute-linked carrier; **TCR**, T cell receptor

the promotion of many age-related diseases. For example, elevated levels of proinflammatory cytokines such as IL-6 and TNF α have been associated with age-related diseases such as cardiovascular disease, type-2 diabetes, osteoporosis and autoimmune diseases such as rheumatoid arthritis and systemic lupus erythematosus [6]. Further, epidemiological studies indicate that up to 25% of all cancers are associated with inflammation [7]. Thus there is a critical need to develop strategies to reduce age-associated chronic inflammation to improve the quality of life for the elderly. While the etiology of immunosenescence and inflamm-aging is multi-faceted, we postulate that nutritional deficits such as zinc deficiency that are prevalent in the elderly may play an important role in contributing to the perturbations in the immune system with age. Further, improving zinc status in the elderly via dietary zinc supplementation may reverse and correct age-related immune dysfunction and chronic inflammation resulting in improved health outcomes.

Zinc is an essential micronutrient required for many biological processes including growth and development, neurological function, reproduction, and immunity. In particular, the importance of zinc for the immune system is underscored by the effects of zinc deficiency, which results in immune dysfunction including thymic atrophy, lymphopenia, impaired adaptive immunity, and chronic inflammation [8–10]. Low zinc status is associated with increased susceptibility to infections and exaggerated inflammatory responses, and zinc imbalance may contribute to the onset and/or progression of diseases including asthma, diabetes, and Alzheimer's diseases [11]. Individuals particularly at risk for zinc deficiency include the elderly, vegetarians, and patients with chronic illnesses. In particular, low zinc status in the elderly has been shown to contribute to age-associated dysregulation of immune function, and may be a contributing factor in age-related inflammation and associated morbidities [12]. However, the precise mechanisms linking zinc, age, and inflammation are unclear. In this review, we will highlight the effects of zinc status on aging, potential mechanisms that contribute to age-related decline in zinc status, and the role of zinc in immune responses with emphasis on age-related immune dysfunction and chronic inflammation. We will also discuss the potential role of epigenetic control on zinc homeostasis, and how epigenetic alteration with age may contribute to age-related zinc deficiency and inflammation.

2 Zinc status and aging

Zinc is a key component for the function of numerous proteins, including zinc-containing metalloenzymes and zinc-associated transcription factors. It is the most abundant trace intracellular element, and is an essential micronutrient required for numerous biological functions, including cell proliferation, differentiation, and reproduction. In addition, zinc is essential for the normal development and function of

the immune system [9]. There is little storage for zinc in the human body, and bioavailable zinc from food or supplements must replenish the zinc pool on a regular basis. Alterations in zinc uptake, retention, or secretion can quickly lead to zinc deficiency and affect zinc-dependent functions. Zinc deficiency can significantly depress adaptive immune response, and susceptible individuals have impaired host defense and are at increased risk to opportunistic infections [8]. At the same time, zinc deficiency has been shown to increase systemic inflammation [13, 14].

Severe zinc deficiency mostly occurs in developing countries and is rare in the United States. On the other hand, marginal zinc deficiency is a potentially widespread problem, as a significant proportion of the US population does not consume adequate zinc. Approximately 12% of the US population does not consume the estimated average requirement (EAR) for zinc, but national surveys show that the prevalence of inadequate zinc intake is even higher among individuals above 50 years of age than any other population [15–18]; 40% of men and 45% of women consume less than the EAR [19]. Studies also show that zinc status, as shown by plasma zinc concentrations, declines with age [20–23]. However, the decline in status may not be solely attributed to decreases in dietary zinc intake. Utilization, distribution, and absorption of zinc may also be altered with age. For example, elderly women consuming zinc supplements did not exhibit increases in plasma zinc levels. In contrast, younger cohorts receiving the same zinc supplements had significant increases in plasma zinc [24]. Zinc deficiency in the elderly has a profound effect on the immune system, rendering them susceptible to infections and chronic inflammation [25, 26]. It is not known why plasma zinc levels fall with aging, but, some propose that it is related to a decline in zinc absorption. However, comparisons of zinc absorption between young and older individuals fail to show a consistent affect of age. Of the five studies done [27–31], three found lower zinc absorption in older than younger adults, two did not. Different study designs, methods, and diets likely account for the divergent results. In a study of the response to zinc loading (100 mg/d for 9 months), Wastney and co-workers found that total zinc absorption increased more in older than younger men with these high doses [32]. Studies of age and zinc absorption in experimental animals are also inconsistent [33, 34]. Thus, although plasma zinc concentrations are consistently lower in older individuals, there is no consistent evidence that this decline is due to a decrease in the ability to regulate zinc absorption. Regardless, there is mounting evidence that subclinical zinc deficiency may be more widespread in an elderly population and there may be an enhanced need for zinc with age. An alternative explanation for the decline in plasma zinc with age may be due to an enhanced uptake of zinc by cells due to altered expression of the zinc binding protein metallothionein (MT) and zinc transporters, resulting in tissue-specific zinc redistribution and/or changes in intracellular zinc pools.

