

Ethanol increases osteoclastogenesis associated with the increased expression of RANK, PU.1 and MITF *in vitro* and *in vivo*

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Abstract. Ethanol has been known to induce osteopenia. However, the cellular and molecular mechanisms responsible for its effect have not been well characterized. This study investigated the effects of ethanol on bone metabolism and osteoclastogenesis using rats fed an ethanol-containing liquid diet (35% of calories from ethanol) for 3 weeks. Ethanol increased the activities of bone tartrate-resistant acid phosphatase (TRAP) and cathepsin K, without affecting the levels of serum osteocalcin or bone alkaline phosphatase activity. Histological analysis showed an increased number of osteoclasts in the proximal tibia, but no significant change in the number of osteoblasts. The mRNA levels of receptor for activation of NF- κ B (RANK), c-fos, c-jun, TRAP and cathepsin K were significantly increased, although those of macrophage colony-stimulating factor and c-fms were unaltered. The mRNA and protein levels of PU.1 and microphthalmia-associated transcription factor (MITF) also increased. Further, the osteoclastic differentiation of bone marrow-derived macrophage/monocyte precursor cells (BMMs) *in vitro* was stimulated by ethanol. The increased osteoclastogenesis of BMMs was associated with increased levels of RANK, PU.1 and MITF expression, activated extracellular signal-regulated kinase (ERK), and reactive oxygen species (ROS). Higher lipid peroxide levels and lower glutathione levels were also observed in the serum of the ethanol-fed rats. These results suggested that ethanol promoted osteoclastogenesis by increasing RANK expression through increases in the production of ROS, activation of ERK and expression of PU.1 and MITF.

Introduction

Alcohol-related damage to bone eventually results in osteopenia, a disease causing substantial morbidity and mortality (1). It is now well accepted that many disorders of bone, including osteopenia and osteoporosis, reflect an imbalance in the differentiation and function of two cell types, the osteoblast and osteoclast, which are responsible for bone formation and bone resorption, respectively (2,3). This study aimed to clarify the effects of ethanol on the function and differentiation of osteoblasts and osteoclasts.

Ethanol dose-dependently reduces cell proliferation and alkaline phosphatase activity in osteoblasts. Moreover, the suppression of osteoblastogenesis is considered a major cause of ethanol-inhibited bone growth, bone loss and deficient bone repair (4). Ethanol has been also demonstrated to stimulate bone resorption (5) and osteoclastogenesis (6).

Osteoclasts, derived from bone marrow hematopoietic stem cells, are highly specialized multinuclear cells capable of resorbing bone (7). The number and activity of osteoclasts are determined by cell lineage allocation, the proliferation and differentiation of osteoclast precursors and the resorptive efficacy of mature osteoclasts (8). Osteoclastic differentiation requires macrophage colony-stimulating factor (M-CSF) and receptor for activation of NF- κ B ligand (RANKL) (9). The binding of M-CSF to c-fms stimulates the expression of receptor for activation of NF- κ B (RANK), the receptor of RANKL, in the hematopoietic osteoclast precursor cells. The binding of RANKL to RANK activates NF- κ B and activator protein-1 (AP-1) and induces osteoclastic differentiation. Recently, we have reported that the suppression of RANK expression was associated with decreased levels of reactive oxygen species (ROS) (10,11). Ethanol was reported to produce ROS in liver Kupffer cells and stellate cells (12), in the lung (13) and in osteoblasts (14). In osteoblasts, ethanol-generated ROS increased the expression of RANKL (14). We hypothesize that ethanol stimulates osteoclastogenesis by increasing RANK expression through direct action on osteoclast precursor cells in addition to the increased RANKL production in osteoblast/stromal cells.

In this study, we investigated the effects of ethanol on bone metabolism and on the gene expression involved in osteoclastogenesis *in vivo* and *in vitro*. Ethanol increased the activities and number of osteoclasts with a normal activity and number of osteoblasts. The increase in the number of osteoclasts was caused by the stimulation of osteoclastogenesis associated

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Abbreviations: ALP, alkaline phosphatase; BMMs, bone marrow-derived macrophage/monocyte precursor cells; CFU-M, colony-forming unit-macrophage; GSH, glutathione; RANK, receptor for activation of NF- κ B; RANKL, receptor for activation of NF- κ B ligand; ROS, reactive oxygen species; TBARS, thiobarbituric acid-reactive substances; TRAP, tartrate-resistant acid phosphatase

Key words: ethanol, osteoclastogenesis, receptor for activation of NF- κ B, PU.1, microphthalmia-associated transcription factor

Table I. Primers used for quantitative real-time PCR.

