Effects of *Melandrium firmum* Rohrbach on RANKL-induced osteoclast differentiation and OVX rats

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Abstract. Osteoporosis is a systemic skeletal disease characterized by reduced bone mineral density (BMD), which results in an increased risk of fracture. Melandrium firmum (Siebold & Zucc.) Rohrbach (MFR), 'Wangbulryuhaeng' in Korean, is the dried aerial portion of Melandrii Herba Rohrbach, which is a member of the Caryophyllaceae family and has been used to treat several gynecological conditions as a traditional medicine. However, to the best of our knowledge, the effect of MFR on osteoclast differentiation and osteoporosis has not been assessed. To evaluate the effects of MFR on osteoclast differentiation, tartrate-resistant acid phosphatase staining, actin ring formation and bone resorption assays were used. Additionally, receptor activator of nuclear factor-kB ligand-induced expression of nuclear factor of activated T cell, cytoplasmic 1 (NFATc1) and c-Fos were measured using western blotting and reverse transcription-PCR. The expression levels of osteoclast-related genes were also examined. To further investigate the anti-osteoporotic effects of MFR in vivo, an ovariectomized (OVX) rat model of menopausal osteoporosis was established. Subsequently, the femoral head was scanned using micro-computed tomography. The results revealed that MFR suppressed osteoclast differentiation, formation and function. Specifically, MFR reduced the expression levels of osteoclast-related genes by downregulating transcription factors, such as NFATc1 and c-Fos. Consistent with the in vitro results, administration of MFR water extract to OVX rats reduced BMD loss, and reduced the expression levels of NFATc1 and cathepsin K in the femoral head. In conclusion, MFR may contribute to alleviate osteoporosis-like symptoms. These results suggested that MFR may exhibit

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potential for the prevention and treatment of postmenopausal osteoporosis.

Introduction

Osteoporosis is a systemic skeletal disease characterized by reduced bone density, which leads to an increased risk of fracture (1). Osteoporosis is considered a public health problem, the prevalence of which is increasing worldwide (2). Bone homeostasis requires maintenance of a balance between osteoclast-induced bone resorption and osteoblast-induced bone formation; however, an increase in osteoclast number and activity can result in an imbalance in bone remodeling (3). This imbalance can result in osteoclast-mediated bone diseases, such as rheumatoid arthritis, periodontal disease, Paget's disease and osteoporosis (4).

Bisphosphonate and hormone replacement therapy with estrogen are the most widely used treatments for postmenopausal osteoporosis (5). However, long-term treatment with bisphosphonate is unsuitable for several patients with osteoporosis owing to serious side effects, such as osteonecrosis of the jaw, atrial fibrillation, esophageal cancer, musculoskeletal pain and atypical fractures (6-9). In addition, hormone replacement therapy has unwanted side effects, such as an increased risk of developing ovarian cancer and breast cancer (10-12). Thus, it is essential to discover novel effective and safe treatments for osteoporosis.

Osteoclasts are multinucleated giant cells derived from monocyte/macrophage hematopoietic precursor cells (13). Receptor activator of nuclear factor (NF)-kB (RANK) ligand (RANKL) is an essential cytokine of osteoclast differentiation (14). Binding of RANKL to RANK induces recruitment of the adaptor protein tumor necrosis factor (TNF) receptor-associated factor (TRAF6). TRAF6 induces the translocation of NF-KB (15,16). These signaling pathways are responsible for the activation of key transcription factors, such as c-Fos and nuclear factor of activated T cell, cytoplasmic 1 (NFATc1), resulting in increased expression of osteoclast-specific genes, including tartrate-resistant acid phosphatase (TRAP/Acp5), ATPase H+ transporting V0 subunit D2 (ATP6v0d2/Atp6v0d2), osteoclast-associated immunoglobulin-like receptor (OSCAR/Oscar), cathepsin K (CTK/Ctsk) and matrix metalloproteinase-9 (MMP-9/Mmp9) (17-20).

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Melandrium firmum (Siebold & Zucc.) Rohrbach (MFR) is the dried aerial portion of Melandrii Herba Rohrbach, a member of the Caryophyllaceae family. MFR is known as 'Wangbulryuhaeng' in Korea and has been traditionally used to treat gynecological conditions, such as breast cancer and lactation disorders (21). Previous studies have shown that the methanol extract of MFR exhibits an anti-inflammatory effect on lipopolysaccharide (LPS)-induced proinflammatory cytokines, such as TNF- α and interleukin (IL)-1 β (22,23). According to previous studies, inhibition of TNF- α and IL-1 β has been associated with metabolic bone disease, such as postmenopausal osteoporosis and rheumatoid arthritis. IL-1ß has been revealed to be regulated by RANKL to increase activity and promote osteoclast formation (24,25). In addition, TNF- α may serve an important role in regulating bone homeostasis by stimulating osteoclast formation and inhibiting osteoblast function (26,27). Furthermore, vitexin, another active compound found in MFR, has been shown to inhibit osteoclastogenesis and prevent LPS-induced osteolysis (28). Based on these previous findings, it was hypothesized that MFR may potentially attenuate osteoclast differentiation, thus preventing bone loss in osteoporosis. However, to the best of our knowledge, the effect of MFR on osteoclast differentiation and osteoporosis is yet to be explained.

In the present study, the effect of MFR on osteoclast differentiation in RANKL-induced mouse macrophage RAW 264.7 cells and bone loss in an ovariectomized (OVX) rat model of osteoporosis was assessed.

Materials and methods

Reagents. DMEM was purchased from Welgene, Inc. Minimum essential medium-α (α-MEM), penicillin/streptomycin (P/S) and Dulbecco's PBS (DPBS) were obtained from Gibco; Thermo Fisher Scientific, Inc. FBS was purchased from Atlas Biologicals. RANKL was purchased from PeproTech, Inc. CellTiter 96 Aqueous non-radioactive cell proliferation (MTS) assay was purchased from Promega Corporation. Bicinchoninic acid (BCA) solution, phosphatase inhibitor cocktail, DAPI, 17β -estradiol (E₂) and alendronate (ALN) were obtained from Sigma-Aldrich; Merck KGaA. Osteo Assay Surface multiple well plates (cat. no. 3989) were obtained from Corning, Inc.. PCR primers were synthesized by Genotech Corp. Acti-stain[™] 488 Fluorescent Phalloidin was purchased from Cytoskeleton, Inc.. The primary antibodies and secondary antibodies used in the present study were: β-actin (cat. no. sc-8432; Santa Cruz Biotechnology, Inc.), c-Fos (cat. no. sc-447; Santa Cruz Biotechnology, Inc.), NFATc1 (cat. no. 556602; BD Biosciences), MMP-9 (cat. no. ab38898; Abcam), CTK (cat. no. ab19027; Abcam), TRAF6 (cat. no. sc-8409; Santa Cruz Biotechnology, Inc.) and peroxidase AffiniPure Goat Anti-Mouse IgG (cat no. 115-035-062; Jackson ImmunoResearch Laboratories, Inc.) and peroxidase AffiniPure Goat Anti-Rabbit IgG (cat no. 115-035-144; Jackson ImmunoResearch Laboratories, Inc). All other reagents used were of analytical grade.

Preparation of MFR extract. MFR was purchased from Omniherb. MFR was prepared by decocting 300 g of the dried herb in 1.5 l boiling distilled water (D.W) for 2 h. The

extract was filtered using filter paper (no. 3; Whatman plc; GE Healthcare Life Sciences) and collected in a rotary evaporator at 55°C. The extract was lyophilized and 19 g dried powder was obtained (yield ratio, 12.7%).

