Release from optimal compressive force suppresses osteoclast differentiation

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Abstract. Bone remodeling is an important factor in orthodontic tooth movement. During orthodontic treatment, osteoclasts are subjected to various mechanical stimuli, and this promotes or inhibits osteoclast differentiation and fusion. It has been previously reported that the release from tensile force induces osteoclast differentiation. However, little is known about how release from compressive force affects osteoclasts. The present study investigated the effects of release from compressive force on osteoclasts. The number of tartrate-resistant acid phosphatase (TRAP)-positive multinucleated osteoclasts derived from RAW264.7 cells was counted, and gene expression associated with osteoclast differentiation and fusion in response to release from compressive force was evaluated by reverse transcription-quantitative polymerase chain reaction. Osteoclast number was increased by optimal compressive force application. On release from this force, osteoclast differentiation and fusion were suppressed. mRNA expression of NFATc1 was inhibited for 6 h subsequent to release from compressive force. mRNA expression of the other osteoclast-specific genes, TRAP, RANK, matrix metalloproteinase-9, cathepsin-K, chloride channel 7, ATPase H+ transporting vacuolar proton pump member 1, dendritic cell-specific transmembrane protein and osteoclast stimulatory transmembrane protein (OC-STAMP) was significantly inhibited at 3 h following release from compressive force compared with control cells. These findings suggest that release from optimal compressive force suppresses osteoclast differentiation and fusion, which may be important for developing orthodontic treatments.

Introduction

Bone remodeling is a balance between bone resorption by osteoclasts and bone formation by osteoblasts (1). When this balance tips toward excess resorption, the risk of osteoporosis is increased, whereas osteoclast dysfunction increases the risk of osteopetrosis (2). Thus, osteoclasts have an important function in bone homeostasis. In orthodontic treatment, osteoclasts are also important for tooth movement (3,4). Promotion of osteoclast differentiation accelerates bone resorption. By contrast, inhibiting osteoclast differentiation accelerates formation. Orthodontic force consists of tensile and compressive force. On the pressure side, osteoclasts are subjected to compressive force. By contrast, on the tension side, osteoclasts are subjected to tensile force. During orthodontic treatment, numerous stimuli are applied to osteoclasts; certain signals promote osteoclast differentiation, whereas others inhibit osteoclast differentiation (3).

Osteoclasts are multinucleated, bone-resorbing cells that are differentiated from the monocyte/macrophage hematopoietic lineage (5). Receptor activator of nuclear factor-κB (RANK)-ligand (RANKL) is an essential factor for osteoclast differentiation. RANKL binding to RANK induces expression of nuclear factor of activated T cells 1 (NFATc1). NFATc1 is the master transcription factor for osteoclast differentiation (6). An increase in the expression of NFATc1 promotes transcription of various osteoclast-specific genes. Numerous studies have previously reported that stimulation of osteoclasts leads to secretion of cytokines associated with bone resorption and formation (7-10).

In a previous study, various types of mechanical stimuli were applied to osteoclasts, and it was reported that these were influential factors in bone remodeling. Mechanical stimuli include tensile force (8,9,11-14), compressive force (7,10,15-17), hydrostatic pressure (18), shear stress (19,20), rotative stress (21) and others (22,23). Stimulation with tensile force using a Flexercell tension system suppresses osteoclast differentiation and fusion. The number of osteoclasts increases

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rapidly after the release of tensile force. Additionally, optimal compressive force induces osteoclast differentiation (9). In this experiment, osteoclasts on slips are reversed and placed on collagen gel layers, and compressive force was adjusted using weights. Optimal compressive force is defined as the weight that induces the largest increase in the number of osteoclasts, and is 280 mg/cm² according to Hayakawa et al (10). Thus, the present study investigated the effects of release from optimal compressive force on osteoclasts.