Zinc transporters comprise a family of multiple transmembrane spanning domain proteins that are encoded by two solute-linked carrier (SLC) gene families: *SLC30* (a.k.a. ZnT) and *SLC39* (a.k.a. Zip). Evaluation of the human genome identified 24 human zinc transporters, consisting of 10 *SLC30* and 14 *SLC39* family members [35, 36]. In general, ZnT and Zip family members have opposing roles in regulating cellular zinc homeostasis. ZnT transporters reduce cytosolic zinc bioavailability by promoting zinc efflux in conditions of excess, while Zip transporters function by increasing cytosolic zinc during deficient states. Members of both families exhibit tissue and cell specific expression and possess differential responsiveness to dietary zinc as well as to physiologic stimuli including cytokines [35, 37]. The expression of zinc transporters has been profiled in immune cell types and their regulation plays an important role in specific immune cell activation. Cousins and colleagues evaluated the gene expression of zinc transporters in peripheral blood monocytes, T-lymphocytes, and granulocytes following modest dietary zinc supplementation in healthy young males [38]. They observed that zinc supplementation induced significant alterations in transcript abundance of individual transporter within different immune cell types and further, that considerable differences exist between immune cell populations (up to 270-fold). In particular, ZnT1 mRNA and metallothionein-1 (MT-1) expression increased with zinc supplementation, while expression of the importers Zip-1, -3 and -8 decreased. Most strikingly, zinc-mediated alteration in zinc transporter gene expression enhanced the response of different leukocyte subsets to relevant immune challenges. Specifically, activated monocytes, granulocytes, and T lymphocytes in zinc supplemented individuals had significantly increased TNF α , IL1 β , and IFN γ transcript levels, respectively. Overbeck et al similarly observed that intracellular zinc homeostasis in different leukocyte subsets is regulated by distinct patterns of zinc exporter expression, and postulated that this leads to different immune subsets having varying susceptibilities to zinc deficiency [39]. Further, recent data indicate that intracellular zinc signaling is involved in immune cells activation, and is regulated via changes in the expression of zinc transporters (to be discussed in further detail in next section) [40–42]. Thus alterations in zinc transporter expression are intimately linked to zinc homeostasis and immune function, and dysregulation of zinc transporter expression may contribute to low zinc status and immune dysfunction observed with aging. Zinc imbalance and perturbed expression of zinc transporters have also been associated with various age-related chronic diseases. Alterations of zinc transporters ZnT1, ZnT4, and ZnT6 have been reported in the brains of preclinical Alzheimer's disease subjects [43]. In diabetes, disease susceptibility is associated with reduced zinc status and zinc transporter dysregulation, and zinc supplementation is protective against both Type 1 and Type 2 diabetes [44]. In pancreatic islets, insulin production by β -cells is tightly regulated

ZnT8. It is postulated that changes in ZnT8 abundance or zinc transport activity will affect zinc accumulation as well as insulin storage and release, resulting in impaired glucose-induced insulin response and promote progression from glucose intolerance to diabetes [45]. Indeed, susceptibility to Type 2 diabetes is associated with ZnT8, and interestingly, ZnT8 is also a major autoantigen in autoimmune Type 1 diabetes [46, 47]. Thus alteration in the expression and regulation of zinc transporters is a potential contributing factor leading to age-associated decline in zinc status, immune dysfunction, and the development of age-related chronic diseases.