High-performance liquid chromatography (HPLC) analysis. Vitexin (cat. no. 49513; Sigma-Aldrich; Merck KGaA) is the active ingredient in MFR (29); therefore, to quantitatively evaluate the MFR extract, HPLC was performed with vitexin used as an internal standard. Vitexin was prepared in DMSO. Analysis was performed using a UV detector (2996 Waters 2695; Waters Corporation); separation was carried out on an Xbridge-C18 with a C18 guard column (250 x 4.6 mm; 5 μ m; Waters Corporation) and proceeded at 30°C for 50 min at a flow rate of 1 ml/min. Vitexin was detected at 335 nm. Samples were injected in a volume of 10 μ l. The mobile phase consisted of acetonitrile (A) and water (B), at a composition of 10% A from 0-10 min and 50% A from 10-30 min.

RAW 264.7 cell culture and cytotoxicity assay. RAW 264.7 cells are mouse monocyte-macrophage like cells (30). RAW 264.7 cells were obtained from Korean Cell Line Bank; Korean Cell Line Research Foundation. RAW 264.7 cells were cultured in DMEM supplemented with 10% FBS and 1% P/S in a cell incubator at 37°C and 5% CO₂. To examine cell viability, the cells were seeded in a 96-well plate at 5x10³ cells/well and incubated for 24 h. RAW 264.7 cells were treated with MFR (12.5, 25, 50 and 100 μ g/ml) at 37°C for 24 h. After treatment, 20 μ l/well MTS solution was added and the cells were incubated at 37°C for 2 h. To determine cell viability of mature osteoclasts, RAW 264.7 cells were seeded in a 96-well plate at 5x10³ cells/well and were treated with or without RANKL (100 ng/ml) for 5 days. Subsequently, cells were treated with MFR (12.5, 25, 50 and 100 μ g/ml) for 1 day, after which, 20 µl MTS solution was added and incubated at 37°C for 2 h. Cell viability was measured using an enzyme-linked immunosorbent assay (ELISA) reader at a wavelength of 490 nm.

TRAP staining and measurement of TRAP levels. A total of 5x10³ RAW 264.7 cells/well were seeded in a 96-well plate. After 24 h, to induce osteoclast differentiation of RAW 264.7 cells, cells were treated with RANKL (100 ng/ml) and MFR (12.5, 25, 50 and 100 μ g/ml) at 37°C for 5 days. The medium was replaced every 2 days. In order to compare the inhibitory effect of MFR and vitexin on osteoclast differentiation, cells were seeded in the same manner as mentioned previously. After 24 h, the medium was replaced with RANKL (100 ng/ml) and MFR (50 and 100 μ g/ml) or vitexin (0.0753 and 0.147 μ g/ml) at 37°C for 5 days. Subsequently, the cells were washed and fixed with 4% formalin at room temperature for 10 min. TRAP staining was performed using a TRAP kit (Sigma-Aldrich; Merck KGaA) according to the manufacturer's protocol. The number of TRAP-positive cells was counted using an inverted light microscope (Olympus Corporation; magnification, x100).

TRAP is secreted in large quantities by osteoclasts and TRAP levels are considered a biochemical marker of osteoclast function (31). Therefore, the present study also analyzed TRAP levels to determine the effect of MFR on osteoclast function. In order to measure the TRAP levels, 50 μ l differentiation medium was obtained and transferred to a new plate. Subsequently, an equal volume of TRAP solution (4.93 mg 4-Nitrophenyl phosphate disodium salt hexahydrate in 850 μ l 0.5 M acetate solution and 150 μ l tartrate solution) was added to the differentiation medium and incubated at 37°C for 1 h. Subsequently, the reaction was terminated using 0.5 M NaOH and TRAP levels were measured using an ELISA reader at a wavelength of 405 nm.

Filamentous actin (F-actin) ring formation and pit formation. To determine F-actin ring formation, $5x10^3$ cells/well were seeded in a 96-well plate, and treated with RANKL (100 ng/ml) and MFR (12.5, 25, 50 and 100 µg/ml) at 37°C for 5 days. The medium was replaced every 2 days. The cells were fixed with 4% paraformaldehyde at room temperature for 20 min and permeabilized with 0.1% Triton X-100 in PBS at room temperature for 5 min. The cells were then stained using Acti-stainTM 488 Fluorescent Phalloidin at room temperature in the dark for 30 min. Cells were washed with PBS and nuclei were counterstained with DAPI. Images were captured using fluorescence microscopy (Cellena; Logos Biosystems; magnification, x200).

To examine pit formation, the cells were seeded in Osteo Assay Surface multiple well plates at 5×10^3 cells/well. Subsequently, the cells were treated with RANKL (100 ng/ml) and MFR (12.5, 25, 50 and 100 μ g/ml) at 37°C for 5 days. The differentiation medium was replaced every 2 days. Subsequently, cells were washed and removed using deionized water with 4% sodium hypochlorite. Images of resorbed areas were captured using an inverted light microscope (Olympus Corporation; magnification, x200). Total pit areas were analyzed using ImageJ version 1.46 (National Institutes of Health).

Western blotting. To examine the effect of MFR on osteoclast differentiation-associated transcription factors, 5x10⁵ RAW 264.7 cells/well were seeded in a 60-mm dish and treated with RANKL (100 ng/ml) and MFR (12.5, 25, 50 and 100 µg/ml) at 37°C for 1 day. In order to compare the inhibitory effect of MFR and vitexin on the expression of NFATc1 and c-Fos, 5x10⁵ RAW 264.7 cells/well were seeded in a 60-mm dish and treated with RANKL (100 ng/ml) and MFR (50 and 100 μ g/ml) or vitexin (0.0753 and 0.147 μ g/ml) at 37°C for 1 day. To prepare whole-cell lysates, the cells were washed with cold DPBS, and the total proteins were extracted using RIPA lysis buffer (50 mM Tris-CI, 150 mM NaCl, 1% NP-40, 0.5% Na-deoxycholate, 0.1% SDS, protease inhibitor cocktail, phosphatase inhibitor cocktail). Protein concentration was determined using a BCA assay. Proteins (30 μ g) were separated by SDS-PAGE on 10% gels and transferred to a nitrocellulose membrane (Whatman plc; GE Healthcare Life Sciences) according to the manufacturer's protocol. The membranes were blocked in 5% skimmed milk for 1 h at 37°C, followed by overnight incubation at 4°C with primary antibodies against NFATc1 (1:1,000), c-Fos (1:1,000), TRAF6 (1:1,000), MMP-9 (1:1,000), CTK (1:1,000) and β-actin (1:1,000). Subsequently, membranes were incubated with secondary antibodies (1:10,000) for 1 h at room temperature. The membranes were visualized using enhanced chemiluminescence reagent (cat no. RPN2106; GE Healthcare Life Sciences) and protein expression levels were semi-quantified using ImageJ (version 1.46; National Institutes of Health).

