

The Alteration of Zinc Transporter Gene Expression Is Associated with Inflammatory Markers in Obese Women

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Abstract Obesity, a chronic inflammatory state, is associated with altered zinc metabolism. ZnT and Zip transporters are involved in the regulation of zinc metabolism. This study examined the relationships among obesity, zinc transporter gene expression, and inflammatory markers in young Korean women. The messenger RNA (mRNA) levels of leukocyte zinc transporters between obese (BMI=28.3±0.5 kg/m², n=35) and nonobese (BMI=20.7±0.2 kg/m², n=20) women aged 18–28 years were examined using quantitative real-time polymerase chain reaction. Inflammatory markers, such as C-reactive protein (CRP), tumor necrosis factor-alpha (TNF-α), and interleukin (IL)-6, were measured in serum by enzyme immunoassay. *ZnT1* and *Zip1* were the most abundantly expressed zinc transporters in leukocytes. The mRNA levels of many zinc transporters (*ZnT4*, *ZnT5*, *ZnT9*, *Zip1*, *Zip4*, and *Zip6*) were significantly lower in obese women, and expression of these genes was inversely correlated with BMI and body fat percentage. In addition, inflammatory markers (CRP and TNF-α) were significantly higher in obese women. The

mRNA levels of *ZnT4*, *Zip1*, and *Zip6* were inversely correlated with CRP ($P<0.05$), and mRNA levels of *ZnT4* and *ZnT5* were inversely correlated with TNF-α ($P<0.05$). In standardized simple regression models, levels of TNF-α and CRP were negatively associated with mRNA levels of zinc transporters such as *ZnT4*, *ZnT5*, *Zip1*, and *Zip6* ($P<0.05$). These results suggest that the expression of zinc transporters may be altered in obese individuals. Changes in zinc transporters may also be related to the inflammatory state associated with obesity.

Keywords Obesity · Zinc transporters · Inflammatory markers

Introduction

Obesity is a worldwide public health problem. The prevalence of obesity has been increasing throughout the world, even in developing countries [1]. Obesity is a chronic inflammatory state characterized by altered adipokine production and increased levels of inflammatory cytokines. In obese individuals, plasma levels of inflammatory markers, such as C-reactive protein (CRP), tumor necrosis factor (TNF)-α, and interleukin (IL)-6, are elevated [2].

Zinc metabolism is also altered in obesity. Obese individuals have lower blood zinc concentrations than nonobese individuals [3, 4]. In obese women, erythrocyte zinc concentration is inversely associated with waist circumference and body mass index (BMI) [5]. Low serum or plasma zinc in obese individuals may be associated with the redistribution of zinc among various tissues [6–8]. In particular, a recent study showed that the reduction of body fat percentage by dietary intervention in obese adolescents led to an increase in the level of intraerythrocytic zinc and a decrease in urinary zinc

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excretion [9]. Zinc redistribution can be regulated by zinc transporters, which fall into two classes: zinc transporters or ZnT [also known as solute-linked carrier (SLC) 30] and Zin and Irt-like proteins, or Zip (also known as SLC39) [10]. ZnT transporters decrease intracellular zinc levels by transporting zinc ions from the cytoplasm to either the extracellular space or intracellular vesicles, whereas Zip transporters increase intracellular zinc levels through the reverse route. Members of the ZnT family and the Zip family (10 and 14, respectively) have been identified in various mammalian tissues [10]. It has been shown that zinc supplementation or depletion alters zinc transporter expression levels in cultured human leukocytes [11, 12] and leukocytes isolated from human blood [11–13]. However, whether zinc transporter expression is altered in obese individuals is not known, and potential mechanisms for such changes remain unclear. To our knowledge, only one study examined zinc transporter expression in obese subjects [14], but the study did not look into the relationship with inflammatory markers, which might be the possible mechanism for the shift of zinc transporters in obesity.

This study examined gene expression levels of zinc transporters between obese and nonobese Korean women and investigated the association of zinc transporter mRNA levels and inflammatory markers.