Target	Forward primer sequence	Reverse primer sequence
Actin	AGCCATGTACGTAGCCATCCA	TCTCCGGAGTCCATCACAATG
cathepsin K	TGTCTGAGAACTATGGCTGTGG	ATACGGGTAACGTCTTCAGAG
c-fm	TAGAGCCAGGTGCAACAGTG	CGCATAGGGTCTTCAAGCTC
c-fos	CTTCACCCCTGCCTCTTCTCA	TCAAAGGGTTCAGCCTTCAG
c-jun	TGAAGCAGAGCATGACCTTG	TAGTGGTGATGTGCCCATTG
M-CSF	CATCCAGGCAGAGACTGACA	TTCGCGCAGTGATAGTGAAC
MITF	TTGGAAGACATCCTGATGGAC	GCTGCTTGTTTTCGAAGCTC
OPG	GAGTGTGCGAATGTGAGGAA	TGCTTTTCGATGACGTCTCAC
PU.1	TGGAGAAGCTGATGGCTTG	CCTTGTGCTTGGACGAGAA
RANK	ATATGCCTGCATCCCCTGAA	TAGCCATCCGTTGAGTTGGA
RANKL	AGCGCAGATGGATCCTAACA	TCGAGTCCTGCAAACCTGTA
TRAP	CAGCCTTATTACCGTTTGC	GAATTGCCACACAGCATCAC

with the increased expression of RANK, PU.1 and microphthemia-associated transcription factor (MITF) in osteoclast precursors and RANKL in osteoblasts/stromal cells.

Materials and methods

Animals and diets. Five-week-old male rats of the Wistar strain were purchased from Japan SLC, Inc. (Shizuoka, Japan) and housed individually in a temperature-controlled room with a 12-h light cycle. After a 1 week period of adaptation, the animals were randomly divided into 2 groups ($n=8$ for each group); a control group and an ethanol-fed group. The ethanol-fed rats were fed a Lieber-DeCarli ethanol-containing liquid diet with 35% of the calories derived from ethanol (F2LEW; Oriental Yeast Co., Ltd., Kyoto, Japan). The controls were pair-fed an alcohol-free isocaloric diet containing maltose-dextrin (F2LCW; Oriental Yeast Co. Ltd., Kyoto, Japan). After 3 weeks, blood and femoral and tibial bone samples were collected under anaesthesia with sodium pentobarbital after overnight access to feed (non-fasting). Serum samples were used to determine the concentrations of ethanol, lipid peroxide, glutathione (GSH) and osteocalcin. After the removal of muscle and tendons, the tibial bone was used for a biochemical analysis, a histological analysis or preparing bone marrow cells. The bone marrow cells were used for the frequency analysis of the clonogenic precursors of osteoblasts and osteoclasts, and the *in vitro* colony formation assay. Animal experiments were performed in accordance with protocols approved by the Animal Care Research Committee of Nara Women's University.

Biochemical analysis. Serum concentrations of ethanol and osteocalcin were measured using a commercial ethanol diagnostic kit (Roche Diagnostics GmbH, Mannheim, Germany) and a Rat osteocalcin EIA kit (Biomedical Technologies, Inc., Stoughton, MA), respectively. Serum lipid peroxide levels were estimated as thiobarbituric acid-reactive substances (TBARS) calculated as malondialdehyde equivalents (15). Serum GSH levels were determined by the method of Ellman (16). The activities of ALP, TRAP and cathepsin K and the amounts of

Ca and hydroxyproline (Hyp) in the proximal tibia (the quarter from the aspect of the knee of the tibia) were determined as reported (17,18).

Histomorphometry. The tibia was fixed in 4% paraformaldehyde, decalcified in 10% EDTA and embedded in paraffin. Sections (4 μm) were stained for TRAP activity using a leukocyte acid phosphatase kit (387A; Sigma) as described (17). Morphometric measurements of trabecular structure (trabecular bone volume, bone surface, thickness and number), and the number of osteoblasts (cuboidal cells on trabecular surfaces) and osteoclasts (TRAP-stained cells with more than 3 nuclei) were carried out at standardized sites (300x300 μm) under the growth plate in the metaphysis of the proximal tibia (19).

Analysis of the frequency of clonogenic precursors of osteoclasts or osteoblasts in bone marrow. The frequency of precursors of osteoclasts or osteoblasts in bone marrow was assessed based on a limiting dilution assay (20) as described previously (21). Briefly, the bone marrow cells were seeded into 96-well plates at 25, 50, 100 or 200 cells/well and cultured for 5 days in the culture medium [MEM containing 10% FCS, penicillin (100 U/ml), streptomycin (100 $\mu\text{g}/\text{ml}$), 1% non-essential amino acid (NEAA) and 1% sodium pyruvate] with M-CSF (5 ng/ml) and RANKL (5 ng/ml) for the assay of clonogenic osteoclast precursors, and at 3×10^5 , 6×10^5 , 10×10^5 , or 20×10^5 cells/well and cultured for 14 days in α -MEM containing 10% FCS, penicillin (100 U/ml), streptomycin (100 $\mu\text{g}/\text{ml}$), 10 mM β -glycerophosphate, and ascorbic acid phosphate salt (50 $\mu\text{g}/\text{ml}$) for the assay of clonogenic osteoblast precursors. The wells containing TRAP-positive multinucleated cells (MNCs) (≥ 3 nuclei/cell) or ALP-positive cells were counted as osteoclast- or osteoblast-positive after the TRAP or ALP staining using the leukocyte acid phosphatase kit (387A; Sigma) and ALP kit (86R; Sigma), respectively.

