Disruption of the mouse Slc39a14 gene encoding zinc transporter ZIP14 is associated with decreased bone mass, likely caused by enhanced bone resorption

Sun Sasaki1, Manami Tsukamoto1, Masaki Saito1, Shintaro Hojyo2, Toshiyuki Fukada3,4,5, Masamichi Takami6 and Tatsuya Furuichi1,7

1 Laboratory of Laboratory Animal Science and Medicine, Co-Department of Veterinary Medicine, Faculty of Agriculture, Iwate University, Morioka, Japan
2 Osteoimmunology, Deutsches Rheuma-Forschungszentrum, Berlin, Germany
3 Faculty of Pharmaceutical Sciences, Tokushima Bunri University, Tokushima, Japan
4 Department of Pathology, School of Dentistry, Showa University, Tokyo, Japan
5 RIKEN Center for Integrative Medical Sciences, Yokohama, Japan
6 Department of Pharmacology, School of Dentistry, Showa University, Tokyo, Japan
7 Department of Basic Veterinary Science, United Graduate School of Veterinary Science, Gifu University, Gifu, Japan

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Correspondence
T. Furuichi, Laboratory of Laboratory Animal Science and Medicine, Co-Department of Veterinary Medicine, Faculty of Agriculture, Iwate University, 3-18-8 Ueda, Morioka, Iwate 020-8550, Japan; Department of Basic Veterinary Science, United Graduate School of Veterinary Science, Gifu University, 1-1 Yanagido, Gifu 501-1193, Japan
Fax: +81-19-621-6222
Tel: +81-19-621-6222
E-mail: furuichi@iwate-u.ac.jp

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Osteoclasts are bone-resorbing cells that play an essential role in maintaining bone homeostasis. Zinc (Zn) has been reported to inhibit osteoclast-mediated bone resorption, but the mechanism of this action has not been clarified. Zn homeostasis is tightly controlled by the coordinated actions of many Zn transporters. The Zn transporter ZIP14/Slc39a14 is involved in various physiological functions; hence, Zip14-knockout (KO) mice exhibit multiple phenotypes. In this study, we thoroughly investigated the bone phenotypes of Zip14-KO mice, demonstrating that the KO mice exhibited osteopenia in both trabecular and cortical bones. In Zip14-KO mice, bone resorption was increased, whereas the bone formation rate was unchanged. Zip14 mRNA was expressed in normal osteoclasts both in vivo and in vitro, but receptor activator of NF-κB ligand (RANKL)-induced osteoclastogenesis was not impaired in bone marrow-derived macrophages prepared from Zip14-KO mice. These results suggest that ZIP14 regulates bone homeostasis by inhibiting bone resorption and that in Zip14-KO mice, bone resorption is increased due to the elimination of this inhibitory regulation. Further studies are necessary to conclude whether the enhancement of bone resorption in Zip14-KO mice is due to a cell-autonomous or a non-cell-autonomous osteoclast defect.

Zinc (Zn) is an essential trace element for various biological activities [1] as it is needed for more than 300 enzymes and it serves as a structural component of at least 3000 proteins in the body. Recently, the Zn ion (Zn^{2+}) has been reported to act as a second messenger that regulates intracellular signal transduction in various cell types [2]. The Zn concentration is relatively high in bone and cartilage [3], and Zn deficiency delays skeletal growth and decreases bone mass [4,5]. These findings indicate that Zn homeostasis is important for skeletal development and maintenance.

Abbreviations
BMM, bone marrow-derived macrophage; IL, interleukin; KO, knockout; LPS, lipopolysaccharide; M-CSF, macrophage colony-stimulating factor; NFATC1, nuclear factor-activated T cells, cytoplasmic, calcineurin-dependent 1; RANKL, receptor activator of NF-κB ligand; sIL-6R, soluble IL-6 receptor; TGF, transforming growth factor; TRAP, tartrate-resistant acid phosphatase.
During adulthood, bone mass is maintained by the coupled activities of osteoblasts and osteoclasts [6,7]. In this physiological process, old bone is resorbed by osteoclasts and then replaced by new bone formed by osteoblasts. An imbalance between these processes leads to bone metabolic diseases, such as osteoporosis and occasionally osteopetrosis. Several in vitro studies illustrated that Zn stimulates osteoblast-mediated bone formation and inhibits osteoclast-mediated bone resorption, thereby positively regulating bone mass [8,9]. However, the details of these Zn-mediated regulatory mechanisms have not yet been clarified.