Methods

Study Design and Subjects

Nonobese and obese women aged 18–28 years (20 and 35, respectively) were recruited from the Daegu and Gyeongbuk areas. Participants answered questions about demographic characteristics, smoking, nutritional supplementation, personal medical history, family history, and medication use in a face-to-face interview. Exclusion criteria included vitamin–mineral or other nutritional supplements and any medications including oral contraceptives, smoking, or participation in a weight-loss program. Obesity [15] was defined as a BMI of ≥ 25 kg/m². Anthropometric measurements, blood pressure (BP), a fasting blood sample, and 24-h urine samples were collected. Also, 3-day diet records were obtained to assess usual dietary intake. Written informed consent was obtained from all participants. All procedures were approved by the Public Institutional Review Board (IRB; PIRB12-040-02) and the IRB of Kyung Hee University (KHSIRB-12-005).

Anthropometric Measurements and BP

All anthropometric measurements were performed by trained research staff using standardized protocols. Height was

measured by anthropometry (TKK-11252, Japan). Body weight and body fat were measured by bioimpedance analysis (Inbody 3.0, Biospace, South Korea). BMI was calculated as weight (kg) divided by height squared (m²). Waist circumference was measured at the midpoint between the lower border of the rib cage and the top of the iliac crest using stretch-resistant tape [16]. Systolic BP (SBP) and diastolic BP (DBP) were measured twice using an automatic sphygmometer (HEM-770A, Japan) after the subject had been resting for 10 min in a sitting position. Average SBP and DBP were used for the analysis [17].

Nutritional Assessment

To estimate usual dietary intake, a three-nonconsecutive-day dietary record including two weekdays and one weekend day was collected. At the first visit, detailed instructions for the 3-day dietary record were given to subjects by trained dietitians. The subjects were instructed to record the type and amount of all foods and beverages they consumed for 3 days. The dietary records were checked for completeness by trained staff. Daily energy and nutrient intakes were calculated using a dietary evaluation program (Can-pro 3.0, Korean Nutrition Society).

Biochemical Analyses

Fasting blood samples and 24-h urine samples were collected. Blood samples were collected using plastic syringes and put on ice for a maximum of 2 h, centrifuged at 1,500×g for 10 min at 4 °C (Allegra 6R, Beckman Colter, USA), and stored at –70 °C prior to analysis. Urine samples were collected in a polyethylene container, weighed, then stored in aliquots at –20 °C until analysis [18]. Serum and urinary zinc were measured using atomic absorption spectrometry (AAS 600, Perkin-Elmer, USA). Urinary creatinine was measured using the Jaffe method [19]. The activity of SOD was measured using an SOD assay kit (Cayman Chemical, USA). Inflammatory markers including CRP, TNF- α , and IL-6 were measured in serum by enzyme immunoassay. Specifically, CRP [20] was measured using a high-sensitivity CRP ELISA kit (Immundiagnostik, Europe). TNF- α [21] was measured using the Quantikine human TNF- α kit (R&D Systems, USA). IL-6 [22] was measured using the Quantikine human IL-6 kit (R&D Systems). Fasting triacylglyceride (TG) and total cholesterol (TC) were measured using an automated analyzer (ADVIA 2400, Japan). Fasting insulin was measured using a chemiluminescent immunoassay. Leptin concentration was measured using a human leptin kit (Linco Research, USA) [23], and adiponectin concentration was determined by immunoassay using a human adiponectin kit (R&D Systems).

Preparation of Leukocyte total RNA and Real-Time PCR Analysis

Total RNA was extracted from 1.5 ml whole blood using the QIAamp RNA blood mini kit (Qiagen, USA) [11]. The concentration and purity of extracted RNA was examined by measuring absorbance at 260- and 280-nm wavelength using a spectrophotometer (Nanodrop, Thermo, USA). RNA samples with OD₂₆₀/OD₂₈₀ values of 1.8 or higher were used in the analysis. cDNA was synthesized from 0.15 µg of total RNA using a PrimeScript™ RT reagent kit (Takara, Japan) according to the manufacturer's protocol. Zinc transporter gene expression levels in leukocytes were assessed via real-time polymerase chain reaction (PCR). Reactions were prepared using SYBR Premix Ex Taq II (Takara) and conducted on a Mini-Opticon (Bio-rad, USA) in duplicate. We examined the expression of a wide range of zinc transporters expressed in leukocytes [7, 12]. The relative expression levels of zinc transporters were determined using glyceraldehyde 3-phosphate dehydrogenase (GAPDH) as reference gene and expressed as $2^{-\Delta Ct}$ ($\Delta Ct = Ct_{\text{target}} - Ct_{\text{reference}}$) per 10^{-4} GAPDH [11].