Semi-quantitative reverse transcription-PCR (RT-PCR). To examine the effect of MFR on osteoclast-related markers, $2x10^5$ cells/well seeded in a 6-well plate, and treated with the RANKL (100 ng/ml) and MFR (12.5, 25, 50 and 100 µg/ml) at 37°C for 4 days. Total RNA of the treated cells was extracted using RNAiso Plus (cat. no. 9109; Takara Bio, Inc.) according to the manufacturer's protocol. A total of $2 \mu g$ RNA was measured using a NanoDrop 2.0 spectrophotometer (NanoDrop; Thermo Fisher Scientific, Inc.). cDNA was synthesized using a RT kit (Invitrogen; Thermo Fisher Scientific, Inc.), according to the manufacturer's protocol. The synthesized cDNA was amplified by PCR using Taq polymerase (Kapa Biosystems; Roche Diagnostics). The PCR analysis conditions were as follows: 26-40 cycles of 30 sec at 94°C (denaturation), 30 sec at 53-58°C (annealing) and 30 sec at 72°C (extension). The primer sequences are listed in Table I. β-actin was used as a loading control. The products of qPCR were assessed on a 2% agarose gel stained with SYBR-Green (Invitrogen; Thermo Fisher Scientific, Inc.). The expression levels of mRNA were semi-quantified using ImageJ version 1.46.

Animal experimental design. To further investigate the anti-osteoporotic effects of MFR in vivo, an OVX rat model of menopausal osteoporosis was used. A number of 48 female Sprague Dawley (SD)-rats (age, 12 weeks; weight, 230-250 g) were obtained from KOATECH. The rats were housed in a standard environment with a controlled temperature of 22±2°C and humidity of 55±5%, under a 12-h light/dark cycle. Animals were provided with ad libitum access to water and food. The protocol for in vivo experiments was approved by the Kyung Hee Medical Center Institutional Animal Care and Use Committee (KHMC-IACUC; approval no. KHMC-IACUC 19-017). All animals were allowed to acclimate for 1 week. For the postmenopausal osteoporosis model, rats were deeply anesthetized using 5% isoflurane (inhaled in 100% oxygen) and the bilateral ovaries of the rats were removed under 2-2.5% isoflurane anesthesia. The sham group underwent the same surgery to ensure they experienced the same stress, but the ovaries were not removed. No animals died during surgery. To prevent infection of the surgical site, the wound was sutured and gentamicin (4 mg/kg) was injected intraperitoneally for 3 days. After 1 week, the rats were randomly divided into six groups (n=8/group): i) Sham group, in which rats underwent the sham operation and were treated with vehicle (water); ii) OVX group, in which OVX rats were treated with vehicle; iii) E₂ group, in which OVX rats were treated with E_2 (100 μ g/kg); iv) ALN group, in which OVX rats were treated with ALN (5 mg/kg); v) MFR-L group, in which OVX rats were treated with a low dose of MFR (16.9 mg/kg); and vi) MFR-H group, in which OVX rats were treated with a high dose of MFR (169 mg/kg) for 8 weeks. E2 and ALN were used as the positive controls. The humane endpoints used in the present study were: Dirty hair and eye discharge; self-injury and anxiety; vomiting and hemoptysis; inactivity; or anxiety and headache. None of the animals exhibited abnormal behavior.

MFR dose was calculated as follows: In Korean medicine, based on an average adult weight of 60 kg, a single dose of 8 g

Gene name	Sequence (5'-3')	Temperature (°C)	Cycle	Accession no.
Nfatc1	F: TGCTCCTCCTCCTGCTGCTC	58	32	NM_198429.2m
	R: CGTCTTCCACCTCCACGTCG			
Fos	F: ATGGGCTCTCCTGTCAACAC	58	40	NM_010234.3
	R: GGCTGCCAAAATAAACTCCA			
Ctsk	F: AGGCGGCTATATGACCACTG	58	26	NM_007802.4
	R: CCGAGCCAAGAGAGCATATC			
Mmp9	F: CGACTTTTGTGGTCTTCCCC	58	30	NM_013599.4
	R: TGAAGGTTTGGAATCGACCC			
Ca2	F: CTCTCAGGACAATGCAGTGCTGA	58	32	NM_001357334.1
	R: ATCCAGGTCACACATTCCAGCA			
Acp5	F: ACTTCCCCAGCCCTTACTACCG	58	30	NM_007388.3
	R: TCAGCACATAGCCCACACCG			
Atp6v0d2	F: ATGGGGCCTTGCAAAAGAAATCTG	58	30	NM_175406.3
	R: CGACAGCGTCAAACAAAGGCTTGTA			
Dcstamp	F: TGGAAGTTCACTTGAAACTACGTG	63	30	NM_001289506.1
	R: CTCGGTTTCCCGTCAGCCTCTCTC			
Oscar	F: CTGCTGGTAACGGATCAGCTCCCCAGA	53	35	NM_001290377.1
	R: CCAAGGAGCCAGAACCTTCGAAACT			
Src	F: TCCAGGCTGAGGAGTGGTACTTTGG	64	40	NM_001025395.2
	R: ATACGGTAGTGAGGCGGTGACACAG			
Prdm1	F: TTCTTGTGTGGTATTG	50	40	NM_007548.4
	R: TTGGGGACACTCTTTG			
Actb	F: TTCTACAATGAGCTGCGTGT	58	30	NM_007393
	R: CTCATAGCTCTTCTCCAGGG			

Table I. Primer	sequences	for reverse	transcription	-PCR analys	sis

Ctsk, cathepsin k; *Mmp9*, matrix metalloproteinase-9; *Ca2*, carbonic anhydrase 2; *Acp5*, tartrate-resistant acid phosphatase; *Atp6v0d2*, ATPase H+ transporting V0 subunit D2; *Dcstamp*, dendritic cell-specific transmembrane protein; *Oscar*, osteoclast-associated receptor; *Pdrm1*, b lymphocyte-induced maturation protein-1; F, forward; R, reverse.

is recommended. MFR (1.016 g; yield, 12.7%) was considered equivalent to 8 g; thus, 16.9 mg MFR was required per 1 kg. Therefore, the MFR-L group was treated with 16.9 mg/kg. The MFR-H group was treated as follows: Rats are well-known to exhibit 6.4-fold faster metabolism than humans (32). In vitro experiments revealed that MFR exhibited a higher inhibition of osteoclast differentiation at high concentrations than at low concentrations. Based on these findings, rats in the MFR-H groups were treated with MFR at concentrations 10-fold higher than those in the MFR-L group to induce higher pharmacological effects. Therefore, the MFR-H group was treated with 169 mg/kg MFR (33-35). Oral administration was performed every morning for 8 weeks. Body weight was measured once a week and the dose was adjusted to weight. After 8 weeks, all animals were anesthetized by 5% isoflurane of inhaled anesthetics in 100% oxygen. Blood was collected using a cardiac puncture following sacrifice by cervical vertebrae dislocation. Subsequently, the uterus and femur were collected and weighed. Femurs samples were collected and fixed in 10% neutral buffered formalin for 1 day at room temperature. Uterus samples were collected and then stored at -80°C.

Serum biochemical analysis. Blood samples were incubated at room temperature for 30 min, and centrifuged at 29,739 x g

for 10 min at 4°C. Serum samples were stored at -80°C until required. Serum levels of alkaline phosphatase (ALP), aspartate aminotransferase (AST) and alanine aminotransferase (ALT) were measured by DKKorea. C-telopeptide of collagen type 1 (CTX-1) levels were measured using an ELISA kit (Elabscience; cat. no. E-EL-R1456) according to the manufacturer's protocol. TRAP levels were measured using a TRAP kit as aforementioned.

Micro-computed tomography (micro-CT) analysis. The femoral head was scanned using micro-CT (SkyScan1176; Bruker Corporation). Bone microarchitecture parameters, including bone mineral density (BMD), trabecular thickness (Tb.Th) and trabecular separation (Tb.sp) were analyzed using NRecon software (SkyScan version 1.6.10.1; Bruker Corporation).