Zn homeostasis is tightly controlled by two major families of Zn transporters: SLC39s/ZIPs and SLC30s/ZnTs [10,11]. ZIP transporters promote Zn influx from extracellular fluid or intracellular vesicles into the cytoplasm, whereas ZnT transporters promote Zn efflux from cells or influx into intracellular vesicles from the cytosol. At least 14 ZIP and 10 ZnT transporters have been identified in mammals. Among them, two Golgi-localized transporters, namely ZIP13 and ZnT5, have been reported to positively regulate osteoblast differentiation and/or function [12–14]. ZIP13 is expressed in osteoblasts, and Zip13-knockout (KO) mice display decreased bone mass and bone formation rate. Furthermore, primary osteoblasts isolated from Zip13-KO mice exhibited impaired expression of the osteoblast marker genes. Znt5-KO mice also display decreased bone mass and bone formation rate, and alkaline phosphatase and mineralization activities are diminished in primary Znt5-KO osteoblasts. The plasma membrane Zn transporter, ZIP1, is reported to be expressed in osteoclasts [15]. Adenoviral overexpression of ZIP1 in osteoclasts reduces bone resorption activity in vitro, suggesting that ZIP1 negatively regulates osteoclast function.

ZIP14/Slc39a14 localizes to cell membranes and promotes Zn influx into cells [16,17]. It has been reported that ZIP14 is expressed ubiquitously and it transports other metals, such as manganese (Mn), iron (Fe), and cadmium (Cd), in addition to Zn [17–19]. In accordance with these findings, Zip14-KO mice have been reported to exhibit multiple phenotypes, including dwarfism, osteopenia, altered glucose homeostasis, low-grade chronic inflammation, and increased body fat [20–23]. In skeletal tissues, ZIP14 is expressed in the growth plate chondrocytes, in which it regulates differentiation [14,20]. In this study, we comprehensively examined the bone phenotypes of Zip14-KO mice. We focused on the ZIP14 function as a Zn transporter because Zn is more important for skeletal development and homeostasis than other metals transported by ZIP14. Zip14-KO mice exhibited an osteopenia phenotype accompanied by enhanced bone resorption, and ZIP14 was expressed in normal osteoclasts. These findings strongly suggest that ZIP14 regulates bone homeostasis by affecting osteoclast-mediated bone resorption.

Materials and methods

Experimental animals

Zip14-KO mice were generated as described previously and maintained on a C57BL/6 background [20]. Mice were housed in a temperature-controlled room with a 12-h/12-h light/dark cycle. Mice had free access to water, and they were fed standard mouse laboratory chow. Genotyping of mice was performed at 4–5 weeks of age by PCR as described previously [20]. All of the animal experiments were performed according to a protocol approved by Iwate University’s Committee on Animal Research and Ethics (Approval Numbers: 201214 and 201512).

X-ray and pQCT analyses

Femurs were dissected from sacrificed mice and fixed with ethanol. Radiographs were obtained using a TRS-1005 soft X-ray apparatus (Sofron, Tokyo, Japan). Femoral cortical bone quality was measured by pQCT analysis using an XCT Research SA+ computed tomography system (Stratec Medizintechnik GmbH, Pforzheim, Germany).

Histological analyses

To assess dynamic histomorphometric indices, 6-week-old mice were injected twice with calcein (15 mg kg⁻¹, i.p.) at 1 and 4 days before sacrifice. Tibiae were fixed with ethanol, and the undecalcified bones were embedded in glycol methacrylate. Sections (3 μm thick) were cut longitudinally in the proximal region of the tibia and stained with toluidine blue O. Histomorphometry was performed using a semiautomatic image analyzing system (Osteoplan II; Carl Zeiss, Thornwood, NY, USA) linked to a light microscope. The histomorphometric measurements were performed at 400 times using a minimum of 17–20 optical fields in the secondary spongiosa area from the growth plate– metaphyseal junction. Nomenclature, symbols, and units were used as recommended by the Nomenclature Committee of the American Society for Bone and Mineral Research. For in situ hybridization, paraffin sections (4–6 μm thick) were prepared from the tibiae of mice at 4 weeks of age. The digoxigenin-labeled RNA probe for Zip14 was used in line with a method described by GEN-OSTAFF CO., LTD (Tokyo, Japan). The sections were counterstained with Kernchrot stain solution. The probe sequences and hybridization conditions are available upon request.
**In vitro assay for osteoclastogenesis**