Statistical Analysis

All data analyses were conducted using SAS 9.3 (SAS Institute, Cary, NC, USA). Differences in anthropometric measurements, zinc status, inflammatory markers, biochemical parameters, and mRNA levels of leukocyte zinc transporters between nonobese and obese women were identified using the Wilcoxon rank-sum test. Spearman correlations between mRNA levels of leukocyte zinc transporters and anthropometric measurements were also calculated. In addition, the relationships among mRNA levels of leukocyte zinc transporters, zinc status, inflammatory markers, and other biochemical parameters were examined using Spearman's partial rank correlation coefficients, with adjustment for BMI. Standardized regression analysis was conducted to explore the effects of BMI and inflammatory markers on mRNA levels of zinc transporters after calculation of *z* scores for each value of BMI and inflammatory markers.

Results

Differences in Anthropometric Measurements, Zinc Status, Inflammatory Markers, and Biochemical Parameters between Nonobese and Obese Women

Anthropometric measurements, zinc status, inflammatory markers, and biochemical parameters are shown in Table 1, stratified by obesity status. BMI, percentage body fat, and waist circumference were significantly higher in obese women ($P < 0.0001$). SBP and DBP were also significantly higher

in obese women ($P < 0.01$). Dietary zinc intake and serum and urinary zinc concentrations were not significantly different between the two groups. Serum SOD activity was significantly lower in obese women ($P = 0.02$), while inflammatory markers, such as CRP ($P < 0.0001$) and TNF α ($P < 0.001$), were significantly higher in obese women. The levels of serum TG, TC, and insulin were significantly higher in obese women as well. Finally, adiponectin was significantly lower ($P = 0.04$) and leptin was significantly higher ($P < 0.0001$) in obese women.

Zinc Transporter mRNA Levels in Nonobese and Obese Women

Zinc transporter mRNA levels in leukocytes are shown in Fig. 1. The zinc transporters with the highest expression levels were *ZnT1* and *Zip1*. The differences in zinc transporter mRNA levels between nonobese and obese women are shown in Table 2. The mRNA levels of zinc transporters, such as *ZnT4*, *ZnT5*, *ZnT9*, *Zip1*, *Zip4*, and *Zip6*, were significantly lower in obese women. *Zip1* mRNA levels were 70.6 % lower in obese women ($P < 0.0001$). By contrast, *ZnT7* mRNA levels were 78.1 % higher in obese women ($P < 0.0001$).

Association of Zinc Transporter Expression Levels with Anthropometric Measurements, Zinc Status, Inflammatory Markers, and Biochemical Parameters

The correlations among leukocyte zinc transporters, anthropometric measurements, zinc status, inflammatory markers, and biochemical parameters are shown in Table 3. The mRNA levels of zinc transporters, such as *ZnT4*, *ZnT5*, *ZnT9*, *Zip1*, *Zip4*, and *Zip6*, which were significantly lower in obese women, were inversely correlated with BMI and percentage body fat. By contrast, *ZnT7* mRNA levels, which were significantly higher in obese women, were positively correlated with BMI and percentage body fat ($P < 0.001$). Zinc transporter mRNA levels were not correlated with either dietary zinc intake or serum zinc concentration. However, the mRNA levels of *ZnT9* and *Zip6* were positively correlated with urinary zinc concentration ($P < 0.05$).

Even after adjustment for BMI, inflammatory markers were significantly associated with the mRNA levels of zinc transporters that had lower expression in obese women. The levels of *ZnT4*, *Zip1*, and *Zip6* mRNA were inversely correlated with CRP ($P < 0.05$), and the mRNA levels of *ZnT4* and *ZnT5* were inversely correlated with TNF- α ($P < 0.05$). However, mRNA levels of other zinc transporters were not associated with inflammatory markers (data not shown).

After adjustment for BMI, TG, and TC were not correlated with zinc transporter mRNA levels. Similarly, serum leptin, adiponectin, and insulin showed no correlation with zinc transporter expression after adjustment for BMI.