Histological analysis. The fixed femur samples were washed using D.W at room temperature for 1 day and decalcified using 10% EDTA at room temperature for 3 weeks. Subsequently, the femur samples were dehydrated at room temperature for 1 day and embedded in paraffin. Paraffin-embedded tissues (5 μ m) were sectioned on a rotary microtome (Carl Zeiss AG) and tissue sections were mounted on slides at room

temperature for 1 day. The sections on the slides were stained with hematoxylin-eosin (H&E); sections were stained with hematoxylin for 10 min and with eosin for 10 sec at room temperature. Subsequently, all slides were sealed using mounting solution and the sections were viewed under a light microscope (Olympus Corporation; magnifications, x40 and x100) for histological evaluation.

Immunohistochemistry (IHC). Paraffin-embedded tissues were deparaffinized in xylene. Endogenous peroxidases were blocked in 0.3% hydrogen peroxide at room temperature for 15 min and proteinase K (0.4 mg/ml) was used for antigen-retrieval at 37°C for 30 min. The sectioned tissues were incubated with 10% normal serum (Gibco; Thermo Fisher Scientific, Inc.) at room temperature for 1 h to block nonspecific binding, then slides were washed with PBS and incubated at 4°C for overnight with primary antibodies against NFATc1 (1:100) and CTK (1:100). The following day, the slides were washed with PBS and incubated with a secondary antibody (1:100; rabbit; cat. no: BA-1000; Vector Laboratories, Inc.) at 4°C for 1 h. The signal was visualized using an ABC kit (Vector Laboratories, Inc.) and 3, 3'-diaminobenzidine solution (Vector Laboratories, Inc.). The stained tissues were imaged using a light microscope (magnifications, x100 and x200).

MC3T3-E1 cell culture and cytotoxic assay. MC3T3-E1 cells were purchased from American Type Culture Collection. MC3T3-E1 cells were cultured in α -MEM without ascorbic acid containing 10% FBS and 1% P/S in a cell incubator at 37°C and 5% CO₂. To examine cell viability, the cells were seeded in a 24-well plate at 1x10⁴ cells/well and incubated at 37°C for 24 h. Subsequently, the cells were treated with α -MEM without ascorbic acid and MFR (12.5, 25, 50 and 100 µg/ml) at 37°C for 24 h. Subsequently, 20 µl/well MTS solution was added to the cells and incubated at 37°C for 2 h. Cell viability was evaluated using an ELISA reader at a wavelength of 490 nm and was expressed as a percentage of the control.

Alizarin red S staining. MC3T3-E1 cells were seeded in a 24-well plate at $1x10^4$ cells/well and incubated for 24 h. Subsequently, osteoblast differentiation was induced using osteogenic medium (α -MEM without ascorbic acid supplemented with 10 mM β -glycerophosphate, 25 μ g/ml ascorbic acid) and MFR (12.5, 25, 50 and 100 μ g/ml) at 37°C for 14 days. The medium was replaced every 2 days. The cells were then washed with DPBS, fixed with 80% ethanol at room temperature for 1 h and stained with Alizarin red S at room temperature for 5 min. To quantify mineralization, the stained dye was extracted using 10% (v/w) cetylpyridinium chloride in sodium phosphate at room temperature for 15 min. The extracted dye was measured using an ELISA reader at a wavelength of 570 nm.

Statistical analysis. Data are presented as the mean ± standard error of the mean of three experiments. Statistical analyses were performed using GraphPad Prism version 5.01 (GraphPad Software, Inc.). *In vitro* and *in vivo* experiments were compared using a one-way ANOVA followed by a Tukey's post hoc analysis. Animal body weight was compared using a two-way ANOVA followed by Bonferroni post hoc test. P<0.05 was considered to indicate a statistically significant difference.

Results

Quantitative analysis of the MFR extract. Vitexin was used as a standard marker of MFR, as described previously (29). The chromatogram of the MFR water extract possessed several peaks at a retention time of 0-30 min, and vitexin was observed at the same retention time as the standard (Fig. 1).

Effects of MFR on TRAP staining, and F-actin ring and pit formation. To analyze the anti-osteoporotic effect of the MFR extract in vitro, TRAP staining was performed using a TRAP kit. Following treatment with RANKL, TRAP staining revealed that the number of TRAP-positive cells was reduced by MFR treatment. Consistent with the results of TRAP staining, MFR decreased TRAP levels in the differentiation medium (Fig. 2A-C). As the actin ring is essential in osteoclast differentiation (36), the effect of MFR on actin ring formation was determined using immunocytochemistry. In addition, the effect of MFR on pit formation was determined using osteo-coated plates. The actin ring structures and the area of bone resorption pits were increased in the RANKL-treated cells, but were reduced following MFR treatment (Fig. 2D and E). Consistent with these results, MFR treatment reduced the number of actin rings in a dose-dependent manner (Fig. 2F). In addition, the area of bone resorption pits was significantly reduced following MFR treatment (Fig. 2G). To confirm whether the concentration of MFR used in the in vitro experiments affected the viability of RAW 264.7 cells, the cells were treated with 12.5-100 μ g/ml MFR. MFR did not affect the viability of RAW 264.7 cells or mature osteoclasts (Fig. 2H and I). In addition, the present study compared and analyzed the inhibitory effect of MFR and vitexin on osteoclast differentiation to determine if the ability of MFR to inhibit osteoclast differentiation was associated with vitexin. The content of vitexin in MFR was 1.47 ppm (0.147%); 0.147 μ g/ml vitexin in 100 μ g/ml MFR and 0.0753 μ g/ml vitexin in 50 μ g/ml MFR. The effects of the two substances (MFR and vitexin) on TRAP staining and TRAP levels in the medium were subsequently assessed. To confirm whether the concentration of vitexin contained in MFR affects the osteoclast inhibitory effect, TRAP staining was performed. As shown in Fig. S1A and B, TRAP-positive cells and TRAP levels were increased in the RANKL-treated cells. Conversely, the number of TRAP-positive cells was considerably decreased by MFR (50 and 100 μ g/ml) treatment compared with vitexin (0.0753 and 0.147 μ g/ml). Similarly, MFR (50 and 100 μ g/ml) further decreased TRAP levels compared with vitexin (0.0753 and 0.147 µg/ml) (Fig. S1B).

To determine whether the concentration of vitexin contained in MFR affects the inhibitory effect on transcription factors, NFATc1 and c-Fos were measured using western blotting. The protein expression levels of NFATc1 and c-Fos were increased by RANKL treatment. MFR (50 and 100 μ g/ml) inhibited the expression of NFATc1 and c-Fos compared with vitexin (0.0753 and 0.147 μ g/ml) (Fig. S1C and D). To confirm whether the concentration of vitexin used in experiments affected the viability of RAW 264.7 cells, the cells were treated with vitexin (0.0753 and 0.147 μ g/ml); vitexin



Figure 1. High-performance liquid chromatography analysis of the MFR extracts. Vitexin was confirmed in (A) the standard and (B) MFR water extract at a wavelength of 335 nm. MFR, *Melandrium firmum* Rohrbach.

not affect cell cytotoxicity (Fig. S1E). Taken together, it was revealed that the osteoclast inhibitory effect of MFR was not mediated by vitexin.