To prepare bone marrow-derived macrophages (BMMs), bone marrow cells were collected from the tibiae and femurs of mice at 6–10 weeks of age and cultured for 16 h in αMEM containing 10% FBS, 50 ng·mL⁻¹ macrophage colony-stimulating factor (M-CSF), and 1 ng·mL⁻¹ transforming growth factor (TGF)-β1 in cell culture dishes. M-CSF (Leukoprol) and TGF-β1 were purchased from Kyowa Hakko Kogyo (Tokyo, Japan) and R&D Systems (Minneapolis, USA), respectively. Supernatants were transferred to Petri dishes and cultured for 72 h. Adherent cells were collected and used as BMMs. For osteoclast formation assays, BMMs (4 × 10⁴ cells per well) were cultured for 3–4 days in 48-well culture plates in the presence of 50 ng·mL⁻¹ M-CSF, 1 ng·mL⁻¹ TGF-β1, and 20 or 100 ng·mL⁻¹ receptor activator of NF-κB ligand (RANKL; R&D Systems). After cultivation, tartrate-resistant acid phosphatase (TRAP) staining and TRAP activity assays were performed as described previously [24]. To evaluate bone-resorbing activity, areas of pits eroded by osteoclasts were measured. BMMs (1.5 × 10⁵ cells per well) were seeded onto Corning OsteoAssay Surface 96-well plates (Corning Incorporated, Corning, NY, USA) in the presence of 50 ng·mL⁻¹ M-CSF, 1 ng·mL⁻¹ TGF-β1, and 100 ng·mL⁻¹ RANKL for 7 days. The culture medium was changed every 3 days. Cells were removed using 5% sodium hypochlorite, followed by washing with distilled water and air-drying. Resorption pits were visualized under a scanning electron microscope, and the resorption area was quantified using ImageJ software (National Institutes of Health, Bethesda, MD, USA).

**RT-PCR**

Total RNA was extracted from BMMs at the indicated time points after 100 ng·mL⁻¹ RANKL treatment using ISOGEN (Nippon Gene, Tokyo, Japan). First-stand cDNA was synthesized using TaqMan Multiscribe Reverse Transcriptase (Applied Biosystems, Foster City, CA, USA) and subjected to amplification using Ex Taq polymerase (TaKaRa, Tokyo, Japan) and the following specific PCR primers: 5’-CAGAGGGCTTTTGCTTCAC-3’ and 5’-CA GACACGTGAAGGAGC-3’ for Zip14; 5’-ACTCTCT GGGATCAACGTCAC-3’ and 5’-GATAGCAGGACAGG GCCAGA-3’ for Oscl; 5’-CCTGCCCATAACCTGGA G-3’ and 5’-TACACCTTTCATCCTGGCCC-3’ for Ctsk; and 5’-AANCTGGGACGACATGGGAAAG-3’ and 5’-GG GGTGTTGGAAGGCTCTAA-3’ for Gapdh. The PCR conditions are available upon request.

**Measurement of serum TRAP levels**

Tartrate-resistant acid phosphatase is used as a serum marker for bone resorption. Serum TRAP levels were determined using a mouse TRAP™ Assay kit (Immunodiagnostic Systems, Boldon, UK).

**Statistical analysis**

Student’s t-test was used to determine the significance of differences between control (Ctrl) and Zip14-KO mice. Significance was defined as P < 0.05.