Materials and methods

Cell culture. The current study used the murine monocyte/macrophage cell line RAW264.7 cells (TIB-71™; American Type Culture Collection, Manassas, VA, USA) as osteoclast precursors. Cells were maintained in Dulbecco’s modified Eagle’s medium (Wako Pure Chemical Industries, Ltd., Osaka, Japan) containing 10% heat-inactivated fetal bovine serum (FBS; Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA) and 66.7 µg/ml kanamycin sulfate (Meiji Seika Kaisha, Ltd., Tokyo, Japan) at 37°C in 5% CO₂ in humidified air. Cells were seeded in 100-mm standard dishes (Corning Incorporated, Corning, NY, USA) and incubated overnight. Subsequently, for osteoclast differentiation, RAW cells (1x10⁴ cells/well) were transferred into 24-well culture plates (Corning Incorporated) and cultured on 12-mm diameter glass cover slips (Fisher Microscope Cover Glass; Thermo Fisher Scientific, Inc.) placed on the 24-well culture plate, in α-minimum essential medium (α-MEM; Wako Pure Chemical Industries, Ltd.) supplemented with 10% heat-inactivated FBS, 2 mM L-alanyl-L-glutamine (Wako Pure Chemical Industries, Ltd.), 284 µM L-ascorbic acid phosphate magnesium salt n-hydrate (Wako Pure Chemical Industries, Ltd.), 66.7 µg/ml kanamycin sulfate, and 50 ng/ml RANKL (Oriental Yeast Co., Ltd., Tokyo Japan) at 37°C under 5% CO₂ in humidified air. Medium was changed every other day.

Preparation of collagen gels. The collagen mixture comprised acid-soluble collagen solution (Cellmatrix; Nitta Gelatin NA Inc., Morrisville, NC, USA) mixed with 10-fold concentrated α-MEM and reconstruction buffer (2.2 g NaHCO₃, + 4.77 g HEPES in 100 ml 0.05 N NaOH; Nitta Gelatin NA Inc.) at a volume ratio of 8:1:1, and supplemented with 10% heat-inactivated FBS (Invitrogen; Thermo Fisher Scientific, Inc.), 284 µM L-ascorbic acid 2-phosphate (Sigma-Aldrich; Merck Millipore), and 50 ng/ml RANKL (Oriental Yeast Co., Ltd.). The collagen mixture was isolated from cultured cells under each set of conditions with TRIzol (Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer’s instructions, and aliquots containing equal amounts of mRNA were subjected to RT-qPCR. First-strand cDNA synthesis was performed using 1 µg total RNA, 25 pmol oligo dT (Toyobo Co., Ltd., Osaka, Japan), 1 mM dNTP (Toyobo Co., Ltd.), 100 U ReverTra Ace® (Toyobo Co., Ltd.), 20 U RNase inhibitor (Toyobo Co., Ltd.) and 4 µl 5X reaction mixture (Toyobo Co., Ltd.) in 20 µl with annealing at 30°C for 10 min, enzyme reaction at 42°C for 20 min, denaturation at 99°C for 5 min and cooling at 4°C. qPCR was performed using the ABI Prism 7300 sequence detection system (Applied Biosystems; Thermo Fisher Scientific, Inc.). The reactions were incubated at 50°C for 2 min and 95°C 10 min, followed by 40 cycles of 95°C for 15 sec and annealing at 60°C for 1 min. The following specific TaqMan probes (Applied Biosystems; Thermo Fisher Scientific, Inc.) for osteoclast-associated genes were used: NFATc1 (ID no. Mm00479445_ml), TRAP (ID no. Mm00475698_ml), matrix metalloproteinase-9 (MMP-9; ID no. Mm00432271_ml), RANK (ID no. Mm00437135_ml), osteoclast stimulatory trans membrane protein (OC-STAMP; ID no. Mm00512445_ml), cathepsin-K (Cath-K; ID no. Mm00484036_ml), chloride channel 7 (CIC-7; ID no. Mm00442400_ml) and ATPase H⁺ transporting vacuolar proton pump member I (ATP6i; ID no. Mm00469395_ml). Levels of mRNA expression were calculated and standardized against the level of glyceraldehyde 3-phosphate dehydrogenase (GAPDH; ID no. Mm99999915_ml) mRNA. Following each PCR run, data were analyzed by the system and amplification plots

Release from compressive forces. After cells had been compressed and incubated for 24 h, weights were removed, and cells on slips were inverted. Subsequently, cells were incubated for 24 h.