In vitro colony formation assay. The numbers of colony-forming unit-macrophages (CFU-M) in bone marrow were determined as described (21). Briefly, bone marrow cells (1×10^4) were cultured in 1 ml of MEM containing 1.2% methylcellulose,

Table II. Effects of ethanol on clinical characteristics.

	Control	Ethanol-fed
Body weight (g)		
Start	164.1±1.4	162.5±0.6
Final	196.0±2.1	200.3±3.6
Food intake (g/day)	46.4 ±0.5	46.2±0.7
Bone length (mm)		
Femur	31.3±0.5	31.2±0.4
Tibia	35.3±0.1	35.0±0.4
Bone weight (g)		
Femur	0.553±0.009	0.524±0.004 ^a
Tibia	0.414±0.004	0.396±0.007 ^a
Proximal tibia	0.183±0.003	0.174±0.003 ^a
Serum		
TBARS (nmol/ml)	1.81±0.21	3.84±0.41 ^a
GSH (μmol/l)	41.42±2.19	35.44±1.04 ^a

Values are means ± SEM for 8 rats. ^aSignificantly different from the control value (P<0.05).

30% FCS, 1% BSA and M-CSF (10 ng/ml) for 7 days, and the colonies (>25 μm in diameter) were counted.

Quantitative real-time RT-PCR. Total-RNA was prepared from the proximal tibia after the washing out of bone marrow cells and homogenizing in the presence of 0.1 M EDTA or from cell lysate using a commercial kit (Sepasol-RNA I Super G; Nacalai Tesque, Inc., Kyoto, Japan). The total-RNA was reverse-transcribed with a first-strand cDNA synthesis kit (Toyobo, Co., Ltd., Tokyo, Japan). Real-time PCR was performed using the cDNA or total-RNA for the negative control, with Thunderbird™ SYBR qPCR Mix (Toyobo, Co., Inc.) and specific primers (Table I) as described (18). Levels of gene expression were determined relative to an internal standard (actin) and expressed relative to the control values.

Western blot analysis. Bone extracts of the proximal tibia (17) or cell lysates were used for the western blot analysis. The protein concentrations were measured using the BCA protein assay kit (Pierce of Thermo Fisher Scientific Inc., Rockford, IL). Equal amounts of protein were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred to membranes. Western blotting and reprobing were performed and the chemiluminescent signals were quantified by a densitometer as reported (21). Antibodies recognizing actin (H-300), RANK (H-300), RANKL (FL-317), osteoprotegerin (OPG) (H-240), PU.1 (H-135), MITF (H-50) and p-ERK (E-4) were purchased from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA). The anti-ERK antibody was obtained from Cell Signaling Technology, Inc. (Hitchin, UK).

Osteoclastic differentiation of the bone marrow-derived macrophage/monocyte precursor cells (BMMs). Bone marrow cells from the femur and tibia were cultured for 16-24 h in the

Table III. Effects of ethanol on bone biochemical markers.

	Control	Ethanol-fed
Proximal tibia		
ALP activity (U/g)	21.12±0.59	21.46±1.66
TRAP activity (U/g)	0.838±0.037	1.266±0.090 ^a
Cathepsin K activity (U/g)	3155.7±150.3	3868.8±203.6 ^a
Ca (mg/g)	88.16±2.56	72.73±2.70 ^a
Hyp (μmol/g)	104.1±2.20	93.2±1.50 ^a
Serum osteocalcin (ng/ml)	79.90±2.74	78.60±0.84

Values are means ± SEM for 8 rats. ^aSignificantly different from the control value (P<0.05).

culture medium and cells at the interface after Ficoll-Paque gradient centrifugation of nonadherent cells were used as BMMs (10). BMMs (1×10⁴ cells/well of a 96-well plate or 1.5×10⁵ cells/35-mm plate) (11) were cultured in the culture medium containing M-CSF (20 ng/ml) and RANKL (10 ng/ml) with or without ethanol. Cultures were maintained with a change of medium every 3 days. After 5 days, cells were used for TRAP staining and the assessment of cell viability with a leukocyte acid phosphatase kit (387A; Sigma) and WST-8 (Cell Counting kit-8; Dojindo Laboratories, Kumamoto, Japan), respectively, as described previously (10).

Determination of ERK activation, levels of RANK, PU.1 and MITF, and production of intracellular reactive oxygen species during osteoclastic differentiation of BMMs. BMMs were pre-cultured in the absence of M-CSF with or without ethanol or DPI (50 μM) for 30 min and stimulated with M-CSF (20 ng/ml). Cells were harvested after stimulation for 5 min or 24 h and levels of phosphorylated ERK or RANK, PU.1 and MITF proteins were determined, respectively. To determine the generation of reactive oxygen species, cells were incubated for 10 min with 2', 7'-dichlorofluorescein diacetate (DCFH-DA) (10 μM) after stimulation for 5 min, and the fluorescence of DCF was detected by fluorescence microscopy as described previously (11).