**Results**

**Both trabecular and cortical bone levels are decreased in Zip14-KO mice**

To examine the role of ZIP14 in bone homeostasis, we compared the bone phenotypes between Ctrl and Zip14-KO mice. Because the phenotypic abnormalities were inherited recessively in Zip14-KO mice and heterozygous Zip14-KO mice were normal [20–23], the Ctrl group consisted of both wild-type and heterozygous Zip14-KO mice. In preliminary studies, we performed X-ray analyses using the distal femurs of both male and female mice at 6 weeks of age. It appeared that Zip14-KO mice were more radiolucent than Ctrl mice, and the increased radiolucency was comparable between male and female Zip14-KO mice (Fig. 1A). This phenotype appeared at 15 months of age (data not shown). The skeleton is composed of two types of bone, namely cortical bone and trabecular bone. Cortical bone has a higher mineral density, thereby protecting the soft tissues and giving the body its shape. Conversely, trabecular bone, also known as cancellous bone, has a lower mineral density, and it is easily resorbed, thereby playing an important role in calcium homeostasis. Histomorphometric analysis revealed that trabecular bone volume, trabecular number, and trabecular thickness were significantly lower in Zip14-KO mice than in Ctrl mice (Fig. 1B,C). pQCT analysis demonstrated that cortical area, cortical thickness, and cortical density were also decreased in Zip14-KO mice (Fig. 1D). In agreement with the previous reports [20,22], Zip14-KO mice displayed an osteopenia phenotype, and we demonstrated for the first time that osteopenia in Zip14-KO mice appeared in both trabecular and cortical bones.

**Bone resorption activity was increased in Zip14-KO mice**

To examine the cause of the osteopenia phenotype in Zip14-KO mice, histomorphometric analysis was performed using calcein-double-labeled tibia sections at 6 weeks of age. Concerning three parameters reflecting...
osteoblast activity (osteoblast surface, mineral apposition rate, and bone formation rate), there were no significant differences between Ctrl and Zip14-KO mice (Fig. 2A–C). Two parameters reflecting osteoclast activity (osteoclast number and osteoclast surface) were increased in Zip14-KO mice compared to the findings in Ctrl mice, albeit without significance (Fig. 2D,E). Conversely, the eroded surface-to-bone surface ratio (ES/BS) was significantly increased in Zip14-KO mice (Fig. 2F). ES/BS is defined as the percentage of the bone surface that exhibits signs of past or present resorption, thereby representing osteoclast activity in vivo. Further, serum TRAP levels were significantly increased in Zip14-KO mice (Fig. 2G). These results revealed that osteoclast-mediated bone resorption is increased in Zip14-KO mice.
Zip14 mRNA was expressed in osteoclasts both in vitro and in vivo

Because bone resorption was increased in Zip14-KO mice, we measured Zip14 mRNA expression in osteoclasts from Ctrl mice. Osteoclasts are large multinucleated cells with highly TRAP activity that arise from the monocyte/macrophage lineage cells [25]. In situ hybridization revealed that Zip14 mRNA was expressed in TRAP-positive multinucleated cells on the bone surface in vivo (Fig. 3A), but it was not expressed in osteoblasts (data not shown). Next, RT-PCR was performed using cDNA prepared from RANKL-treated Ctrl BMMs in vitro (Fig. 3B). RANKL is a critical cytokine for osteoclast differentiation and activation [26–28]. Two osteoclast maker genes, Oscar and Ctsk, were abundantly expressed 48 and 96 h after RANKL simulation, indicating that BMMs could differentiate into osteoclasts. Zip14 mRNA was modestly expressed in untreated BMMs, and its expression gradually increased after RANKL treatment. These results indicate that ZIP14 is expressed in osteoclasts and suggest the involvement of ZIP14 in RANK-induced osteoclast differentiation.

Osteoclastogenesis was not impaired in Zip14-KO BMMs in vitro

Finally, we examined the effect of ZIP14 disruption on the osteoclastogenesis in RANKL-treated BMMs. BMMs prepared from Ctrl and Zip14-KO mice were cultured in the presence of M-CSF, TGF-β1, and RANKL. TRAP-positive cells with more than three nuclei were identified as osteoclasts, and the area of resorption pits formed by osteoclasts on inorganic bone was measured to evaluate bone-resorbing activity. The images of TRAP-stained cells were comparable between Ctrl and Zip14-KO mice (Fig. 4A). No significant differences were noted for any parameters, including TRAP activity, osteoclast number, and pit area, between Ctrl and Zip14-KO mice (Fig. 4B,C,E). These results indicate that RANKL-induced osteoclastogenesis is not impaired in Zip14-KO BMMs in vitro and intimate that ZIP14 disruption in osteoclasts is not a major cause of the increased bone resorption in Zip14-KO mice.