Tartrate-resistant acid phosphatase (TRAP) staining. After cells were cultured for a 2, 3, 4, 5 or 6 days, and fixed with 10% neutral formalin for 30 min at room temperature, they were washed with distilled water and treated with TRAP staining solution (pH 5.0) supplemented with Fast Red Violet LB Salt (Sigma Aldrich; Merck Millipore) (24). TRAP staining solution contained acetate buffer (pH 5.0; Sigma-Aldrich; Merck Millipore), naphthol AS-MX phosphate (Sigma-Aldrich; Merck Millipore) as a substrate, red violet LB (Sigma-Aldrich; Merck Millipore) as a stain in the presence of 50 mM sodium tartrate (Wako Pure Chemical Industries, Ltd.). TRAP-positive cells with ≥2 nuclei were counted under the microscope as osteoclasts. TRAP-positive cells with 2-7 nuclei were considered to be small osteoclasts, and those with ≥8 nuclei to be large osteoclasts (8,9).

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Cells were incubated for 1, 3, 6, 12 and 24 h in 24-well culture plates, after compression for 24 h. Total RNA was isolated from cultured cells under each set of conditions (TRIzol; Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer’s instructions, and aliquots containing equal amounts of mRNA were subjected to RT-qPCR. First-strand cDNA synthesis was performed using 1 µg total RNA. 5 pmol oligo dT (Toyobo Co., Ltd., Osaka, Japan), 1 mM dNTP (Toyobo Co., Ltd.), 100 U ReverTra Ace® (Toyobo Co., Ltd.), 20 U RNase inhibitor (Toyobo Co., Ltd.) and 4 µl 5X reaction mixture (Toyobo Co., Ltd.) in 20 µl with annealing at 30°C for 10 min, enzyme reaction at 42°C for 20 min, denaturation at 99°C for 5 min and cooling at 4°C. qPCR was performed using the ABI Prism 7300 sequence detection system (Applied Biosystems; Thermo Fisher Scientific, Inc.). The reactions were incubated at 50°C for 2 min and 95°C 10 min, followed by 40 cycles of 95°C for 15 sec and annealing at 60°C for 1 min. The following specific TaqMan probes (Applied Biosystems; Thermo Fisher Scientific, Inc.) for osteoclast-associated genes were used: NFATc1 (ID no. Mm00479445_ml), TRAP (ID no. Mm00475698_ml), matrix metalloproteinase-9 (MMP-9; ID no. Mm00432271_ml), RANK (ID no. Mm00437135_ml), osteoclast stimulatory trans membrane protein (OC-STAMP; ID no. Mm00512445_ml), cathepsin-K (Cath-K; ID no. Mm00484036_ml), chloride channel 7 (CIC-7; ID no. Mm00442400_ml) and ATPase H⁺ transporting vacuolar proton pump member I (ATP6i; ID no. Mm00469395_ml). Levels of mRNA expression were calculated and standardized against the level of glyceraldehyde 3-phosphate dehydrogenase (GAPDH; ID no. Mm99999915_ml) mRNA. Following each PCR run, data were analyzed by the system and amplification plots
were obtained. qPCR results were calculated using the $2^{-\Delta\Delta C_q}$ method (26).

**Statistical analysis.** Values represent the mean ± standard deviation. Comparisons between two groups were analyzed by 2-tailed unpaired Student's t test. P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Number of osteoclasts on various culture days.** In order to determine when osteoclast differentiation and fusion were activated, the number of osteoclasts and the nuclei in osteoclasts were observed for 6 days. Osteoclasts induced from RAW264.7 cells were cultured with 50 ng/ml RANKL on 24-well culture plates for 6 days. The number of osteoclasts increased rapidly on day 5, and marginally on day 6 (Fig. 1A). Small osteoclasts (2-7 nuclei) did not increase in number, but large osteoclasts (≥8 nuclei) increased on day 6 (Fig. 1B). Even in small osteoclasts, osteoclasts with 2-3 nuclei decreased, however osteoclasts with 4-7 nuclei did not increase (Fig. 1C).