To date, little is known regarding the effects of age on the expression and function of zinc transporters. Age-related epigenetic changes has been proposed as one potential mechanism that may result in age-related dysregulation of zinc transporter expression with age (to be discussed in further detail in later section). Epigenetic alterations associated with aging includes a progressive decline in DNA methylation in cells and tissues (global DNA hypomethylation), coupled with site-specific promoter hypermethylation at select genes, resulting in gene silencing [48, 49]. It has been postulated that aberrant methylation of zinc transporter promoter regions may modulate the expression and function of select zinc transporters [50]. The effects of age on zinc transporter expression, particularly in immune tissues, remains to be investigated. This will help determine the underlying molecular mechanisms contributing to altered zinc metabolism during aging, and will clarify whether low zinc status observed with age results from diminished zinc absorption, and/or a redistribution of zinc to different tissues/cellular compartments due to aberrant zinc transporter expression and function.

3 Zinc in immune responses

It has been well established that zinc is necessary for the normal function of the immune system. The clinical manifestation of severe zinc deficiency is highlighted in patients suffering from acrodermatitis enteropathica (AE), a rare genetic disorder resulting in defective zinc uptake and severe zinc deficiency [51]. AE patients suffer from dermatitis, diarrhea, and growth retardation. In addition, AE patients have characteristic immune defects including thymic atrophy and lymphopenia. Both conditions lead to impaired host defense, resulting in recurrent viral, bacterial, and fungal infections [52]. Nutritional zinc deficiency results in similar immune dysfunction. For example, plasma zinc level in children is a predictor of diarrheal and respiratory morbidity [53]. Children with low plasma zinc concentration have increased diarrhea prevalence and severity, and have a higher mean prevalence rate of acute lower respiratory tract infections compared to children with normal plasma zinc. In hemodialysis patients, impaired immune response to Diphtheria vaccination is associated

with decreased serum zinc concentration [54]. In the elderly, patients with reduced zinc status have higher frequencies of respiratory infections [55–57] and gastrointestinal disease [58]. A multitude of defects in both adaptive and innate immunity contributes to impaired immune response resulting from zinc deficiency.

Changes in zinc status affect multiple immune cell types involved in both innate and adaptive immunity and has been extensively reviewed [9, 59]. Innate immunity is the immune mechanism that provides immediate but non-specific defense against pathogens and infections, and involves natural killer cells, mast cells, eosinophils, basophils, and phagocytic cells such as macrophages and neutrophils. In vivo, zinc deficiency has been shown to affect the functions of cells involved in innate immune response. For example, mouse peritoneal macrophages from zinc deficient mice have reduced phagocytosis and parasite killing capacities [60]. Zinc deficiency also reduces the activity of natural killer cells, and decreases chemotaxis and oxidative burst by neutrophils [12, 61]. In addition, zinc is involved in mast cell degranulation and cytokine production, and zinc depletion inhibits mast cell activation [62]. Zinc deficiency also affects immune cells involved in adaptive immunity. Induction of an adaptive immune response involves the recognition of antigen-specific lymphocytes to antigen, such as during viral infection or immunization, and the subsequent development of immunological memory [63, 64]. T and B cells are the primary mediators of adaptive immunity, and are involved in cell-mediated and antibody immune responses, respectively. Significant amount of work have demonstrated the importance of zinc in adaptive immunity [9]. Zinc deficiency causes a reduction in B cell lymphopoiesis resulting in reduced number of pre-B and immature B cells due to accelerated apoptosis, but the absolute numbers of mature B cells are not significantly affected. However, antibody response, particularly against T cell-dependent antigens is affected by zinc deficiency [65, 66]. Numerous defects are observed in the T cell compartment with zinc deficiency. T cell lymphopoiesis takes place in the thymus and is significantly affected by zinc deficiency. Thymic atrophy is a hallmark of zinc deficiency, caused by substantial loss of pre-T cells due to enhanced apoptosis [67]. The activity of thymulin, a hormone that regulates the differentiation of immature T cells in the thymus and the function of mature T cells in the periphery, is also decreased with zinc deficiency [68]. In the periphery, zinc deficient T cells have reduced cell proliferation upon mitogen stimulation, reduced precursor number and activity of CD8⁺ cytotoxic T cells, and impaired CD4⁺ T cells functions, resulting in altered T helper (Th) cells function, and an imbalance of Th1/Th2 cytokines secretion [69, 70]. While zinc deficiency results in the impairment of antibody-mediated and cell-mediated immune responses, increased autoreactivity has been observed in zinc deficient animals due to impaired immune tolerance and inefficient removal of autoreactive T cells [71].