Effect of MFR on protein and mRNA expression levels of NFATc1 and c-Fos. NFATc1 and c-Fos are important transcription factors required for mature osteoclast differentiation in RAW 264.7 cells (37,38). RANKL-induced expression of NFATc1 and c-Fos was measured using western blotting and RT-qPCR. Treatment with RANKL significantly increased the expression levels of NFATc1, whereas MFR decreased the protein and mRNA expression levels of NFATc1 (Fig. 3A and B). In addition, c-Fos expression was significantly increased by RANKL treatment, whereas this increase was reversed by MFR treatment (Fig. 3C and D). Furthermore, RANKL stimulation increased TRAF6, but this finding was not significant, and treatment of MFR attenuated the increased expression of TRAF6. In particular, in cells treated with 25 μ g/ml MFR, the expression of TRAF6 was significantly decreased compared with that in RANKL-only treated cells (Fig. 3E and F).

Effect of MFR on RANKL-induced expression of bone resorption and osteoclast-related markers. The inhibitory effects of the MFR extract on bone resorption markers were examined using western blotting and RT-qPCR. Treatment with RANKL significantly increased the expression levels of MMP-9 and CTK, whereas MFR decreased the protein expression levels of MMP-9 and CTK (Fig. 4A and B). Consistent with the results of western blotting, MFR reduced the mRNA expression levels of Mmp9, Ctsk and carbonic anhydrase 2 (CA2/Ca2) (Fig. 4C and D). In addition, the effects of MFR on RANKL-stimulated changes in osteoclast-specific genes, including Acp5, Atp6v0d2, dendritic cell-specific transmembrane protein (DC-STAMP/Dcstamp), Oscar, c-Src (Src) and B lymphocyte-induced maturation protein-1 (Blimp-1/Prdm1), were determined, as they are crucial in promoting osteoclastogenesis (39). The mRNA expression levels of osteoclast-specific genes were increased by RANKL treatment, whereas MFR attenuated the RANKL-induced increase in the mRNA expression levels of osteoclast-specific genes, including Acp5, Atp6v0d2, Dcstamp, Oscar, Src and Prdml, in a dose-dependent manner (Fig. 4E-H).

Effect of MFR on body, uterus and femur weights, and on the levels of hepatotoxicity, bone formation and bone resorption markers. As shown in Fig. 5A, the OVX group exhibited a significant increase in body weight compared with that in the sham group from 4 weeks. The E_2 group had a significantly lower increase in body weight after 3 weeks compared with



Figure 2. Effect of MFR on RANKL-induced osteoclast differentiation, F-actin ring formation and pit formation. (A) RANKL-induced osteoclasts were stained using a TRAP kit. Magnification, x100; scale bar, 200 μ m. (B) Number of TRAP-positive cells with >3 nuclei were counted. (C) TRAP levels was measured using an enzyme-linked immunosorbent assay reader. (D) F-actin rings were stained using fluorescent phalloidin and (E) a bone resorption assay was performed using osteo-coated plates. Magnification, x100; scale bar, 200 μ m. (F) Number of F-actin rings was counted and (G) pit area was measured using ImageJ. (H) RAW 264.7 cell viability was determined using the MTS assay. (I) Cell viability of mature osteoclasts was examined using the MTS assay. All data are presented as the mean ± standard error of the mean of three independent experiments. Data were analyzed using one-way ANOVA followed by Tukey's post hoc test. ^aP<0.05 vs. normal group (untreated cells); ^bP<0.05 vs. RANKL treatment group; ^cP<0.05 vs. MFR 12.5 μ g/ml treatment group; ^dP<0.05 vs. MFR 25 μ g/ml treatment group; ^cP<0.05 vs. MFR 50 μ g/ml treatment group. MFR, *Melandrium firmum* Rohrbach; RANKL, receptor activator of nuclear factor- κ B ligand; F-actin, filamentous actin; TRAP, tartrate-resistant acid phosphatase.

the OVX group; however, MFR-L, MFR-H and ALN treatment did not affect body weight. In addition, the OVX group exhibited significantly decreased uterus weight compared with that in the sham group, whereas E_2 treatment prevented uterus weight loss. MFR-L, MFR-H and ALN groups did not exhibit any protective effects on uterus weight loss (Fig. 5B).



Figure 3. Effect of MFR on the protein and mRNA expression levels of NFATc1, c-Fos and TRAF6. (A) NFATc1 expression was determined by western blotting and reverse transcription-PCR, and (B) normalized to β -actin. (C) Protein and mRNA expression levels of c-Fos were determined and (D) normalized to β -actin. (E) Protein expression levels of TRAF6 were measured and (F) normalized to β -actin. Data are presented as the mean \pm standard error of the mean of three independent experiments. All data were analyzed using one-way ANOVA followed by Tukey's post hoc test. ^aP<0.05 vs. normal group (untreated cells); ^bP<0.05 vs. RANKL treatment group; ^cP<0.05 vs. MFR 12.5 μ g/ml treatment group; ^dP<0.05 vs. MFR 25 μ g/ml treatment group; ^cP<0.05 vs. MFR 50 μ g/ml treatment group. MFR, *Melandrium firmum* Rohrbach; RANKL, receptor activator of nuclear factor- κ B ligand; NFATc1, nuclear factor of activated T-cells; TRAF6, tumor necrosis factor receptor-associated factor 6.

While E_2 and ALN groups exhibited a preventive effect on the reduction of femur weight compared with the OVX group, MFR-L and MFR-H groups did not (Fig. 5C). To investigate the extent of hepatotoxicity following treatment with MFR, E_2 and ALN, the serum levels of AST and ALT were measured. Previous studies have shown that levels above 150 U/l for AST and 40 U/l for ALT indicate hepatotoxicity in rats (40,41). The levels of AST and ALT in the MFR-L, MFR-H, E_2 and ALN groups in the present study did not exceed these values; therefore, it was indicated that MFR-L, MFR-H, E_2 and ALN did not induce hepatotoxicity (Fig. 5D and E). To determine the effect of MFR on bone formation, the serum levels of ALP were measured. ALP was significantly increased in the OVX group compared with that in the sham group; moreover, the MFR-L



Figure 4. Effect of MFR on the expression levels of osteoclast-related genes. (A) Western blot analysis of CTK and MMP-9. (B) Protein expression levels were normalized to β -actin. (C) mRNA expression levels of *Mmp9*, *Ctsk* and *Ca2* were analyzed using RT-PCR. (D) mRNA expression levels were normalized to *Actb*. (E) RT-PCR analysis was used to determine the mRNA expression levels of *Acp5*, *Atp6v0d2* and *Dcstamp*, and (F) expression was normalized to *Actb*. (G) RT-PCR analysis was performed to determine the mRNA expression levels of *Oscar*, *Src* and *Prdm1*, and (H) expression was normalized to *Actb*. All data are presented as the mean ± standard error of the mean of three independent experiments. Data were analyzed using one-way ANOVA followed by Tukey's post hoc test. ^aP<0.05 vs. normal group (untreated cells); ^bP<0.05 vs. RANKL treatment group; ^cP<0.05 vs. MFR 12.5 µg/ml treatment group; ^dP<0.05 vs. MFR 25 µg/ml treatment group; ^eP<0.05 vs. MFR 50 µg/ml treatment group. MFR; *Melandrium firmum* Rohrbach; RANKL, receptor activator of nuclear factor-κB ligand; CTK/*Ctsk*, cathepsin k; MMP-9/*Mmp9*, matrix metalloproteinase-9; CA2/*Ca2*, carbonic anhydrase 2; RT-PCR, reverse transcription-PCR; *Acp5*, tartrate-resistant acid phosphatase; *Atp6v0d2*, ATPase H+ transporting V0 subunit D2; *Dcstamp*, dendritic cell-specific transmembrane protein; *Oscar*, osteoclast-associated receptor; *Pdrm1*, B lymphocyte-induced maturation protein-1.