Table 1 Anthropometric measurements, zinc status, inflammatory markers, and biochemical parameters of subjects

	Nonobese (<i>n</i> =20)			Obese (<i>n</i> =35)			<i>P</i> values ^a
	Mean	SEM	(Range)	Mean	SEM	(Range)	
<i>Anthropometric measurements</i>							
Age (year)	20.7	0.4	(19–24)	20.9	0.4	(18–28)	NS
Waist circumference (cm)	69.9	0.9	(65.0–78.0)	89.8	1.5	(74.0–118.0)	<0.0001
BMI (kg/m ²)	20.7	0.2	(18.7–22.7)	28.3	0.5	(25.6–39.1)	<0.0001
Body fat (%)	14.8	0.6	(8.7–20.6)	28.0	0.9	(22.3–42.9)	<0.0001
SBP (mmHg)	108.8	2.3	(87.0–124.5)	118.2	1.6	(100.0–135.0)	<0.01
DBP(mmHg)	69.9	1.7	(58.0–82.0)	78.1	1.2	(64.0–89.5)	<0.01
<i>Dietary intake</i>							
Energy (kcal)	1490.0	97.2	(913.5–2,679.5)	1700.8	78.0	(1,002.7–2,571.4)	0.08
Zinc intake (mg/day)	7.0	0.8	(2.5–16.6)	7.4	0.4	(3.5–14.2)	NS
<i>Zinc status indicators</i>							
Serum zinc (μmol/l)	13.7	0.5	(9.6–17.3)	13.2	0.4	(8.1–17.3)	NS
Urine zinc (μg/mg creatinine)	0.45	0.05	(0.18–0.89)	0.36	0.03	(0.12–0.91)	NS
SOD (U/ml)	12.1	1.7	(4.9–30.3)	7.6	0.3	(1.0–12.7)	0.02
<i>Inflammatory markers</i>							
hs-CRP (nmol/l)	2.4	0.2	(1.0–3.9)	26.9	6.3	(0.4–157.6)	<0.0001
TNFα (pg/ml)	9.4	0.8	(3.2–17.9)	14.8	1.1	(3.4–36.0)	<0.001
IL-6 (pg/ml)	0.71	0.17	(0–2.4)	0.97	0.21	(0–3.9)	NS
<i>Biochemical parameters</i>							
Triacylglycerol (mmol/l)	0.72	0.07	(0.33–1.67)	1.27	0.2	(0.58–9.05)	<0.01
Total cholesterol (mmol/l)	4.2	0.1	(3.1–5.4)	4.7	0.2	(2.8–6.9)	<0.01
Insulin (pmol/l)	35.9	4.2	(6.8–80.2)	87.7	7.6	(17.6–193.4)	<0.0001
Adiponectin (μg/ml)	7.9	1.3	(1.0–17.7)	4.6	0.6	(0.7–13.8)	0.04
Leptin (μg/l)	7.1	0.7	(2.8–15.0)	19.8	1.0	(6.1–34.7)	<0.0001

NS not significant, SOD superoxide dismutase, ALP alkaline phosphatase, hs-CRP high sensitivity C-reactive protein, TNF-α tumor necrosis factor-alpha, IL-6 interleukin-6

^a The differences of variables between nonobese women and obese women were examined by Wilcoxon rank-sum test

For further analysis, the results of standardized simple regression analyses to examine the influence of BMI and

inflammatory markers on mRNA levels of zinc transporters are shown in Table 4. BMI and inflammatory markers were

Fig. 1 The mRNA levels of leukocyte zinc transporters in total subjects (*n*=55)

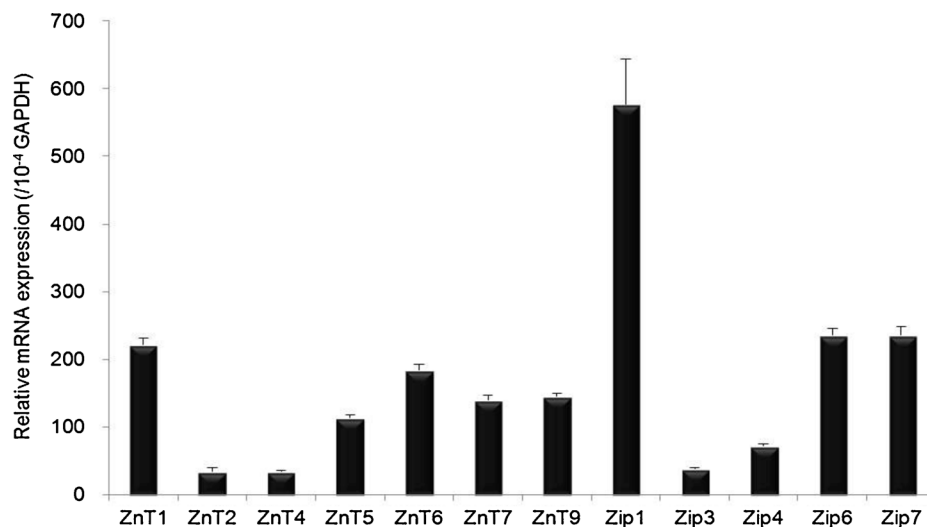


Table 2 The mRNA levels of leukocyte zinc transporters between nonobese women and obese women^a