Recent studies indicate that intracellular zinc homeostasis is critically involved in the signaling events in immune cells, and is regulated by changes in zinc transporter expression [42, 72]. Two classes of intracellular zinc signaling have been observed in immune cells, involving transcription-independent early zinc signaling in which intracellular zinc levels change rapidly (within minutes) upon stimulation, and transcription-dependent late zinc signaling where intracellular zinc levels are altered several hours after stimulation [10]. In T cells, the regulation of T cell receptor (TCR) signaling is controlled by activation-induced zinc influx, where cytoplasmic zinc concentration increases within 1 min after TCR engagement. Zinc influx is mediated by the zinc transporter Zip6, and Zip6 gene silencing inhibits T cell activation [42]. In monocytes, immediate increase in intracellular zinc was observed upon lipopolysaccharide (LPS) treatment, and preventing zinc influx with the zinc-specific chelator TPEN blocks LPS-induced monocyte activation [40]. The function of dendritic cells (DC), a potent antigen presenting cell that is involved in driving the inflammatory process, is also regulated by zinc homeostasis. In DC, a reduction in intracellular zinc mediated via downregulation of Zip6 and upregulation of ZnT1 expression, is required for upregulation of major histocompatibility complex class II and costimulatory molecules [41]. Alteration of zinc transporter expression and/or zinc homeostasis in immune cells may represent one mechanism by which age-associated zinc deficiency contributes to immune dysfunction. Mocchegiani and collaborators have postulated that intracellular zinc levels are compromised during aging due to increased expression of MT [73]. MT is a major storage protein for zinc, thus age-related increases in MT expression could result in increased sequestration of zinc. Taken together, zinc is intimately involved at multiple levels in the development and function of the immune system, including lymphopoiesis, the development of adaptive and innate immune response, and controlling intracellular signaling events and immune cell activation. It is noteworthy that immune dysfunction observed with age (thymic atrophy, reduced antibody and cellular immune responses, increased inflammation and autoimmunity) bear remarkable similarities to the hallmarks of zinc deficiency, and low zinc status associated with age is likely a contributing factor in age-related immune dysfunction.

4 Zinc as an anti-inflammatory agent

Similar to aging, the decline of adaptive and innate immunity with zinc deficiency is accompanied by a heightened proinflammatory response. In both in vitro and in vivo models, zinc deficiency has been associated with increases in inflammatory cytokines and other markers of inflammation. In an in vitro prostate cancer model, depletion of intracellular zinc increases the expression of

proinflammatory cytokines IL-6 and IL-8, and may contribute to tumor progression [74]. In a mouse model of polymicrobial sepsis, zinc deficiency increases systemic inflammation, resulting in increased organ damage and mortality [14]. In allergic airway inflammation, a murine model of asthma, zinc deficient mice have increased allergic inflammation, as determined by higher airway hyperresponsiveness and eosinophilia [75]. In addition, zinc deficiency also increases oxidative stress and DNA damage [76–82], which also play a role in promoting the inflammatory process. Zinc may modulate the proinflammatory response by targeting Nuclear Factor KappaB (NFκB), a transcription factor that is the master regulator of proinflammatory responses [83]. Activation of NFκB results in the phosphorylation and degradation of the inhibitory protein IκB mediated by IκB kinase, allowing nuclear translocation and subsequent nuclear binding of the NFκB complex to consensus sequences and the induction of gene transcription, including the production of proinflammatory cytokines. Both dietary zinc deficiency and the depletion of intracellular zinc with a zinc-specific chelator result in increased activation of NFκB, and the upregulation of NFκB-controlled proinflammatory cytokine expression [13, 74]. In aging, dysregulated NFκB signaling is proposed to be a culprit that result in age-related chronic inflammation [84, 85], and may potentially be exacerbated by low zinc status in the elderly.

While zinc deficiency results in systemic inflammation, zinc supplementation has been shown to have potent anti-inflammatory effects. For example, zinc supplementation significantly decreases proinflammatory immune cell infiltration into the lung during allergic inflammation, and normalizes Zip1 and Zip14 expression that are altered with acute inflammation [75]. In another study, short term zinc repletion reverses sepsis-induced proinflammatory response and reduces associated tissue damage in zinc deficient mice [14]. Moreover, zinc supplementation suppresses sepsis-induced NFκB activation [13]. The suppressive effects of zinc on NFκB activation is mediated by blocking IκB kinase activity and inhibiting the phosphorylation and degradation of IκB [86]. In addition, the ability of zinc supplementation to inhibit activation of NFκB may be related to the activation of A20, an endogenous negative regulator of NFκB [87, 88]. In humans, several zinc supplementation studies have shown to be associated with improved immune functions and decreased inflammation. For example, zinc supplementation in the elderly reduced the incidence of infections [23], increased delayed type hypersensitivity reaction [89, 90], and increased T cell function [91]. Moreover, zinc supplementation in humans exerts anti-inflammatory effects. Zinc supplementation has been associated with decreased inflammatory responses in populations susceptible to zinc deficiency, including sickle cell patients and in children [92, 93]. In the elderly, age-related increase in proinflammatory markers including C-reactive proteins and inflammatory cytokines, are corrected by zinc supplementation [94–96]. However, the beneficial effects of zinc