group exhibited decreased serum levels of ALP, whereas the MFR-H, E_2 and ALN groups did not (Fig. 5F). To examine the effect of MFR on bone resorption, serum levels of CTX-1

and TRAP were measured. CTX-1 levels were increased in the OVX group compared with those in the sham group, whereas they were significantly decreased in the MFR-L, E_2 and ALN



Figure 5. Effect of MFR on body, uterus and femur weights, and serum levels of ALP, AST and ALT. (A) Body weight was measured weekly for 8 weeks. (B) Uterus and (C) femur weights were measured after sacrificing the rats. The serum levels of (D) AST, (E) ALT, (F) ALP, (G) CTX-1 and (H) TRAP were measured using enzyme-linked immunosorbent assays. All data are presented as the mean \pm standard error of the mean (n=8 per group). (A) Data were analyzed using two-way ANOVA followed by Bonferroni post hoc test. (B-H) Data were analyzed using one-way ANOVA followed by Tukey post hoc test. ^aP<0.05 vs. sham group; ^bP<0.05 vs. OVX group; ^cP<0.05 vs. MFR-L group; ^dP<0.05 vs. MFR-H group; ^cP<0.05 vs. E₂ group. MFR, *Melandrium firmum* Rohrbach; OVX, ovariectomized; ALP, alkaline phosphatase; AST, aspartate aminotransferase; ALT, alanine aminotransferase; CTX-1, C-telopeptide of type I collagen; TRAP, tartrate-resistant acid phosphatase; MFR-L, low dose of MFR; MFR-H, high dose of MFR; E, β -estradiol; ALN, alendronate.

groups compared with those in the OVX group; there were no significant changes in the MFR-H group (Fig. 5G). As shown in Fig. 5H, TRAP levels were increased due to OVX, but the difference was not significant, whereas oral administration of MFR-L, E_2 and ALN reduced TRAP levels. Furthermore, the TRAP levels were decreased in the MFR-H group; however, this finding was not significant.

Effect of MFR on the bone microarchitecture of ovariectomy-induced osteoporosis in rats. In order to investigate the protective effects of MFR on ovariectomy-induced bone loss, micro-CT was used. As shown in Fig. 6A, micro-CT images of the femoral head indicated trabecular bone loss in OVX rats. BMD loss was significantly inhibited in the MFR-L, E_2 and ALN groups compared with that in the OVX group (Fig. 6B). Furthermore, Tb.Th in the OVX group was

reduced compared with that in the sham group, whereas Tb.Th was significantly increased in the ALN group. However, Tb.Th in the MFR-L, MFR-H and E_2 groups did not differ significantly from that in the OVX group (Fig. 6C). As shown in Fig. 6D, Tb.sp was significantly increased in the OVX group compared with that in the sham group, whereas Tb.sp levels were decreased in the E_2 and ALN groups. However, Tb.sp levels were not affected in the MFR-L and MFR-H groups.

Effect of MFR on bone loss in the OVX rat model of osteoporosis. To investigate the histological changes in the femoral head, H&E staining was performed on the bone tissue (Fig. 7A). The trabecular area in the femoral head of OVX rats was significantly decreased compared with that in the sham group, whereas this reduction was prevented by treatment with MFR-L, E_2 and ALN (Fig. 7B).



Figure 6. Effects of MFR on bone microarchitecture of OVX-induced osteoporosis rats. (A) Femurs were analyzed using micro-CT imaging. (B) BMD, (C) Tb. Th and (D) Tb.sp were analyzed using micro-CT image analysis. All data are presented as the mean \pm standard error of the mean (n=8/group). Statistical analyses were performed using one-way ANOVA followed by Tukey post hoc test. ^aP<0.05 vs. sham group; ^bP<0.05 vs. OVX group; ^cP<0.05 vs. MFR-L group; ^dP<0.05 vs. MFR-H group; ^cP<0.05 vs. E₂ group. MFR, *Melandrium firmum* Rohrbach; OVX, ovariectomized; BMD, bone mineral density; Tb.Th, trabecular thickness; Tb.sp, trabecular spacing; micro-CT, micro computed tomography; MFR-L, low dose of MFR; MFR-H, high dose of MFR; E₂, β -estradiol; ALN, alendronate.



Figure 7. Effect of MFR on trabecular area in OVX-induced rats. (A) Bone tissues were stained with H&E. Magnifications, x40 and x100; scale bar, 500 and 200 μ m, respectively. (B) Trabecular area was analyzed in the H&E-stained sections. All data are presented as the mean ± standard error of the mean (n=8/group). Statistical analyses are performed using one-way ANOVA followed by Tukey post hoc test. ^aP<0.05 vs. sham group; ^bP<0.05 vs. OVX group; ^cP<0.05 vs. MFR-L group; ^dP<0.05 vs. MFR-H group. MFR, *Melandrium firmum* Rohrbach; OVX, ovariectomized; H&E, hematoxylin and eosin; MFR-L, low dose of MFR; MFR-H, high dose of MFR; E₂ β-estradiol; ALN, alendronate.



Figure 8. Effects of MFR on the expression levels of CTK and NFATc1 in OVX-induced rats. (A) NFATc1 and (B) CTK-positive expression in bone tissue was assessed using IHC. NFATc1 and CTK-positive expression is highlighted by the red arrowheads. Mean number of (C) NFATc1 and (D) CTK-positive cells in femur tissues was counted. Magnifications, x100 and x200; scale bar, 200 and 100 μ m, respectively. All data are presented as the mean ± standard error of the mean (n=8/group). Statistical analyses are performed using one-way ANOVA followed by Tukey post hoc test. ^aP<0.05 vs. sham group; ^bP<0.05 vs. OVX group; ^oP<0.05 vs. MFR-L group. MFR, *Melandrium firmum* Rohrbach; CTK, cathepsin k; NFATc1, nuclear factor of activated T-cells, cytoplasmic 1; OVX, ovariectomized; IHC, immunohistochemistry; MFR-L, low dose of MFR; MFR-H, high dose of MFR; E₂ β -estradiol; ALN, alendronate.

Effect of MFR on the expression levels of NFATc1 and CTK in femoral tissues in the in vivo osteoporosis model. To determine the effect of MFR on the expression levels of NFATc1 and CTK in the rat model of osteoporosis, IHC was performed (Fig. 8A and B). The OVX group exhibited increased expression levels of NFATc1 compared with those in the sham group. By contrast, the MFR-L and E_2 groups exhibited significantly reduced expression levels of NFATc1 expression; no changes were observed in the MFR-H and ALN groups compared with the OVX group (Fig. 8C). In OVX rats, the expression levels of CTK were increased compared with those in the sham group. Conversely, MFR-L, MFR-H, E_2 and ALN effectively reduced the expression levels of CTK (Fig. 8D).

Effect of MFR on mineralization of MC3T3-E1 cells. MC3T3-E1 cells derived from mouse calvaria are characterized by proliferation, differentiation and mineralization of osteoblasts, and are used in studies related to bone formation (42). Alizarin red staining can be used to detect the formation of mineralization (43). Therefore, to confirm the effect of MFR on osteoblast differentiation and mineralization, Alizarin red S staining was performed. The control cells exhibited significantly increased mineralization compared with that in the normal cells. Treatment with low concentrations of MFR (12.5 and 25 μ g/ml) promoted osteoblast differentiation, but high concentrations of MFR (50 and 100 μ g/ml) inhibited osteoblast differentiation (Fig. 9A). To examine the effects of the MFR extract on MC3T3-E1 cell viability, cells were treated with 12.5-100 μ g/ml MFR. Notably, MFR did not affect the viability of MC3T3-E1 cells (Fig. 9B). Quantification of Alizarin red staining revealed a significantly reduced effect of MFR in a dose-dependent manner (Fig. 9C).