Statistical analysis. All statistical analyses were performed by Welch's method using the Microsoft Excel data analysis program. The differences were considered significant at P<0.05. All data are expressed as the mean ± SEM.

Results

Effects of ethanol on clinical characteristics and bone biochemical markers. The ethanol-fed rats consumed an average of 2.3±0.03 g (mean ± SEM) of ethanol/day. Plasma ethanol concentrations were 111.9±10.6 mg/dl, ranging from 80 to 142 mg/dl. Body weight, food intake and bone length were similar to those in the control group, but the weights of the femur and tibia were significantly lower (Table II). The serum levels of TBARS and GSH were higher and lower than the control values, respectively.

Table IV. Bone histomorphometry.

	Control	Ethanol-fed
Trabecular bone volume (%)	36.69±1.49	28.13±1.35 ^a
Trabecular bone surface (mm/mm ²)	24.26±1.07	21.70±0.55 ^a
Trabecular thickness (μm)	33.16±1.35	25.85±0.63 ^a
Trabecular number (N/mm)	9.78±0.75	10.85±0.28
Osteoblast index (N of Ob/mm trabecular bone length)	55.92±0.92	53.23±1.19
Osteoclast index (N of Oc/mm trabecular bone length)	7.09±0.51	13.29±0.55 ^a

N, number; Oc, osteoclast; Ob, osteoblast. Values are the mean ± SEM for 8 rats. ^aSignificantly different from the control value (P<0.05).

Table V. Frequency analysis of osteoclast and osteoblast precursors and CFU-M.

	Control	Ethanol-fed
Osteoclast precursor (cells/100 BMs)	0.175±0.017	0.480±0.028 ^a
CFU-M (cells/10 ⁴ BMs)	68.0±5.9	71.3±6.8
Osteoblast precursor (cells/10 ⁶ BMs)	0.167±0.016	0.183±0.029

BMs, bone marrow cells. Values are the mean ± SEM for 8 rats. ^aSignificantly different from the control value (P<0.05).

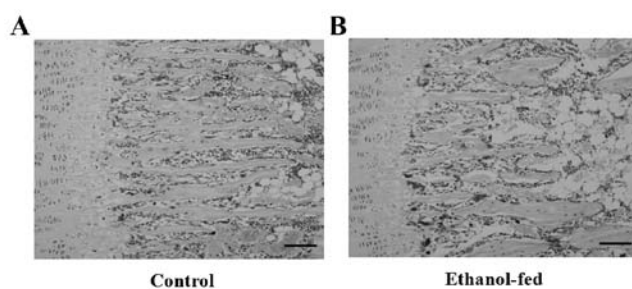


Figure 1. TRAP staining of the proximal tibia in (A) control and (B) ethanol-fed rats. Paraformaldehyde-fixed, decalcified and paraffin-embedded tibia obtained from (A) control and (B) the ethanol-fed rats, was processed for TRAP staining. TRAP-positive cells appeared red. Hematoxylin counterstaining. Magnification x200. Bar, 100 μm. The results presented here are typical of 4 rats of each group.

The ALP activity in the proximal tibia and serum osteocalcin levels were similar to those in the control group (Table III). The activities of TRAP and cathepsin K increased significantly to 1.5- and 1.2-fold the control level, respectively. The levels of hydroxyproline and Ca decreased to about 90% and 80% of the control value, respectively.

Histological analysis. The results of histochemical staining of the tibia for TRAP, a marker of osteoclasts, are shown in Fig. 1. Morphometric measurements showed that the number of osteoclasts in the ethanol-fed rats increased to 1.9-fold the control value (Table IV). The number of osteoblasts was unchanged. Trabecular bone volume, surface and thickness were decreased, but trabecular numbers were not significantly affected.

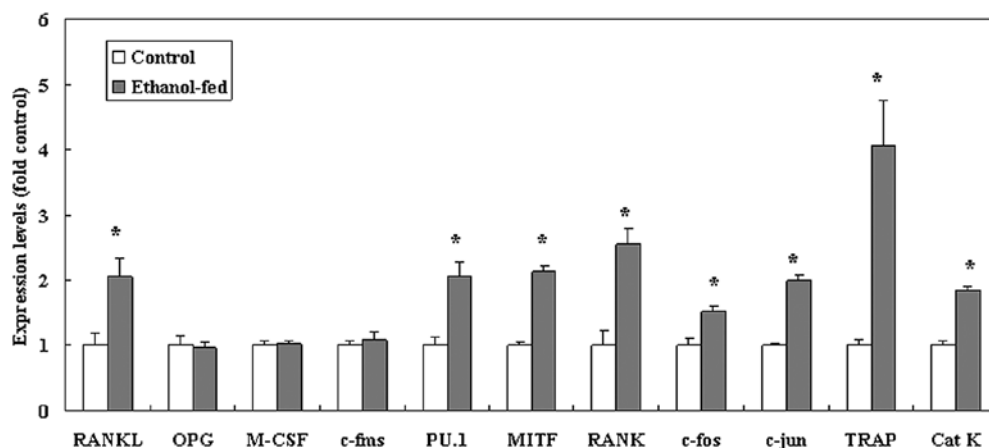


Figure 2. Expression of genes involved in osteoclastic differentiation in the proximal tibia. Total-RNA was extracted from the proximal tibia of control and ethanol-fed rats and the mRNA levels of RANKL, OPG, M-CSF, c-fms, PU.1, MITF, RANK, c-fos, c-jun, TRAP and cathepsin K (Cat K) were assessed by quantitative real-time PCR as described in Materials and methods. Values are the mean ± SEM for 8 rats. Levels are expressed relative to the control value (fold-increase). *Significantly different from the control value (P<0.05).