Discussion

Zip14-KO mice exhibited osteopenia with increased bone resorption activity. Although ZIP14 was expressed in normal osteoclasts, osteoclastogenesis was not impaired in Zip14-KO BMMs in vitro. It has been reported that ZIP14 is expressed ubiquitously and it transports other metals (Mn, Fe, and Cd), in addition to Zn [17–19]. In agreement with these findings, Zip14-KO mice exhibit multiple phenotypes, indicating that ZIP14 is an ion transporter involved in various physiological responses. However, the main cause of the increased bone resorption in Zip14-KO mice was not precisely determined.

There are two possible explanations for the observation of normal osteoclastogenesis in Zip14-KO BMMs in vitro. One possibility is that ZIP14 disruption in osteoclasts increases bone resorption autonomously in

![Fig. 3. Zip14 mRNA was expressed in osteoclasts both in vitro and in vivo.](image-url)
Zip14-KO mice; however, this defect cannot be reproduced under in vitro culture conditions. Occasionally, cultured cells do not have the same physiological responses as the original cells. It has been reported that the percentage of serum, the source of serum, and presence and absence of serum itself influence the Zn-dependent responses of many cell types [29]. Compensatory upregulation of other Zn transporters may be associated with the normal osteoclastogenesis in Zip14-KO BMMs. If this is true, then ZIP14-mediated Zn influx into osteoclasts should negatively regulate osteoclastogenesis; that is, the bone resorption is increased in Zip14-KO mice because of the elimination of this negative regulation.

Many in vitro studies demonstrated that Zn has inhibitory effects on osteoclast-mediated bone resorption [8,9,30], and the present study suggests that ZIP14-mediated Zn influx is involved in this effect. Osteoclasts differentiate from the monocyte/macrophage lineage upon stimulation by two essential cytokines, M-CSF and RANKL [26–28]. Activation of transcription factors such as c-Fos, NF-κB, and nuclear factor-activated T cells, cytoplasmic, calcineurin-dependent 1 (NFATc1) is required for sufficient osteoclast differentiation. In particular, NFATc1 serves as a master transcriptional regulator of osteoclast differentiation. Oral Zn administration was reported to decrease osteoclastogenesis by inhibiting RANKL expression in Zn-adequate rats [31]. Park et al. reported that Zn inhibited osteoclast differentiation in vitro by inhibiting Ca^{2+}–calcineurin–NFATc1 signaling pathway [32]. Phospho-NFATc1 is dephosphorylated by activated calcineurin, which leads to nuclear translocation of the protein and the induction of NFATc1-mediated gene transcription. Zn inhibits calmodulin activity by competing with Ca^{2+} for binding to calmodulin, resulting in the inhibition of NFATc1 translocation to the nucleus [33]. Yamaguchi et al. reported that Zn inhibited osteoclast differentiation in vitro by inhibiting NF-κB activation [30]. Adenoviral overexpression of ZIP1 also inhibited NF-κB activation leading to impaired osteoclast function [15]. ZIP14-mediated Zn influx may influence Ca^{2+}–calcineurin–NFATc1 signaling and/or NF-κB activation during osteoclast differentiation.

