

The role of tissue-nonspecific alkaline phosphatase in the phosphate-induced activation of alkaline phosphatase and mineralization in SaOS-2 human osteoblast-like cells

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Abstract Tissue-nonspecific alkaline phosphatase (TNAP) plays a key role in mineralization by degrading inorganic pyrophosphate and providing free inorganic phosphate. We have previously reported that TNAP is induced by β -glycerophosphate and NaH_2PO_4 in short-term cultures of SaOS-2 human osteoblast-like cells and that PHEX (phosphate-regulating gene with homologies to endopeptidase on the X chromosome) mRNA is also induced after TNAP induction. In the present study, we have investigated the effects of levamisole, a TNAP inhibitor, and phosphonoformic acid (PFA), a type III sodium-phosphate cotransporter inhibitor, on the phosphate-induced expression of TNAP and mineralization. Levamisole inhibited β -glycerophosphate-induced mineralization, TNAP and PHEX expression, and the increase in enzymatic activity of NPP1 (5'-nucleotide pyrophosphatase phosphodiesterase 1), but did not inhibit NaH_2PO_4 -induced mineralization. PFA completely inhibited NaH_2PO_4 -induced mineralization and NPP1 enzymatic activation, and partly inhibited β -glycerophosphate-induced mineralization, but did not affect the increase in TNAP activity. These results suggest that phosphate derived from TNAP-induced hydrolysis of β -glycerophosphate yields signals that induce TNAP expression and mineralization, and that PHEX expression may be linked to TNAP expression. However, luciferase assays failed to detect any transcriptional activation of the promoter region of the human TNAP gene by

β -glycerophosphate or NaH_2PO_4 , suggesting that the effects of these phosphates may be indirect.

Keywords Tissue-nonspecific alkaline phosphatase · Mineralization · Levamisole · Phosphonoformic acid · PHEX

Introduction

Alkaline phosphatase (ALP, EC.3.1.3.1) is a biomarker of bone formation. Human enzymes are classified into four types: tissue-nonspecific, intestinal, placental, and germ cell, of which the tissue-nonspecific type is involved in bone formation [1, 2]. Tissue-nonspecific alkaline phosphatase (TNAP) is encoded by a gene located on chromosome 1p36.1-34 and is more than 50 kb in size [3]. TNAP plays a key role in bone mineralization by initiating and/or promoting the formation of hydroxyapatite crystals in the matrix vesicles of osteoblasts and hypertrophic chondrocytes and then propagates them into the extracellular matrix [4, 5]. Matrix vesicles are extracellular membrane-invested particles that serve as the initial site of mineralization [6]. On the outer surface of the matrix vesicle membrane, TNAP degrades inorganic pyrophosphate (PPi) in the extracellular matrix that inhibits hydroxyapatite formation, and also yields free inorganic phosphate (Pi) which reacts with calcium to form hydroxyapatite [4, 6]. PPi in the extracellular matrix is formed by NPP1 (an isoenzyme of 5'-nucleotide pyrophosphatase phosphodiesterase, PC-1) on the matrix vesicle membrane [7] and is also delivered by ANKH (human homolog of mouse progressive ankylosis (Ank)), a transporter of PPi on the hypertrophic chondrocyte membrane [8]. TNAP, NPP1 and ANKH together regulate

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hydroxyapatite formation [9–11]. The role of TNAP in bone mineralization has been confirmed in studies of patients with hypophosphatasia, a genetic defect of the TNAP gene [12–17], and knockout mice [5, 18, 19]. Hypophosphatasia is characterized by hypomineralization of hard tissues, and knockout mice have a phenotype that is similar to that of humans with hypophosphatasia [20]. Patients with hypophosphatasia have elevated serum levels of PPI [20]. Primary osteoblasts obtained from TNAP knockout mice have impaired capacity to undergo mineralization due to an inability to hydrolyze PPI [21]. These findings indicate that TNAP promotes mineralization through hydrolysis of PPI. The TNAP gene is also associated with bone mineral density in elderly women [22]. We have recently demonstrated that TNAP gene expression in SaOS-2 osteoblast-like cells is stimulated by β -glycerophosphate and slightly stimulated by sodium phosphate, both of which have been found to stimulate mineralization in an in vitro cell culture system [23].