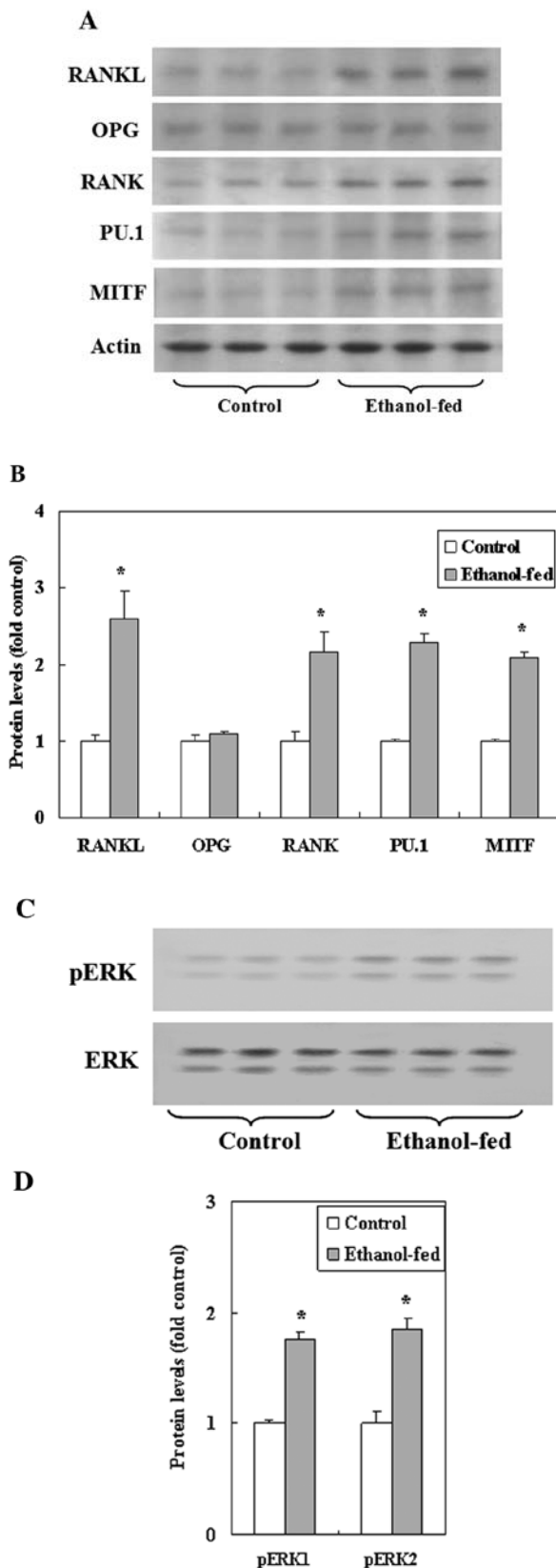


Figure 3. Protein levels of RANKL, OPG, RANK, PU.1, MITF and phosphorylated ERK in the proximal tibia. Bone extracts from the proximal tibia of control and ethanol-fed rats were resolved by SDS-PAGE. After transfer, the blot was probed or re-probed with antibody and detected by ECL as described in Materials and methods. Western blot analyses; (A) RANKL, OPG, RANK, PU.1 and MITF; (C) phosphorylated ERK. The protein levels were quantified by densitometry and represented graphically (B, D). Data represent the mean \pm SEM for 8 rats. *Significantly different from the control value ($P < 0.05$).

Effects of ethanol on the population of osteoclast and osteoblast precursors and CFU-M in bone marrow. Based on the frequency analysis, the population of osteoclast precursor cells in bone marrow (frequency) in the ethanol-fed rats increased to 2.7-fold the control level, although no significant difference was observed in the population of osteoblast precursors (Table V). The numbers of CFU-M in bone marrow were similar among the 2 groups.

Expression of genes involved in osteoclastic differentiation in the proximal tibia. The gene expression levels of the osteoclastogenesis-related factors, RANKL, OPG, M-CSF, c-fms, PU.1, MITF, RANK, c-fos, and c-jun and osteoclast-specific proteins, TRAP and cathepsin K, relative to the internal control, actin, are shown in Fig. 2. The mRNA level of RANKL was about 2-fold the control value (Fig. 2). The expression of OPG, M-CSF and c-fms were not different from the control values. However, levels of RANK, PU.1, MITF, c-fos and c-jun in the ethanol-fed rats were 2.5-, 2-, 2-, 1.5- and 2-fold the control values, respectively. The expression of TRAP and cathepsin K mRNA also increased to 4- and 1.8-fold the control values, respectively.