	Nonobese (n=20)		Obese (n=35)		P values ^b
	Mean	SEM	Mean	SEM	
ZnT1	212.1	25.8	227.2	7.3	NS
ZnT2	41.8	13.1	30.0	3.9	NS
ZnT4	49.5	3.3	25.3	1.1	<0.0001
ZnT5	153.2	7.6	90.3	2.1	<0.0001
ZnT6	189.2	18.6	181.1	9.2	NS
ZnT7	93.1	5.9	165.9	9.2	<0.0001
ZnT9	162.6	9.2	134.6	5.6	<0.01
Zip1	1046.4	127.0	307.8	12.5	<0.0001
Zip3	34.7	3.7	40.6	2.1	NS
Zip4	89.5	5.1	61.6	1.9	<0.0001
Zip6	269.8	18.4	215.3	10.8	0.02
Zip7	216.7	26.5	244.9	12.9	NS

NS not significant

^a The levels of zinc transporters mRNA were examined using GAPDH as reference gene (per 10⁻⁴ GAPDH)

^b The significant differences between nonobese women and obese women were examined by Wilcoxon rank-sum test

inversely associated with mRNA level of zinc transporters. BMI had the negative association with mRNA level of *ZnT4* (standardized $\beta=-0.29$, $P<0.0001$), *ZnT5* (standardized

$\beta=-0.20$, $P<0.0001$), and *Zip1* (standardized $\beta=-0.43$, $P<0.0001$). The level of TNF- α was negatively associated with mRNA level of *ZnT4* (standardized $\beta=-0.24$, $P<0.0001$), *ZnT5* (standardized $\beta=-0.15$, $P<0.001$), and *Zip1* (standardized $\beta=-0.30$, $P<0.001$), even though they were a little smaller than those of BMI. Furthermore, TNF- α level was negatively associated with mRNA level of *ZnT4* (standardized $\beta=-0.12$, $P=0.03$) even after adjustment for BMI in a standardized multiple regression model (data not shown). In addition, CRP had the largest negative association with mRNA level of *Zip6* (standardized $\beta=-0.11$, $P=0.01$).

Discussion

In this study, we found that zinc transporter expression was altered in the leukocytes of obese women. Leukocyte mRNA levels of *ZnT4*, *ZnT5*, *ZnT9*, *Zip1*, *Zip4*, and *Zip6* were significantly lower in obese women, and expression of these zinc transporters was inversely correlated with BMI and body fat. Furthermore, expression levels of some zinc transporters (*ZnT4*, *ZnT5*, *Zip1*, and *Zip6*) were inversely associated with inflammatory markers such as TNF- α and CRP. These results suggest that zinc metabolism is altered in obese individuals via changes in the expression of various zinc transporters; such changes in zinc metabolism could be related to the elevated level of inflammation found in obesity.

Table 3 Spearman's correlation coefficients among the mRNA levels of leukocyte zinc transporters, anthropometric measurements, zinc status, inflammatory markers, and biochemical parameters in subjects (n=55)