Recent studies of phosphate metabolism have identified novel regulators of plasma Pi. In particular, FGF23 (fibroblast growth factor 23) may be a central regulator [24], and PHEX (phosphate-regulating gene with homologies to endopeptidases on the X chromosome) and MEPE (matrix extracellular phosphoglycoprotein) have also been shown to regulate phosphate metabolism and mineralization [25, 26]. The relationship between these factors and TNAP, however, remains unclear. We have previously showed that gene expression of PHEX and MEPE is increased by β -glycerophosphate after induction of the TNAP gene in SaOS-2 cells [23].

Since hydroxyapatite is formed initially within matrix vesicles, intravesicular phosphate ion accumulation may precede calcium phosphate deposition [6]. Extracellular Pi is incorporated into the cells by the type III sodium-phosphate (Na/Pi) cotransporter, which is expressed on hypertrophic chondrocyte membranes [27] and in isolated matrix vesicles from murine chondrogenic cells and plays a role in matrix mineralization [28]. In order to clarify the role of TNAP in mineralization, it is important to understand the relationships among TNAP, the type III Na/Pi cotransporter and mineralization processes. In the present study, we used inhibitors of TNAP and the type III Na/Pi cotransporter to examine the effects of these inhibitors on the expression of TNAP, NPP1, and PHEX in human osteoblast-like cells, SaOS-2. In addition, we evaluated the influence of the inhibitors on mineralization in the cell culture system. Furthermore, we examined the effect of phosphates on transcription activity of the TNAP promoter and TNAP mRNA stability, and examined whether glycerol, a product of β -glycerophosphate hydrolysis, affects the enzymatic activity.

Materials and methods

Cell culture

The human osteoblastic osteosarcoma cell line SaOS-2 was obtained from the RIKEN cell bank (Tsukuba, Japan). Cells were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal calf serum (FCS), 50 U/ml penicillin, and 50 μ g/ml streptomycin in an atmosphere of 5% CO₂. Stock solutions of 1 M β -glycerophosphate disodium and 100 mM sodium dihydrogenphosphate were prepared in DMEM/10% FCS and adjusted to pH 7.3–7.4. In order to examine the effect of the phosphates, β -glycerophosphate and NaH₂PO₄, the stock solutions were added to the culture medium to yield a final concentration of 10 mM. Levamisole, an inhibitor of TNAP, and phosphonoformic acid (PFA), an inhibitor of the type III Na/Pi cotransporter, were also prepared as 100 mM stock solutions and added to the medium to yield a final concentration of 1 mM. Similarly, media containing a series of concentrations of glycerol were prepared to examine the effect of glycerol on ALP and NPP1 activity. In order to assess ALP and NPP1 activity, cells were seeded in 12-well plates and after the designated culture times they were washed with Tris-buffered saline (TBS; 10 mM Tris-HCl and 0.15 M NaCl, pH 7.4). The cells were then harvested in TBS containing 1% (v/v) Triton X-100 and 1 mM phenylmethylsulfonyl fluoride, sonicated, and centrifuged at 10,000 \times g for 10 min. The supernatant was collected and analyzed for the enzyme activity and protein concentration.

Alkaline phosphatase activity

ALP activity was assessed at 37°C using 10 mM p-nitrophenylphosphate as a substrate in 0.1 M 2-amino-2-methyl-1,3-propanediol-HCl buffer (pH 10.0) containing 5 mM MgCl₂. Protein concentration was determined using a Protein DC Assay kit (BioRad Laboratories, Hercules, CA, USA). ALP activity was expressed as μ mol/min/mg protein.

5'-Nucleotide pyrophosphatase phosphodiesterase (NPP1) activity

5'-Nucleotide pyrophosphatase phosphodiesterase 1 (NPP1) activity was assessed at 37°C using 1 mM *p*-nitrophenyl thymidine-5'-monophosphate as a substrate in 50 mM Tris-HCl, pH 8.9. NPP1 activity was expressed as nmol/h/ μ g protein.