supplementation on immune function in the elderly have been inconsistent. Some of the discrepancies may be attributed to differences in zinc dose and timing (for review see [12, 97]). It has also been postulated that polymorphisms in inflammatory genes such as IL-6, a pro-inflammatory cytokine and regulator of MT expression, may alter responses to zinc supplementation in the elderly [97]. This research suggests that genetic variation in pro-inflammatory genes such as IL-6 renders individuals more susceptible to intracellular zinc loss and these are the subjects that may benefit the most from zinc supplementation [98]. Regardless, improving zinc status in the elderly population may be of significant benefits in improving overall immune function, and reducing age-related inflammation and associated morbidities.

5 Zinc, aging and epigenetics

It is clear that nutrient-gene interactions may exist that alter an individual's susceptibility to zinc deficiency and associated immune defects with age. In addition, there are possible epigenome alterations that may play a role in age-related zinc decline. The role of epigenetic alteration during the aging process has gained increasing attention in recent years. Epigenetics is the study of the regulation of gene activity that is not dependent on nucleotide sequence; this may include heritable changes in gene activity and expression, but also long-term alterations in the transcriptional potential of a cell that are not heritable [99]. Epigenetic phenomena include, but are not limited to, covalent modifications of histones, methylation of cytosines in DNA, and gene regulation by non-coding RNA. These features are potentially reversible and affect the expression of genes. For example, methylation of CpG islands in promoter elements is a major epigenetic controlling event for gene silencing [99–101]. In recent years, it has become increasingly clear that in addition to genetic changes, the epigenome is equally as critical as the DNA to controlling gene expression and healthy human development. In the context of the immune system, epigenetic modifications, including DNA methylation and histone modifications, have been shown to control immune function [102]. In particular, epigenetic modifications appear to regulate the differentiation of various immune cell populations critical in the development, regulation, and maintenance of immune response, including memory T cells [103], T-helper cells [104], regulatory T cells [105], and DC [106]. Aberrant DNA methylation can lead to hyperactivation or silencing of genes while histone modifications can result in either transcriptional activation or repression, depending on the histone mark. Accumulating evidence indicates epigenetic dysregulation is a common feature of aging, characterized by global DNA hypomethylation and gene-specific promoter hypermethylation or hypomethylation, as well as alteration in histone modifications [49, 107, 108]. Furthermore, alteration to the

epigenome has been proposed to be involved in the pathogenesis of various human diseases, including various age-related diseases such as chronic inflammation, autoimmunity, and cancer [3, 109–111]. For example, Agrawal et al recently demonstrated that epigenetic modifications in aged human DNA increase its immunogenicity and increases DC activation, and may be a potential mechanism leading to age-associated increase in autoimmune and proinflammatory responses [109]. In another study, promoter methylation status of IL-6 in rheumatoid arthritis patients has been shown to be significantly hypomethylated compared to control subjects, indicating that differential methylation may be involved in IL-6 regulation and disease pathogenesis [112]. In addition to age-related epigenetic modifications, specific nutrients can also modulate epigenetic regulation and alter disease susceptibility. Several studies revealed that maternal exposures to nutritional deficiencies lead to significant and persistent effects in their offspring. For example, studies using data from the WWII Dutch winter famine found that under-nutrition during the mid to late gestational periods was associated with increased risk of glucose intolerance and diabetes in middle age [113, 114]. Importantly, maternal zinc deficiency contributes to an increased risk for several chronic diseases, such as metabolic syndrome and diabetes later in life in the offspring [115, 116]. Marginal zinc deficiency in pregnant rats resulted in sustained defects in immune function in the offspring, that do not readily reverse with zinc supplementation and the defect can last through several generations [117]. Maternal zinc deficiency also caused persistent defects in insulin secretion and glucose intolerance in the offspring of rats. This work has prompted interest in the possibility that nutritional deficiencies, such as zinc, may cause epigenetic modifications resulting in “fetal programming” and increased susceptibility to disease.