	ZnT4	ZnT5	ZnT7	ZnT9	Zip1	Zip4	Zip6
<i>Anthropometric measurements</i>							
BMI (kg/m ²)	-0.70***	-0.59***	0.57***	-0.31***	-0.65***	-0.50***	-0.27*
Body fat (%)	-0.67***	-0.60***	0.58***	-0.29***	-0.61***	-0.53***	-0.20
<i>Zinc status^a</i>							
Zinc intake (mg/day)	0.02***	-0.02***	0.11***	0.03***	0.18***	0.08***	0.16
Serum zinc (μ mol/l)	-0.04***	-0.11***	0.00***	-0.04***	0.12***	0.06***	0.00
Urine zinc (μ g/mg creatinine)	0.24***	0.26***	0.21***	0.32***	0.07***	0.18***	0.31*
SOD (U/ml)	-0.02***	0.09***	-0.16***	0.00***	0.15***	0.10***	-0.08
<i>Inflammatory markers^a</i>							
hs-CRP (nmol/l)	-0.30***	-0.23***	-0.08***	-0.10***	-0.32***	0.04***	-0.29*
TNF α (pg/ml)	-0.30***	-0.28***	-0.11***	-0.05***	-0.25***	-0.01***	-0.11
IL-6 (pg/ml)	0.19***	0.10***	-0.03***	0.16***	0.20***	0.22***	0.07
<i>Biochemical parameters^a</i>							
Triacylglycerol (mmol/l)	-0.03***	-0.13***	0.27***	-0.11***	-0.17***	0.03***	0.05
Total cholesterol (mmol/l)	0.10***	0.01***	0.22***	-0.19***	-0.19***	0.16***	0.01
Insulin (pmol/l)	0.01***	0.07***	0.00***	0.00***	-0.19***	0.18***	-0.01
Adiponectin (μ g/ml)	-0.05***	-0.01***	-0.05***	-0.02***	0.01***	0.03***	-0.02
Leptin (μ g/l)	-0.20***	-0.18***	0.21***	0.05***	-0.03***	-0.01***	0.12

SOD superoxide dismutase, hs-CRP high sensitivity C-reactive protein, TNF α tumor necrosis factor-alpha, IL-6 interleukin-6

^a Adjusted for BMI by Spearman's partial correlation analysis * $P<0.05$, *** $P<0.001$

Table 4 Standardized simple regression models of predicting mRNA levels of zinc transporters with BMI and inflammatory markers in total subjects ($n=55$)

Dependent	Parameter	R^2	Standardized β	SE	P values	Dependent	Parameter	R^2	Standardized β	SE	P values
ZnT4	BMI	0.45	-0.29	0.04	<0.0001	Zip1	BMI	0.43	-0.43	0.07	<0.0001
	TNF- α	0.31	-0.24	0.05	<0.0001		TNF- α	0.21	-0.30	0.08	<0.001
	hs-CRP	0.17	-0.18	0.05	<0.01		hs-CRP	0.12	-0.23	0.09	0.01
ZnT5	BMI	0.42	-0.20	0.03	<0.0001	Zip6	BMI	0.07	-0.08	0.04	NS
	TNF- α	0.23	-0.15	0.04	<0.001		TNF- α	0.08	-0.09	0.04	0.04
	hs-CRP	0.14	-0.12	0.04	<0.01		hs-CRP	0.12	-0.11	0.04	0.01

NS not significant, TNF α tumor necrosis factor-alpha, hs-CRP high sensitivity C-reactive protein

The zinc transporters with the highest level of expression in leukocytes were *ZnT1* and *Zip1*. Previous studies report similar findings. Aydemir et al. [13] reported that in the monocytes of three young, healthy men aged 19–31 years, mRNA levels of *ZnT1* and *Zip1* were much higher than those of all other zinc transporters examined (*ZnT5*, *ZnT7*, *Zip3*, and *Zip8*). Another study, by Overbeck et al. [12], also found that, among all *ZnT* proteins, *ZnT1* mRNA was the most highly expressed in peripheral blood mononuclear cells from seven young, healthy donors.

In our study, with the exception of *ZnT7*, expression levels of zinc transporters were lower in obese women. Moreover, expression of these zinc transporters was inversely correlated with BMI and body fat, suggesting zinc metabolism might be altered in obese subjects. Several studies have demonstrated altered zinc status in obese individuals [4, 5, 24]. For example, among 73 premenopausal women aged 20–50 years, erythrocyte zinc concentration was lower in obese women, displaying an inverse association with BMI and waist circumference [5]. Similarly, among 48 children aged 5–15 years [25], monocyte zinc concentration was lower in obese individuals, showing a significant inverse correlation with body weight and arm fat area. In dietary intervention study through nutritional education about balanced diet with 15 obese adolescent girls aged 14–18 years, the decreased body fat resulted in the increased levels of erythrocyte zinc with no change of zinc intake. That is, the change of zinc status in obese was affected by the proportion of body fat, not zinc intake [9]. The altered zinc status reported in obesity could be related to the changes in zinc transporter expression levels we identified in leukocytes. To the best of our knowledge, only one prior study [14] has examined zinc transporter expression in obese subjects. It compared the expression of eight *ZnT* (*ZnT* 1–8) and eight *Zip* (*Zip* 1–8) levels in the adipose tissue of 12 obese women (mean BMI 44.5 ± 5.3 kg/m²) and 12 nonobese women (mean BMI 23.1 ± 2.6 kg/m²). Consistent with our findings, mRNA levels of *ZnT2*, *ZnT3*, *ZnT6*, and *ZnT8* were significantly lower in the subcutaneous fat of obese women. Moreover, mRNA levels of all *Zip* transporters examined (*Zip* 1–8) were significantly lower in the subcutaneous fat of obese women,

indicating that changes in zinc transporter expression are an important feature of obesity.