Mineralization assay

SaOS-2 cells were cultured with 10 mM β -glycerophosphate or 10 mM NaH₂PO₄ plus 1 mM levamisole or 1 mM

PFA for 5 days, and then mineralization was evaluated using Alizarin Red S staining [29]. Briefly, confluent cells in a 96-well culture plate were rinsed with PBS twice and fixed with ice-cold 70% ethanol for 1 h. The cells were then stained with 0.5% Alizarin Red S for 10 min at room temperature with rotation. After being washed with PBS, photographs of the cells were obtained. Any mineralized materials were solubilized by incubation with 100 mM cetylpyridinium chloride in 10 mM sodium phosphate (pH 7.0) for 1 h at room temperature, and optical density at 570 nm was measured. Mineralization was expressed as U/mg protein, where 1 U was equal to 1.00 unit of OD at 570 nm. Protein concentration was determined in the cells cultured in parallel by using a Protein DC Assay kit. The optical density and the protein concentration were both assayed in triplicate.

Assessment of cell proliferation

After cells were cultured in a 96-well culture plate for 5 days, they were stained with MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrasodium bromide) at 37°C for 3 h, then incubated with 0.04 N HCl in isopropanol for 10 min at room temperature to dissolve the formazan. Cell proliferation was calculated as a percentage as follows: $100 \times (\text{OD at } 595 \text{ nm} / \text{OD at } 595 \text{ nm in control cells})$.

Reverse transcription-polymerase chain reaction (RT-PCR)

Total RNA was extracted from cells using an RNeasy Mini Kit (Qiagen, Hilden, Germany), after which RT-PCR was carried out. Bone-type TNAP, ENPP1 (ecto-nucleotide pyrophosphatase phosphodiesterase, which encodes NPP1), PHEX, MEPE, and ANKH mRNAs were amplified using primers described elsewhere [23]. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA was also amplified as an internal standard. Quantitative real-time RT-PCR was carried out to estimate mRNA levels. In order to estimate the levels of TNAP mRNA, 5 ng of cDNA was transcribed using random hexamers, then primers and a FAM-labeled TaqMan probe described elsewhere [30] were used for amplification. Fluorescence was then detected using an ABI PRISM 7700 Sequence Detector (Applied Biosystems, Foster City, CA, USA). Levels of ENPP1, PHEX and MEPE mRNAs were estimated using the TaqMan Gene Expression Assays (Inventoried) kit with FAM-labeled MGB (minor groove binder) probes. Levels of ENPP1 mRNA were estimated using 10 ng of cDNA with the Assay ID Hs00196445_m1, those of PHEX mRNA were estimated using 20 ng of cDNA with the Assay ID Hs00166525_m1, and those of MEPE mRNA

were estimated using 10 ng of cDNA with Hs00220237_m1. These mRNA levels were normalized according to the levels of GAPDH mRNA, which were estimated using primers and the VIC-labeled TaqMan Probe described elsewhere [30]. For the mRNA stability assay, we used 18S ribosomal RNA (rRNA) as an internal standard.

Luciferase assays

Plasmids containing the TNAP gene promoter region and the pGL3 basic vector (pGL3BV; Promega, Madison, WI, USA) with the firefly luciferase reporter gene were constructed as described elsewhere [30]. SaOS-2 cells were seeded in 6-well plates at a density of 6×10^5 cells/well, and were transfected for 24 h with 4 μg of the luciferase reporter plasmids containing portion of the TNAP promoter region extending from $-4,556$ to -122 or from $-1,313$ to -122 using Lipofectamine 2000 (10 μl /well; Invitrogen, Carlsberg, CA, USA) according to the manufacturer's instructions. Eight nanograms of the pRL-Tk Renilla luciferase reporter plasmid was cotransfected into the cells as a transfection efficiency control. Thereafter, the cells were cultured for 48 h with or without 10 mM β -glycerophosphate or NaH_2PO_4 , harvested, and lysed using the Picca-gene Dual kit (Toyo Ink, Tokyo, Japan). Twenty-microliter aliquots of lysate were added to the Picca-gene substrate solution for dual luciferase assays. Luciferase activity was measured using a Lumat LB 9506 luminometer (EG & G Berthold, Bad Wildbad, Germany) and normalized according to the levels of Renilla luciferase activity.

mRNA stability

SaOS-2 cells were cultured overnight and then treated with 10 mM β -glycerophosphate or 10 mM NaH_2PO_4 for 48 h. In order to assess mRNA stability, a final concentration of 7.5 $\mu\text{g}/\text{ml}$ of actinomycin D was added, cells were harvested at 0, 4, 8, and 24 h, and then total RNA was extracted. The remaining TNAP mRNA levels were determined using quantitative real-time RT-PCR with 18S rRNA as an internal standard. TNAP mRNA levels were normalized according to 18S rRNA levels as estimated using the TaqMan Ribosomal RNA Control Reagents kit (Applied Biosystems). The half-life of the mRNA was calculated from a linear regression of decay of the mRNA levels.