**Number of osteoclasts following release from compressive force during osteoclast differentiation.** The effect of the release from compressive force on osteoclasts was also examined. Following release from compressive force, osteoclast differentiation and fusion were decreased in comparison with the control groups. The number of osteoclasts release from compressive force increased 1.7 fold. However, the number in the control groups increased by 2.4 fold. The total number of osteoclasts in the control groups was significantly greater than the cells released from compressive force (P=0.02; Fig. 3A). The number of small osteoclasts (2-7 nuclei) and large osteoclasts (≥8 nuclei) in the control group was increased compared with the cells released from compressive force (P=0.04 and P=0.03, respectively; Fig. 3B).

**Effects of release from compressive force on the expression of osteoclast differentiation genes.** How the expression of osteoclast differentiation genes was altered by release from compressive force was investigated using RT-qPCR analysis...
for osteoclast-specific genes (NFATc1, TRAP, RANK, MMP-9, Cath-K, CIc7 and ATP6i) and fusion-associated factors (DC-STAMP and OC-STAMP). Analysis demonstrated that NFATc1 mRNA levels in the control groups were increased at 0, 1, 3 and 6 h compared with the cells released from compressive force (P=0.0003, P=0.003, P=0.0005 and P=0.05, respectively). However, TRAP, RANK, MMP-9, Cath-K, CIc7, ATP6i, DC-STAMP, OC-STAMP mRNA levels peaked at 3 h in control groups; and were significantly increased compared with the cells that had been released from compressive force (all P<0.01; Fig. 4).

Discussion

Orthodontic tooth movement is achieved by the remodeling of periodontal ligament and alveolar bone in response to mechanical pressure and tension (3,4). During tooth movement, osteoclasts remove bone on the pressure side and osteoblasts create new bone on the tension side of the tooth. Following the release of orthodontic force from teeth, orthodontic relapse occurs. Orthodontic relapse can be defined as the tendency for teeth to return to their pre-treatment position. This is considered to be due to gingival fibers and unbalanced lip-tongue force (27). However, effects of release from orthodontic force on osteoclasts are unknown.

In a previous study, osteoclasts were subjected to various types of force in vitro. During orthodontic treatment, osteoclasts are subjected to compressive force on the pressure side, and tensile force on the tension side. The number of osteoclasts was observed to decrease following the application of tensile force using a Flexercell tension system (9). Osteoclast differentiation was upregulated subsequent to release from tensile force (8). By contrast, optimal compressive force induced osteoclast differentiation and fusion (10). Thus, the aim of the present study was to investigate the effects of release from compressive force on the pressure side of tooth movement in vitro.

In order to investigate the effects of release from optimal compressive force on osteoclast differentiation, the number of TRAP-positive cells was counted and the expression of NFATc1, TRAP, MMP-9, Cath-K, CIc7, ATP6i, DC-STAMP and OC-STAMP mRNA was examined.

The results presented in Fig. 1A suggest that osteoclastogenesis advances rapidly after 4-5 days. Osteoclastogenesis continued to increase up to day 6, therefore, indicating that the most appropriate time to apply compressive force was at days 4-5 and to release compressive force at day 5.

As described above, the optimal compressive force was defined as 280 mg/cm², which induced the greatest increase in osteoclasts in a previous experiment using collagen gels (10).
Figure 4. Effects of release from optimal CF on mRNA levels of osteoclast-associated genes for 1-24 h. mRNA levels of osteoclast-associated genes (A) NFATc1, (B) TRAP, (C) RANK, (D) MMP-9, (E) Cath-K, (F) ClC7, (G) ATP6i, (H) DC-STAMP, and (I) OC-STAMP were evaluated by reverse transcription-quantitative polymerase reaction. Results are presented as the mean ± standard deviation (n=4).