Protein Levels of RANKL, OPG, RANK, PU.1, MITF and phosphorylated ERK in the proximal tibia. The results of the western blot analysis are shown in Fig. 3. The protein levels of RANKL and RANK in the ethanol-fed group increased to 2.5- and 2-fold the control values, respectively, while the OPG level was similar to the control value (Fig. 3A and B). The PU.1 and MITF protein levels also increased to about 2-fold the control value. The level of phosphorylated ERK1/2 in the EtOH rats was 1.8-fold the control value (Fig. 3C and D).

Effects of ethanol on osteoclastic differentiation and the protein levels of RANK, PU.1, MITF and phosphorylated ERK in BMMs. To clarify the direct effects of ethanol on osteoclast precursor cells, an experiment using BMMs was performed. The addition of ethanol stimulated the osteoclastic differentiation induced by M-CSF and RANKL in BMMs as shown in Fig. 4A. Ethanol at 25 and 50 mM increased the number of TRAP-positive cells to 1.4- and 1.6-fold the control level, respectively (Fig. 4B). Cell viability was not affected by the presence of ethanol (Fig. 4C).

The levels of RANK protein increased to 2.5-fold the control level in the presence of 25 or 50 mM ethanol (Fig. 4D). PU.1 levels increased to 1.6-, 3- and 3-fold on exposure to 10, 25 and 50 mM ethanol, respectively. The levels of MITF increased to 2.5- and 3-fold the control in the presence of 25 and 50 mM ethanol, respectively.

As M-CSF-induced RANK expression was mediated through the activation of ERK (10), the levels of phosphorylated ERK were examined in the presence of ethanol. The phosphorylation of ERK 1/2 increased at 5 min after the addition of M-CSF (Fig. 4E). The presence of 10, 25 and 50 mM of ethanol increased the phosphorylation of ERK1 1.5-, 3- and 3.6-fold, and ERK2 levels 1.6-, 2.6- and 3-fold, respectively, relative to the control value, respectively.

Effects of ethanol on the generation of ROS during the osteoclastogenesis of BMMs. The intracellular production