Statistical analysis

All data except for responses to glycerol are expressed as mean \pm SEM, and glycerol response data are expressed as

mean \pm SD. The statistical significance of differences between the groups was estimated using unpaired *t*-test. Values obtained from cells cultured in control medium and those from cells cultured with β -glycerophosphate or NaH_2PO_4 were compared. Similarly, values obtained from cells cultured with β -glycerophosphate or NaH_2PO_4 and those from cells cultured with the inhibitors were compared. For the glycerol response experiment, differences among the groups were analyzed using one-way analysis of variance (ANOVA). A *P* value less than 0.05 denoted the presence of a statistically significant difference.

Results

Effects of inhibitors on mineralization

Figure 1 shows the levels of mineralization in cell cultures incubated with the various phosphates and inhibitors. Addition of β -glycerophosphate resulted in a marked increase in mineralization relative to the control, which was completely inhibited by addition of levamisole, a TNAP inhibitor, and partly inhibited by addition of PFA, a Pi transporter inhibitor. The increase in mineralization relative to the control after addition of inorganic NaH_2PO_4

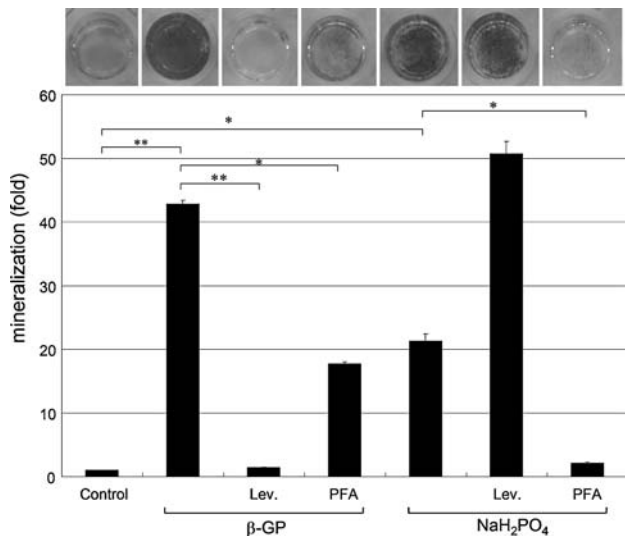


Fig. 1 Effects of the inhibitors levamisole and PFA on mineralization. Mineralization was assessed 5 days after addition of the phosphates (β -glycerophosphate and NaH_2PO_4) and inhibitors by using Alizarin Red S staining. The upper panel shows representative stained cell cultures for each treatment. Values are expressed as a ratio relative to mineralization unit of the cells cultured without β -glycerophosphate or NaH_2PO_4 . Data are expressed as mean \pm SEM for three independent experiments, and each experiment was carried out in triplicate. Differences between the groups were evaluated using unpaired *t*-test. ** *P* < 0.01, * *P* < 0.05. Lev.: levamisole, β -GP: β -glycerophosphate

was smaller than that produced by β -glycerophosphate. This increase was inhibited completely by PFA but not by levamisole. Although addition of levamisole to the medium containing NaH_2PO_4 appeared to increase the levels of mineralization, no statistical difference in the levels of mineralization between addition of NaH_2PO_4 and that of NaH_2PO_4 plus levamisole was exhibited. This seeming increase in the levels of mineralization may be due to low protein concentration that was caused by reduced cell viability as shown in Fig. 2.

Assessment of cell proliferation

Cell viability was evaluated using an MTT assay 5 days after addition of the phosphates and inhibitors. Addition of β -glycerophosphate or NaH_2PO_4 reduced number of viable cells. Furthermore, addition of PFA to the medium containing β -glycerophosphate as well as addition of levamisole to the medium containing NaH_2PO_4 further reduced cell viability (Fig. 2). In particular, addition of levamisole to the medium containing NaH_2PO_4 exhibited low protein concentration.