Discussion

The aim of the present study was to investigate the inhibitory effect of MFR extract on osteoclast differentiation and bone loss in ovariectomy-induced postmenopausal osteoporotic rats. *In vitro*, MFR inhibited RANKL-induced osteoclast differentiation, formation and function. In addition, MFR inhibited the TRAF6 axis and inhibited the expression of NFATc1/c-Fos, which is known as a key factor for osteoclast differentiation. Finally, MFR suppressed the expression of osteoclast-related genes, such as *Acp5*, *Mmp9*, *CtsK*, *Atp6v0d2*, *Dcstamp* and *Src*;



Figure 9. Effects of MFR on the production of mineralization in MC3T3-E1 cells. (A) MFR treatment suppressed mineralization in MC3T3-E1 cells. (B) Cell viability was determined using the MTS assay. (C) Quantitative analysis of mineralization was performed using an enzyme-linked immunosorbent assay reader. Data were analyzed using one-way ANOVA followed by Tukey's post hoc test. $^{\circ}P<0.05$ vs. normal group (untreated cells); $^{\circ}P<0.05$ vs. osteogenic medium treatment group; $^{\circ}P<0.05$ vs. MFR 12.5 μ g/ml treatment group; $^{\circ}P<0.05$ vs. MFR 25 μ g/ml treatment group; MFR, *Melandrium firmum* Rohrbach.

this potential mechanism of suppression is shown in Fig. 10. *In vivo*, MFR-L considerably increased BMD, trabecular area, and decreased the expression levels of NFATc1 and CTK in OVX-induced models.

Murine macrophage RAW 264.7 cells have a monocyte/macrophage like cell lineage; they are derived from BALB/c mice (44) and are suitable as a cell experimental model in measuring osteoclast differentiation (45). TRAP staining is the standard method used to detect osteoclast formation and levels (31,46,47). In the present study, MFR suppressed RANKL-induced TRAP-positive cells and TRAP levels. The F-actin ring is the cytoskeletal structure that is required for formation of the sealing zone, and permits the firm adhesion of mature osteoclasts to the bone surface (48). The pit formation assay is widely used to examine bone resorption and is an important indicator of bone resorption from mature osteoclasts (48-50). In the present study, MFR reduced the number and size of the F-actin rings and the area of pit formation in RANKL-induced cells. These results suggested that MFR may inhibit osteoclast formation and bone resorption.

The RANK/RANKL signaling pathway has been reported to induce the activation of TRAF6. TRAF6 deficiency in mice is known to induce severe osteopetrosis and impairment in osteoclast function (51-53). Activated TRAF6 can stimulate c-Fos, which serves a crucial role in the induction of transcription factors, including NFATc1 (54-56). In a previous study, c-Fos-knockout mice developed osteopetrosis due to attenuated osteoclast function (20). In addition, NFATc1-deficient adult mice exhibited reduced bone loss in the absence of osteoclast activity (17,19). In the present study, MFR was shown to reduce the protein and mRNA expression levels of c-Fos and NFATc1. These results suggested that MFR may decrease the production of osteoclast differentiation, bone resorption and F-actin ring formation by suppressing NFATc1 and c-Fos signaling pathways. However, in contrast to the results regarding NFATc1/c-Fos, the present study revealed that MFR significantly inhibited TRAF6 at low concentrations but not at high concentrations. The reason for the difference in the expression levels of each factor can be inferred due to the following reasons. RANKL binds to RANK, increases the expression and ubiquitination of TRAF6, and leads to the accumulation of the TRAF6/transforming growth factor- β -activated kinase 1 (TAK1) complex, which activates sub-signaling molecules, such as phosphoinositide 3-kinases, MAPK and NF-KB (57,58). It was hypothesized that treatment with a high concentration of MFR in the present study could inhibit the ubiquitination of TRAF6 or the accumulation of the TRAF6/TAK1 complex, rather than affecting the expression of TRAF6. However, additional research needs to be conducted to assess this hypothesis.

NFATc1 regulates several important osteoclast-related genes, such as *Mmp9*, *Ctsk* and *Ca2*, which are markers of bone resorption (17). *Mmp9* and *Ctsk* are well-known as bone resorption enzymes. Notably, *Mmp9* is expressed during the transformation of early osteoclasts to mature osteoclasts (59) and deficiency of *Ctsk* leads to impairments in bone resorption in mice and humans (60). Furthermore, *Ca2* is an enzyme that is upregulated by c-Fos signaling, and is expressed in the early



Figure 10. Inhibitory mechanisms of MFR on osteoclast differentiation. MFR, *Melandrium firmum* Rohrbach; RANKL, receptor activator of nuclear factor-κB ligand; RANK, receptor activator of nuclear factor-κB; TRAF6, tumor necrosis factor receptor-associated factor 6; NF-κB, nuclear factor-κB; NFATc1, nuclear factor of activated T cells, cytoplasmic 1; MMP-9, matrix metalloproteinase-9; ATP6v0d2, ATPase H+ transporting V0 subunit D2; DC-STAMP, dendritic cell-specific transmembrane protein; TRAP, tartrate-resistant acid phosphatase.

stages of osteoclast differentiation and influences the acidity of the bone surface (61). In the present study, MFR suppressed the expression levels of *mmp9*, *ctsk* and *ca2*. These results suggested that MFR may possess anti-osteoporotic activity by regulating the expression of bone resorption markers. Atp6vod2 and Destamp serve an important role in cell-cell fusion during RANKL-stimulated osteoclast differentiation. In addition, Atp6vod2 and Dcstamp are important for F-actin ring formation (39,62). Notably, gene-knockout mice of Atp6vod2 and Destamp have been reported to develop osteopetrosis due to defects in osteoclastogenesis (62,63). In the present study, MFR suppressed the expression levels of *Atp6vod2* and *Dcstamp*. These results indicated that the inhibitory effect of MFR on F-actin ring formation may be associated with the suppression of Atp6vod2 and Dcstamp. Oscar serves an important role in the bone-specific regulation of osteoclast differentiation, and activation of *Oscar* results in differentiation of early osteoclasts to mature osteoclasts. In addition, Oscar can induce calcium activation, resulting in increased expression of NFATc1 (64). Src signaling is important in regulating the osteoclast cytoskeleton (65). Prdml serves an essential role in the differentiation and functions of macrophages and lymphocytes (66). In addition, Prdml controls osteoclast development and bone homeostasis (67). In the present study, MFR decreased the expression levels of Oscar, Src and Prdm1. These results suggested that MFR may inhibit the expression of osteoclast-related genes by suppressing NFATc1 signaling. Notably, several genes were expressed in the form of two bands in the present study; as a result of comparing the primer size using a DNA ladder marker, the band at the top is considered the target, and the bottom is inferred as the dimer remaining after the reaction was completed.