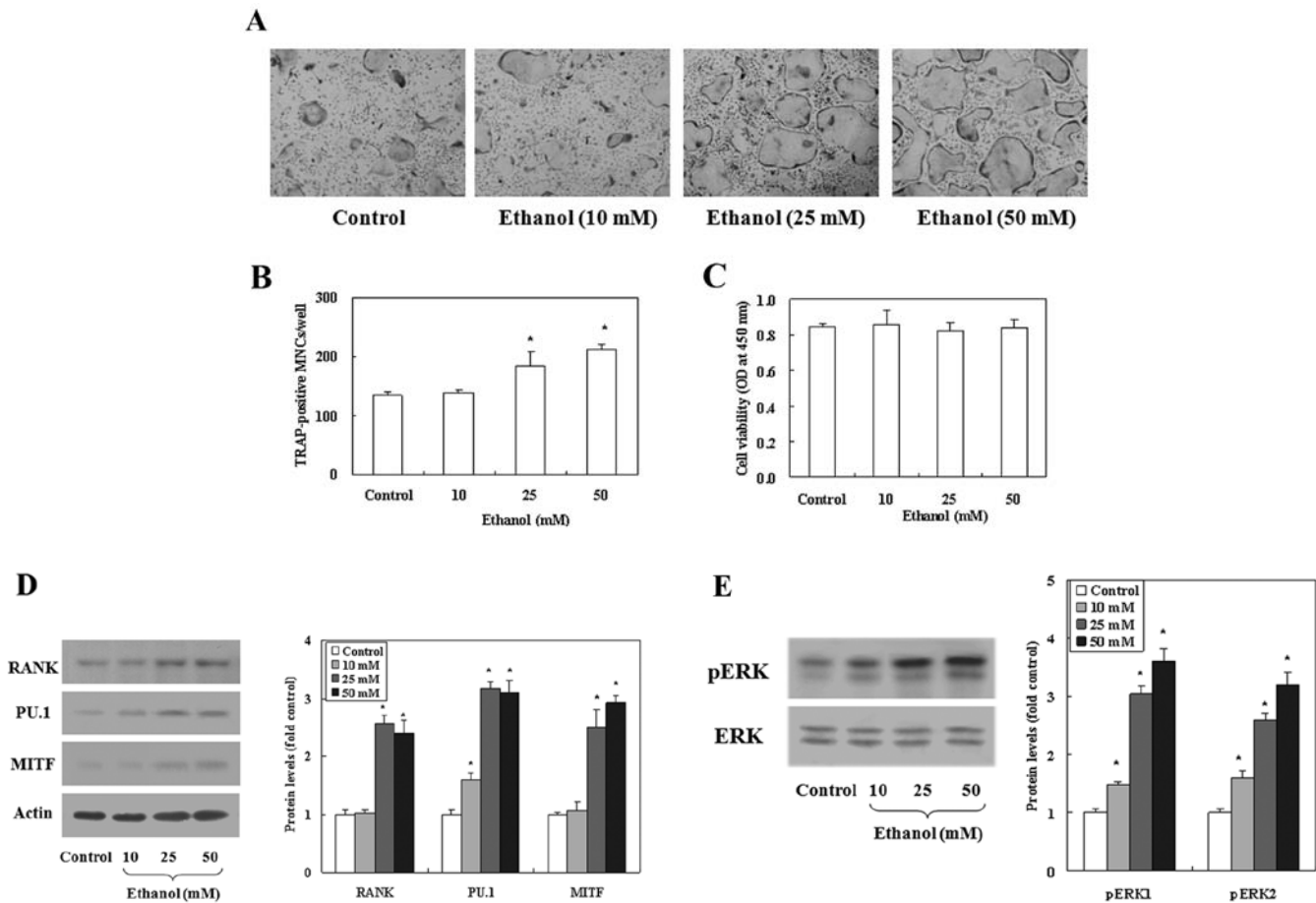


Figure 4. Effects of ethanol on osteoclastic differentiation of BMMs and the protein levels of RANK, PU.1, MITF and phosphorylated ERK. (A-C) BMMs were cultured with M-CSF and RANKL in the absence or presence of ethanol (10, 25 or 50 mM). After the culture of BMMs for 5 days, the cells were used for TRAP staining and the assessment of cell viability as described in Materials and methods. The photographs (x100 magnification) are representative of 6 experiments. (D) After the culture of BMMs for 24 h, the cells were harvested and used for western blotting for RANK, PU.1 or MITF. (E) After the culture of BMMs for 5 min, the cells were harvested and used for western blotting for ERK1/2 or phosphorylated ERK. The protein levels are shown relative to the control value. Values are the mean \pm SEM for 6 experiments. *Significantly different from the control value ($P < 0.05$).

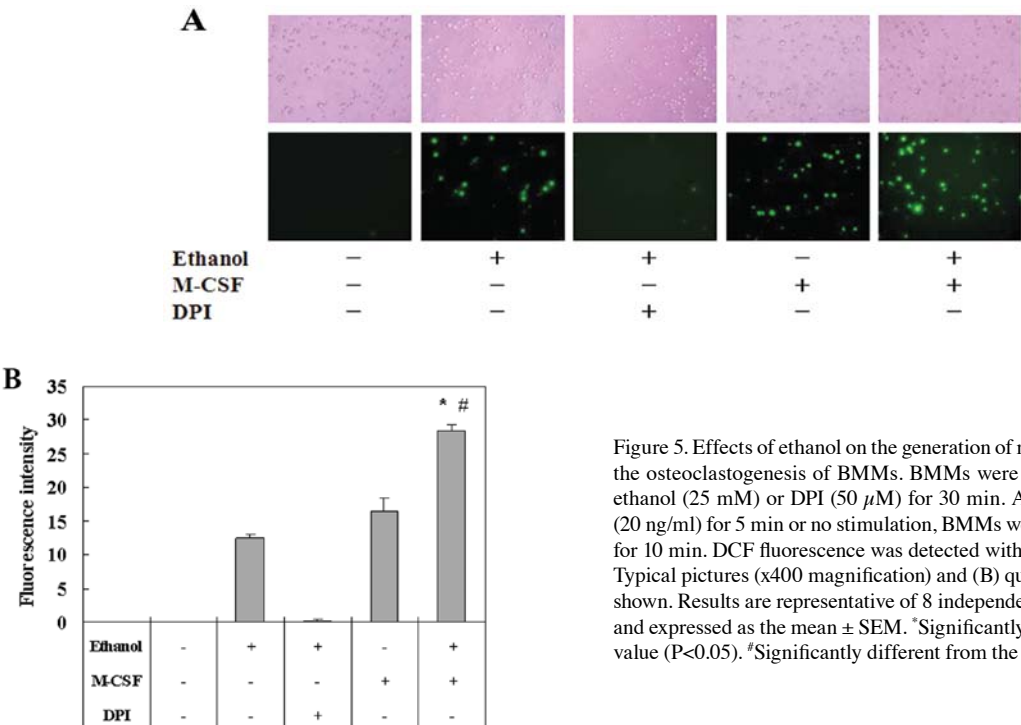


Figure 5. Effects of ethanol on the generation of reactive oxygen species during the osteoclastogenesis of BMMs. BMMs were pre-cultured with or without ethanol (25 mM) or DPI (50 μ M) for 30 min. After stimulation with M-CSF (20 ng/ml) for 5 min or no stimulation, BMMs were incubated with DCFH-DA for 10 min. DCF fluorescence was detected with fluorescence microscopy. (A) Typical pictures (x400 magnification) and (B) quantitative calculation data are shown. Results are representative of 8 independent sets of similar experiments and expressed as the mean \pm SEM. *Significantly different from the (+M-CSF) value ($P < 0.05$). #Significantly different from the (+Ethanol) value ($P < 0.05$).

of ROS was measured with DCFH. Exposure to 25 mM of ethanol resulted in an increase in the intensity of fluorescence (Fig. 5A). The presence of DPI, a Nox inhibitor, suppressed the increase in fluorescence induced by ethanol. The stimulation of BMMs with M-CSF increased the intensity of DCF fluorescence. Exposure to ethanol combined with the stimulation by M-CSF induced an additive increase in fluorescence intensity (Fig. 5).