Effects of inhibitors on TNAP activity and mRNA levels

ALP enzymatic activity was determined at 5 days and TNAP mRNA levels were determined at 3 days after addition of the phosphates and inhibitors. β -Glycerophosphate-induced TNAP mRNA expression and ALP enzymatic activity were inhibited by levamisole but not PFA (Fig. 3). Although levamisole inhibited NaH_2PO_4 -induced TNAP

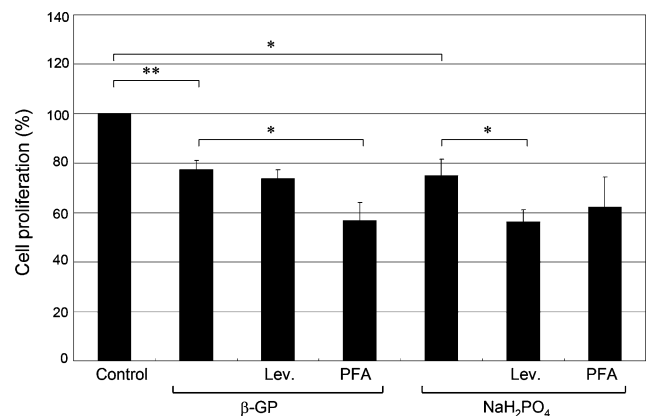


Fig. 2 Assessment of cell proliferation. Cell proliferation was assessed 5 days after addition of the phosphates (β -glycerophosphate and NaH_2PO_4) and inhibitors by using MTT assay. Values are expressed as a percentage described in the Materials and methods section. Data are expressed as mean \pm SEM for four independent experiments, and each experiment was carried out in triplicate. Differences between the groups were evaluated using unpaired *t*-test. ** *P* < 0.01, * *P* < 0.05

mRNA expression, the enzymatic activity did not change. Low protein concentration in levamisole plus NaH_2PO_4 condition may account for this discrepancy.

Effects of inhibitors on NPP1 activity and mRNA levels

NPP1 enzymatic activity was determined at 5 days and ENPP1 mRNA levels were determined at 3 days after addition of the phosphates and inhibitors. Both β -glycerophosphate and inorganic NaH_2PO_4 induced NPP1 enzymatic activity. β -Glycerophosphate-induced NPP1 activity was inhibited by levamisole but not PFA, whereas

NaH_2PO_4 -induced NPP1 activity was inhibited by PFA. However, no statistically significant changes in ENPP1 mRNA levels were observed after addition of the phosphates, suggesting that the changes in NPP1 activity was posttranscriptional (Fig. 4).

Effects of inhibitors on PHEX and MEPE mRNA levels

RT-PCR analysis revealed an apparent increase in PHEX mRNA expression 5 days after the addition of β -glycerophosphate, which seemed to be inhibited by levamisole