Osteoporosis is the most common type of bone disease worldwide, which is characterized by a low bone mass (68). Estrogen deficiency is the primary cause of postmenopausal osteoporosis (69). Rats with ovariectomy-induced osteoporosis have been used as a postmenopausal osteoporosis model and share clinical characteristics with human osteoporosis (3). It has been established that an increase in body weight is a symptom observed in OVX rats. In addition, reduction in uterus weight is indicative of successful establishment (70). In the present study, the body weight of animals was significantly increased 3 weeks after OVX and uterus weight was significantly decreased in the OVX group compared with that in the sham group. These results suggested that a postmenopausal osteoporosis model was successfully established.

ALP is a marker commonly associated with bone formation, which is produced during the early differentiation of osteoblasts (71). Wu et al (72) demonstrated that ALP activity was increased in estrogen-deficient mice. Excessive osteoclast activity causes an imbalance in bone metabolism, resulting in increased osteoblast activity and ALP expression. In the present study, the serum ALP levels were significantly increased in the OVX group compared with those in the sham group, whereas ALP levels were significantly decreased in the MFR-L group. These results suggested that the MFR-L group reduced the levels of ALP, which were increased due to excessive osteoclast activity. AST and ALT are the most commonly used factors for assessment of hepatotoxicity (73). The results of the present study suggested that the levels of AST and ALT were not affected in OVX rats. In addition, MFR-L, MFR-H, E₂ and ALN did not exert hepatotoxic effects. CTX is the most widely used indicator to measure bone resorption, and CTX-1 and TRAP are bone resorption markers of osteoclasts that have been reported to be increased in OVX models (74). In the present study, CTX-1 levels were significantly increased in the OVX group, whereas they were reduced in the MFR-L, E2 and ALN groups; CTX-1 levels in the MFR-H group were unchanged. In addition, TRAP levels were increased in the OVX group; however, this difference was not significant. By contrast, TRAP levels were reduced in the MFR-L, E2 and ALN groups; TRAP levels were also reduced in the MFR-H group, but this change was not significant.

Micro-CT has been used to study bone tissues and in orthopedic studies (75). The advantage of micro-CT is the lack of damage to the sample and ease of reconstruction using image sections (76). BMD refers to the amount of mineral content in bone tissue and is indicative of the strength of bones; BMD is based on calcium content and is an important measure in the evaluation of the OVX rat model (77). In the present study, the BMD in the OVX group was significantly decreased compared with that in the sham group, whereas BMD loss was reduced in the MFR-L, E₂ and ALN groups. Tb.Th and Tb.sp are indicators used in assessment of the 3D image structure of cancellous bone (78). In the present study, the results revealed that the OVX group exhibited decreased Tb.Th and increased Tb.sp compared with those in the sham group. Tb.Th was increased following ALN treatment compared with that in the OVX group, but no significant effect was observed on Tb.Th in the OVX rats treated with MFR-L, MFR-H and E₂. Tb.sp

was decreased in the E_2 and ALN groups compared with that in the OVX group, but no significant effect was observed in the MFR-L and MFR-H groups. These findings indicated that oral administration of MFR-L reduced bone loss.

Decreased trabecular area is a symptom commonly observed in patients with osteoporosis and is the primary cause of an increased risk of fracture; thus, the trabecular area has been used as an important index of anti-osteoporotic activity in a previous study (79). H&E staining can also be used to assess the trabecular area in the femur alongside micro-CT (80-84). Bone loss of the femoral head was enhanced in the OVX rats, whereas MFR-L, E_2 and ALN reduced trabecular area loss in the OVX-induced rat models. These results suggest that MFR-L, E_2 and ALN inhibited OVX-induced bone loss. IHC demonstrated that the expression levels of NFATc1 were decreased following treatment with MFR-L and E_2 . In addition, administration of MFR-L, MFR-H, E_2 and ALN reduced the expression levels of CTK in the OVX rats.

The present study confirmed the effect of MFR on osteoblast differentiation, which has an important role in bone metabolism. MFR (12.5 and 25 μ g/ml) promoted osteoblast differentiation, whereas MFR (50 and 100 μ g/ml) suppressed osteoblast differentiation. These results indicated that high concentrations of MFR can simultaneously inhibit osteoclast and osteoblast differentiation. These results may be related to the insignificant effect of the MFR-H group compared to the MFR-L in bone loss in the OVX-induced rat model.

In conclusion, the effects of MFR on RANKL-induced osteoclast differentiation and ovariectomy-induced bone loss in a SD-rat model were determined. *In vitro*, MFR reduced RANKL-induced osteoclast differentiation, function and formation. MFR downregulated the expression levels of master transcription factors, such as NFATc1 and c-Fos. In addition, MFR reduced the expression levels of MMP-9, CTK, CA2, TRAP, ATP6v0d2, DC-STAMP, OSCAR, c-Src and Blimp-1 through the downregulation of NFATc1 and c-Fos signaling. *In vivo*, MFR-L increased BMD in the OVX-induced bone loss model. MFR-H exhibited insignificant effects on the OVX-induced bone loss model compared with MFR-L.

The present study had several limitations, as follows: i) In vitro, MFR inhibited RANKL-induced expression of NFATc1 and c-Fos signaling. MAPK and NF-KB signaling pathways are also involved in NFATc1 and c-Fos signaling; however, the effect of MFR was not determined on these factors. To further assess the mechanisms underlying the inhibitory effects of MFR on osteoclast differentiation via the NFATc1 and c-Fos pathway, further studies regarding the involvement of MAPKs and NF-KB are required. ii) When comparing the pharmacological effects of MFR in the in vitro experiments and in vivo experiments, MFR showed a relatively low pharmacological effect in in vivo experiments compared with that in the *in vitro* experiments. Therefore, further research on the effects of MFR on osteoclast-mediated bone diseases, such as in the LPS-induced bone loss model, senile osteoporosis and osteoporosis caused by steroid side effects, are necessary. iii) In the micro-CT test results, MFR-L increased BMD; however, it did not significantly affect the bone micro-architecture. In general, an increase in BMD is associated with changes in the bone micro-architecture; however, this is not the case in this study. Further studies are required to determine the reason behind this. iv) The present study focused on the effects of MFR on inhibition of osteoclast differentiation; According to previous studies, MFR contains a variety of components (85), and constituents of MFR, such as vitexin, linarin, ecdysterone and ursolic acid, were found to be effective in inhibiting osteoclast differentiation. Vitexin inhibited osteoclast differentiation and osteolysis (28), and Wang et al (84) revealed the inhibitory activity of linarin on osteoclastogenesis through the RANKL-induced NF-KB pathway. In addition, ecdysterone, another active compound in MFR, prevented LPS-induced osteoclastogenesis (86). Ursolic acid has also been shown to inhibit osteoclastogenesis and titanium particle-induced osteolysis (87). However, to the best of our knowledge, the effects of most of the components of MFR on osteoclast differentiation have not been studied. Therefore, analyzing the effect of each component of MFR on osteoclast differentiation will help to understand the underlying anti-osteoporotic mechanism of MFR.

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Availability of data and materials

All data generated or analyzed during this study are included in this published article.

Authors' contributions

YS designed the study. MK and BK prepared the extract. MK, JHK and EYK performed the *in vitro* experiments and analyzed the data. MK, JHK, SH and BK performed the *in vivo* experiments and analyzed the data. MK, JHK and HSJ contributed to the statistical analysis and helped interpret the results. YS supervised the experiments in discussion with JHK and MK. MK and JHK wrote the manuscript. MK and JHK confirm the authenticity of all the raw data. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The protocols for this experiment were approved by the Kyung Hee Medical Science Research Institute Animal Care and Use Committee (approval no. KHMC-IACUC 19-017).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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