Lipopolysaccharide (LPS), a component of the outer membranes of Gram-negative bacteria, is capable of inducing bone resorption in both in vitro and in vivo studies [34,35]. LPS induces the production of pro-inflammatory cytokines such as tumor necrosis factor-

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**Fig. 4.** Osteoclastogenesis was not impaired in Zip14-KO BMMs in vitro. (A–C) BMMs were cultured with M-CSF (50 ng·mL⁻¹), TGF-β1 (1 ng·mL⁻¹), and RANKL (20 or 100 ng·mL⁻¹). Osteoclast formation was evaluated by TRAP staining image (A), TRAP activity (B), and osteoclast number (C, count only when cultured with 100 ng·mL⁻¹ RANKL). TRAP-positive cells with more than three nuclei were identified as osteoclasts. (D,E) BMMs were cultured with M-CSF (50 ng·mL⁻¹), TGF-β1 (1 ng·mL⁻¹), and RANKL (100 ng·mL⁻¹) on Corning OsteoAssay Surface 96-well plates. After cultivation, cells were removed using 5% sodium hypochlorite. Resorption pits were visualized under a scanning electron microscope (D), and the resorption area was quantified using IMAGEJ software (E). (B,C,E) Data are presented as mean ± SE (n = 5). NS: no statistical difference by Student’s t-test.
α, interleukin (IL)-1β, and IL-6, which can directly stimulate osteoclast differentiation. LPS alone can induce osteoclast differentiation in RAW264.7 macrophage cells but not in BMM culture [36,37]. LPS was reported to strongly induce ZIP14 mRNA in primary macrophage cells prepared from human blood [38]. Therefore, upregulation of ZIP14 expression may be involved in LPS-induced osteoclastogenesis.

The second possible explanation for the normal osteoclastogenesis in Zip14-KO BMMs is that ZIP14 regulates osteoclast-mediated bone resorption in a non-cell-autonomous manner. Among the defects observed in Zip14-KO mice, elevated IL-6 expression with chronic inflammation is the most likely cause of increased bone resorption. It has been reported that osteoclast formation is triggered by IL-6 in the presence of soluble IL-6 receptor (sIL-6R), and RANKL expression is induced by IL-6/sIL-6R via the JAK/STAT signaling pathway [39,40]. IL6-KO mice are protected against ovariectomy-induced osteoporosis via a mechanism that prevents osteoclast activation [41]. Accordingly, anti-IL-6R antibody inhibits osteoclast formation in animal models and patients with rheumatoid arthritis [42,43]. Anti-IL-6R antibody treatment of Zip14-KO mice would resolve the involvement of elevated IL-6 levels in the increased bone resorption activity.

Many in vitro studies demonstrated that ZIP14 transports other metals (Mn, Fe, and Cd). Among them, Fe has been reported to be involved in osteoclastogenesis [44]. Transferrin receptor 1-mediated Fe uptake promotes osteoclast differentiation and bone resoring, which are associated with the induction of mitochondrial respiration and the production of reactive oxygen species. Assuming that ZIP14 transports Fe into osteoclasts, Fe levels are predicted to be decreased in Zip14-KO osteoclasts. Therefore, Fe uptake-mediated promotion of osteoclast differentiation is not likely to be related to the increased bone resorption observed in Zip14-KO mice. Recently, loss-of-function mutations in human ZIP14 were identified in patients with hypermanganesemia and progressive parkinsonism–dystonia [19]. In these patients, blood Mn levels are drastically increased, without affecting Zn, Fe, and Cd levels. Consequently, the authors claimed that the major role of ZIP14 is to transport Mn, and ZIP14 dysfunction reduces biliary Mn elimination, causing hypermanganesemia. Strause et al. reported that Mn deficiency decreased osteoclast activity in rats fed Mn-depleted diets [45]. It is necessary to examine Mn metabolism in Zip14-KO mice and the relationship between hypermanganesemia and osteoclastogenesis.

In conclusion, the present study illustrates that ZIP14 is involved in the negative regulation of osteoclastogenesis and supports the fact that Zn exerts an inhibitory effect on osteoclast-mediated bone resorption. Both Zn and other metals transported by ZIP14 may be involved in the ZIP14-mediated regulation of osteoclastogenesis. Further identification of Zn transporters involved in osteoclastogenesis will help to clarify the role of Zn in bone homeostasis and the pathological mechanism of skeletal abnormalities produced by Zn deficiency.

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Author contributions
SS, MT, and MS performed the main experiments and analyzed the data. SH and TF generated Zip14-KO mice. MT prepared cDNA from RANKL-treated BMMs and advised about in vitro assay for osteoclastogenesis. TF planned the experiments, analyzed data, and wrote the manuscript.

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