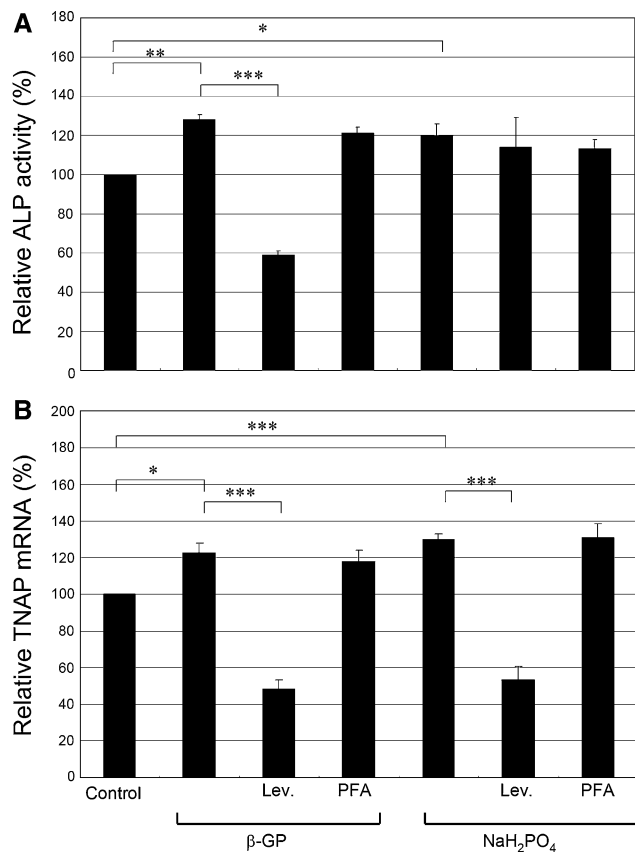


Fig. 3 Effects of the inhibitors levamisole and PFA on expression of TNAP. (a) ALP enzymatic activity was assessed 5 days after addition of the phosphates (β -glycerophosphate and NaH_2PO_4) and inhibitors. Values are expressed as a percentage relative to activity in cells cultured without β -glycerophosphate or NaH_2PO_4 . Data are expressed as mean \pm SEM for four independent experiments, and each experiment was carried out in triplicate. Differences between the groups were evaluated using unpaired *t*-test. *** $P < 0.001$, ** $P < 0.01$, * $P < 0.05$. (b) Levels of TNAP mRNA were assessed 3 days after addition of the phosphates and inhibitors by using a quantitative real-time RT-PCR analysis. Values are expressed as a percentage relative to the levels of TNAP mRNA in cells cultured without β -glycerophosphate or NaH_2PO_4 . Data are expressed as mean \pm SEM for five independent experiments, and each experiment was carried out in duplicate. Differences between the groups were evaluated using unpaired *t*-test. *** $P < 0.001$, * $P < 0.05$

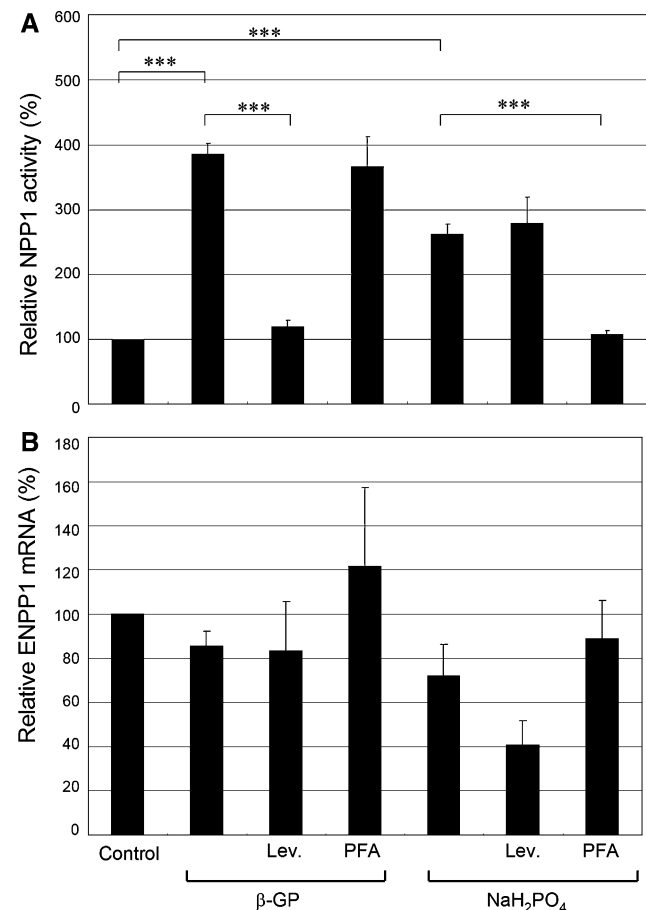


Fig. 4 Effects of the inhibitors levamisole and PFA on expression of NPP1. (a) NPP1 activity was assessed 5 days after addition of the phosphates (β -glycerophosphate and NaH_2PO_4) and inhibitors. Values are expressed as a percentage relative to activity in cells cultured without β -glycerophosphate or NaH_2PO_4 . Data are expressed as mean \pm SEM for four independent experiments, and each experiment was carried out in triplicate. Differences between the groups were evaluated using unpaired *t* test. *** $P < 0.001$. (b) Levels of ENPP1 mRNA were assessed 3 days after addition of the phosphates and inhibitors by using quantitative RT-PCR analysis. Values are expressed as a percentage relative to the levels of ENPP1 mRNA in cells cultured without β -glycerophosphate or NaH_2PO_4 . Data are expressed as mean \pm SEM for three independent experiments, and each experiment was carried out in duplicate. No statistical differences between the groups were observed