The link of zinc deficiency to alterations in the epigenome is apparent with several epigenetic mechanisms. In particular, the strongest evidence is for the role of zinc in DNA methylation. Methylation of cytosines in CpG rich DNA is catalyzed by DNA methyltransferase (DNMT) enzymes. Emerging data suggests a requirement for zinc in pathways that both create and regulate DNA methylation and histone modification events. DNMTs, as well as chromatin modifying enzymes such as histone methyltransferase and histone deacetylases are zinc-containing enzymes [118, 119]. Zinc also acts as key cofactor in several enzymes involved in the methionine/transsulfuration pathway; a key pathway for generating methyl donation equivalents such as S-adenosyl-methionine and betaine. Betaine-homocysteine methyltransferase and methionine synthase are also zinc enzymes [120]. Serine hydroxymethyltransferase, a key enzyme in folate metabolism and helps transfer methyl units from serine into the methionine cycle is regulated by zinc-dependent transcription factors including metal-regulatory transcription factor 1 (MTF-1). Together this strongly suggests that zinc likely has an important function in

maintaining methylation status in the cell. Thus, zinc deficiency may cause a methyl deficiency, similar to other methyl donors like folate, resulting abnormal gene expression and developmental defects. In rats, zinc deficiency decreased turnover of S-adenosyl-methionine and depressed DNA and histone methylation in the liver [121]. In agouti mice, supplementation with methyl donors (choline, betaine, folate, B12, methionine) and zinc epigenetically regulated the expression of agouti in the offspring [122]. In addition, oxidative stress also can lead to alterations in DNA methylation [123]. Replacement of guanine with the oxygen radical adduct 8-hydroxyguanine markedly alters methylation of adjacent cytosines [124]. Interestingly, several zinc transporters such as ZnT5 and Zip8 contain CpG islands in their promoter elements and are susceptible to epigenetic regulation. For example, hypermethylation of the CpG island of Zip8 gene causes epigenetic silencing and down-regulation of Zip8 expression, resulting in a decrease in cadmium accumulation [125], and promoter methylation of ZnT5 is associated with reduced expression of an associated reporter gene [50]. Coneyworth et. al. have proposed that age-related methylation of ZnT5 contributes to the decline of zinc status with age and downstream health consequences [50]. It is possible that age-related epigenetic modifications, either with age itself or in combination with age-related zinc deficiency could impact key regulatory mechanisms that exacerbate intracellular zinc loss and immune dysregulation. In particular, epigenetic dysregulation of zinc transporters during the aging process may be one contributing factor resulting in age-related zinc deficiency. It is a likely possibility that age-related alterations in the epigenome contribute to age-related zinc decline, and together age and zinc deficiency cause additional epigenetic changes that result in immune dysfunction and other related age-related pathologies. Further delineation of age-related gene changes at the epigenome level, and their link to nutritional deficiencies is an emerging area of research.

6 Concluding remarks

Elderly individuals over 65 years of age represent 13% of the U.S. population; they are the most rapidly growing population segment. National surveys show that their dietary zinc intakes are inadequate. Coupled with impaired utilization of zinc with age, the elderly are a population highly susceptible to zinc deficiency and that many of their health problems are conditions associated with poor zinc nutriture. The importance of zinc in the normal development and function of the immune system and the negative impact of zinc deficiency on immune response is well established. Immune defects associated with aging, including immunosenescence and chronic inflammation, are remarkably similar to the hallmarks of zinc deficiency. Furthermore, imbalance in zinc homeostasis and zinc transporter dysregulation may be one of the underlying etiologies in

age-related diseases. Thus the decline in zinc status with age may play a significant role in immune dysfunction and the progression of age-related chronic diseases. However to date, the precise factors contributing to age-related zinc deficiency remain poorly defined. We postulate that alteration in the expression and regulation of zinc transporters, in particular epigenetic dysregulation, during the aging process is an important contributing factor towards age-associated decline in zinc status. As the elderly population increases, demands on the health care system and related costs will increase exponentially. The identification of the underlying molecular mechanisms by which zinc status alters immune function in the elderly population will aid in defining optimal zinc nutrition for this susceptible population and may help establish age-specific requirements for zinc, reduce disease burden, and improve the quality of life for the elderly.

7 References

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