Discussion

This study clearly demonstrated that ethanol increased the activity and number of osteoclasts. No significant change in the activity or number of osteoblasts was observed after 3 weeks of ingesting ethanol. This is consistent with a report that osteoblast function in cancellous bone of rapidly growing rats (at 6 weeks or younger) appeared to be unaffected during 2-8 weeks of exposure to ethanol (22). The histological analysis showed that the increase in osteoclastic activity was associated with the number of osteoclasts, although the osteoblast number was unchanged. The frequency assay also showed an increase in osteoclast precursors, but no significant change in osteoblast precursors, in bone marrow of the ethanol-fed rats. These results suggested that ethanol did not affect the numbers of osteoblasts/precursors but increased the numbers of osteoclasts and their precursors. However, a significant difference in CFU-M was not observed between the two groups. In bone marrow, the phenotype of progenitors for osteoclasts was comparable with that of CFU-M (23-25). These results suggested that ethanol had no effect on the population of precursors which responded to M-CSF, but reduced that of the precursors which differentiated into osteoclasts in response to RANKL.

The precursors of osteoclasts are derived from the hematopoietic stem cells in bone marrow. The differentiation into osteoclasts, however, occurs on the bone surface *in vivo* (2,24). Therefore, the gene expression of osteoclastogenesis-related factors was examined in the bone. Consistent with the frequency analysis, the mRNA levels of M-CSF and c-fms, the receptor of M-CSF, in the ethanol-fed rats were similar to the control values. However, the mRNA and protein levels of RANKL were significantly increased, although the OPG level was unchanged. RANKL plays a critical role in the differentiation of osteoclast precursors into osteoclasts. Osteoclastic differentiation is principally stimulated by an increase in the biological availability of RANKL, assessed by the ratio of RANKL to its decoy receptor, OPG (26-28). The ratio of RANKL to OPG in the ethanol-fed rats was significantly increased. Consistent with previous reports (6,14), ethanol stimulated osteoclastogenesis through an increase in RANKL expression in osteoblasts. However, in the present study, the mRNA and protein levels of RANK, the receptor of RANKL, also increased. These findings suggested that the increase in osteoclastogenesis caused by ethanol was due to an increase in the expression of RANKL in osteoblasts/stromal cells and also RANK in osteoclast precursors. In this study, for the first time, an increase in the mRNA and protein levels of RANK as well as RANKL due to ethanol was shown. Further, the increase in RANK expression was associated with increases in the mRNA and protein levels of PU.1 and MITF. The gene expression of RANK was shown to be regulated by the tran-

scription factors PU.1 and MITF (29). These results suggested that ethanol increased the expression of RANK through the increase in PU.1 and MITF.

To investigate whether ethanol acted directly on the osteoclast progenitor cells, experiments were performed *in vitro* using BMMs. Ethanol stimulated the osteoclastic differentiation of BMMs and simultaneously increased the expression of RANK, PU.1 and MITF. These results indicated that ethanol acted directly on osteoclast precursors and stimulated osteoclastic differentiation through the increased expression of RANK. Further, the increased expression of RANK was associated with the activation of ERK in agreement with our previous report (10). Indeed, the activation of ERK was also observed here in the proximal tibia of the ethanol-fed rats, in this study, although it is not clear where it occurred, in osteoclasts, osteoblasts or their precursors, *in vivo*. It was reported that ethanol activated ERK in osteoblasts (14). These results suggested that the activation of ERK occurred in osteoclasts/precursors as well as osteoblasts of the ethanol-fed rats.

ROS was reported to play an important role in the activation of ERK in osteoblasts (14,30). In the present study, exposure to ethanol induced the production of ROS in BMMs. The Nox inhibitor DPI inhibited ROS production, indicating the NADPH-dependent generation of ROS in BMMs. Nox has been implicated in the activation of several signaling cascades, including ERK-signaling pathways (31). In fact, ethanol-induced ROS generation via NADPH was also reported in osteoblasts (14). These results suggested that ethanol generated ROS via a NADPH-dependent pathway, activated ERK, and induced the expression of RANK in osteoclast precursors.

Bone formation has long been believed to be the primary target of alcohol's deleterious effects on bone metabolism (1). However, early on, the ethanol acted directly on osteoclast precursors. This study have provided evidence that ethanol stimulates osteoclastogenesis through the increased expression of RANK mediated by the production of ROS and activation of ERK in osteoclast precursor cells in addition to an increase in RANKL in osteoblast stromal cells.

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