* P<0.05, ** P<0.01, comparison indicated by brackets. cont, control; CF, compressive force; NFATc1, nuclear factor of activated T cells 1; TRAP, tartrate-resistant acid phosphatase; RANK, receptor activator of nuclear factor-κB; MMP-9, matrix metalloproteinase-9; Cath-K, cathepsin-K; ClC7, chloride channel 7; ATP6i, ATPase H+ transporting vacuolar proton pump member I; DC-STAMP, dendritic cell-specific transmembrane protein; OC-STAMP, osteoclast stimulatory transmembrane protein.
Optimal compressive force had been applied to osteoclasts for 24 h; the number of TRAP-positive multinucleated osteoclasts was increased in the compressive force cells compared with the control groups. This suggests that optimal compressive force promoted osteoclast differentiation and fusion.

Subsequent to the release from optimal compressive force, the number of TRAP-positive multinucleated osteoclasts were counted. The number of osteoclasts in the control group was increased compared with the cells that were released from compressive force. This suggests that release from optimal compression suppresses osteoclast differentiation. Thus, how the expression of osteoclast differentiation genes was altered by release from compressive force was subsequently investigated using RT-qPCR analysis.

NFATc1 mRNA levels in the control groups were increased at 0, 1, 3 and 6 h compared with cells that were release from compressive force. Thus, NFATc1 mRNA expression was inhibited for 6 h subsequent to release from compressive force. NFATc1 is the master switch for osteoclast differentiation (28). The increasing of expression of NFATc1 lead to expression of other osteoclast-specific genes increased in the control group. TRAP, RANK, MMP-9, Cath-K, CIC7, ATP6i, DC-STAMP and OC-STAMP mRNA levels peaked at 3 h in the control groups and were significantly increased compared with the cells released from compressive force. Although the NFATc1 mRNA level in the control group was higher than the compressive force group at 0 h, these data were consistent with the results of Hayakawa et al (10) and it was noted that compressive force affected the expression of NFATc1 at 0 h. However, release from compressive force also affected the expression of NFATc1 at 1, 3 and 6 h.

The inhibition of the expression levels of TRAP, RANK, MMP-9, Cath-K, CIC7, ATP6i, DC-STAMP and OC-STAMP mRNA peaked at 3 h after release from compressive force. TRAP and RANK are histochemical markers of osteoclasts (29). Inhibition of TRAP and RANK expression indicates a reduction in the number of osteoclasts. MMP-9 expression is essential for the migration of osteoclasts through collagen in the periostium and developing marrow cavity of primitive long bones (30,31). Bone resorption may be significantly reduced by inhibition of MMP-9 (32,33).

Cath-K, CIC7 and ATP6i directly affect bone resorption within the ruffled border of osteoclasts (34-36). This expression is associated with bone resorption and was inhibited following the release from compressive force. In addition to inhibition of MMP-9 mRNA expression, osteoclast migration and resorption may be decreased.

DC-STAMP and OC-STAMP modulate cell-cell fusion in osteoclasts, and are induced by the RANKL-NFATc1 axis (37). Decreased expression of DC-STAMP and OC-STAMP inhibit the fusion of osteoclasts. Thus, the number of large osteoclasts was decreased. The current study demonstrated that optimal compressive force promoted osteoclast differentiation and fusion; however, release from this compressive force suppressed osteoclast differentiation and fusion.

The major causes of orthodontic relapse are considered to be gingival fibers and unbalanced lip-tongue force. We hypothesize that the suppression of osteoclast differentiation and fusion following release from optimal compressive force also has a role in orthodontic relapse. Release from compressive force suppresses osteoclast differentiation, accelerating bone formation, whereas, release from tensile force promotes osteoclast differentiation, accelerating bone resorption (8).

This mechanism may be a factor involved in orthodontic relapse, however, further research is necessary to clarify the mechanism of the relapse following active orthodontic treatment.

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