We found that, unlike other *ZnT* transporters, *ZnT7* mRNA levels were higher in the leukocytes of obese individuals. Huang et al. [26] found low weight gain and fat mass in *ZnT7*-knockout mice, suggesting *ZnT7* is involved in the regulation of body composition including fat mass. In the present study, *ZnT7* mRNA levels were positively correlated with BMI and percentage body fat. Therefore, an increase in body fat mass may be associated with elevated expression of *ZnT7* in obese women.

Importantly, we found that mRNA levels of some zinc transporters (*ZnT4*, *ZnT5*, *Zip1*, and *Zip6*) were inversely correlated with inflammatory markers. This result suggests that obesity-related inflammation could be linked to alterations in zinc transporter expression. In the present study, mRNA levels of *ZnT4* and *ZnT5* were significantly lower (~50 %) in obese women and were significantly inversely correlated with serum TNF α and/or CRP. Consistent with our findings, a marked downregulation of *ZnT4* and *ZnT6* has been reported in mouse models of allergic inflammation [27]. The *ZnT* family primarily functions as a zinc exporter protein, decreasing cytosolic zinc concentrations [10, 28]. For example, in mammary and intestinal epithelial cells, a mutation in *ZnT4* can result in reduced expression, leading to decreased zinc secretion [29, 30]. In light of the anti-inflammatory effects of zinc, it is possible that reduced expression of *ZnT4* and *ZnT5* might exert a protective role against inflammation by increasing retention of intracellular zinc [31, 32]. By contrast, expression of *ZnT1* and *ZnT2* was not associated with inflammatory markers in this study. These transporters are chiefly responsible for intracellular compartmentalization, playing a crucial role in the delivery of zinc to zinc-dependent proteins important for cell function such as SOD and alkaline phosphatase [30]. It is likely that expression of these crucial transporters is not downregulated by obesity-related inflammation.

The mechanism by which obesity-related inflammation is associated with *Zip1* and *Zip6* expression is unclear. Some *Zip* proteins are located in the *trans*-Golgi network, where they are

involved in the influx of zinc from the Golgi apparatus to the cytoplasm [33]. Thus, the downregulation of these Zip proteins could facilitate the conservation of intracompartmental zinc. Similar to our findings, Zip6 is downregulated in response to LPS treatment in dendritic cells [34]. Zip1 and Zip6 can also be located in the plasma membrane, where they are involved in zinc influx from the extracellular space. In this case, decreased *Zip* expression could lead to a reduction in intracellular zinc concentrations. Further studies are needed to better understand the relationship between changes in *Zip* expression and cellular zinc concentration in obesity.

Although serum or urine zinc concentration did not differ between obese women and nonobese women in the present study, the activity of superoxide dismutase, an index of zinc status, was significantly lower in obese women than in nonobese women. No changes in serum zinc concentration may be due to zinc homeostasis. Serum concentrations are maintained stable within a narrow range even when dietary zinc levels fluctuate [35].

The present study has several limitations. First, zinc transporter protein levels were not determined. While a number of *ZnT/Zip* transporter genes are transcriptionally regulated [36], posttranscriptional and translational regulation cannot be ruled out and should be investigated. Second, the study cannot reveal causal relationships because of its cross-sectional design. A longitudinal study of obesity and zinc transporter expression, with a larger sample size, would be helpful in clarifying the direction of causality.

In conclusion, the present study clearly demonstrated that expression of multiple zinc transporters was significantly lower in the leukocytes of obese women. In addition, expression of several zinc transporters (*ZnT4*, *ZnT5*, *Zip1*, and *Zip6*) were inversely associated with inflammatory markers. These findings suggest zinc transporters as an important link between altered zinc metabolism and inflammation in obesity. Further intervention studies are warranted to fully elucidate the role of zinc transporters in modulating zinc metabolism and inflammation associated with obesity.

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