(Fig. 5). These findings were confirmed by quantitative RT-PCR analysis (Fig. 6). Levamisole completely inhibited and PFA partly inhibited β -glycerophosphate-induced PHEX expression. MEPE mRNA expression seemed to

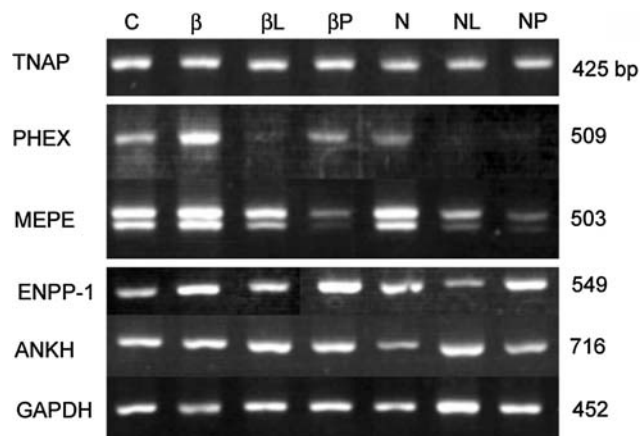


Fig. 5 Effects of the inhibitors levamisole and PFA on expression of the phosphate regulating genes mRNA (RT-PCR). Total RNA was extracted 5 days after addition of the phosphates (β -glycerophosphate and NaH_2PO_4) and inhibitors, and then RT-PCR was carried out using the primers described in the Materials and methods section. The PCR products were analyzed on a 1.5% agarose gel. C: control; β : β -glycerophosphate; βL : β -glycerophosphate + levamisole; βP : β -glycerophosphate + PFA; N: NaH_2PO_4 ; NL: NaH_2PO_4 + levamisole; NP: NaH_2PO_4 + PFA

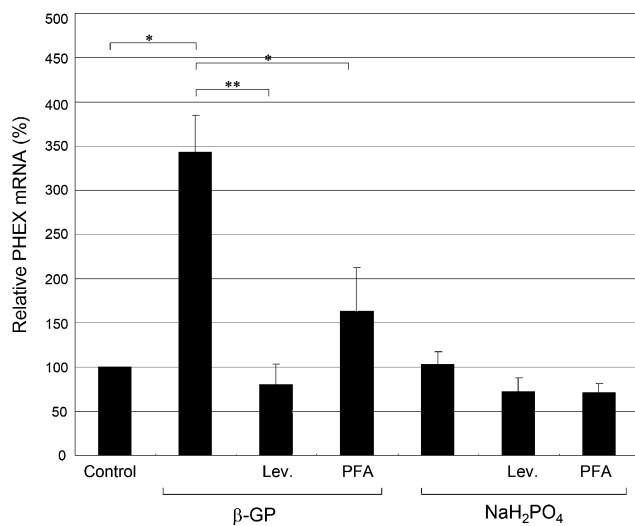


Fig. 6 Effects of the inhibitors levamisole and PFA on expression of PHEX mRNA (quantitative RT-PCR). Quantitative real-time RT-PCR was performed 5 days after addition of the phosphates (β -glycerophosphate and NaH_2PO_4) and inhibitors. PHEX mRNA levels are expressed as a percentage relative to the levels of PHEX mRNA in cells cultured without β -glycerophosphate or NaH_2PO_4 . The PHEX mRNA level was normalized according to the level of GAPDH mRNA. Data are expressed as mean \pm SEM for three independent experiments, and each experiment was carried out in duplicate. Differences between the groups were evaluated using unpaired *t*-test. ** $P < 0.01$, * $P < 0.05$

increase with addition of NaH_2PO_4 , but the increase could not be confirmed by quantitative RT-PCR analysis (data not shown). Although quantitative RT-PCR analysis of TNAP mRNA carried out at 3 days after addition of various phosphates and inhibitors showed changes by phosphates and inhibitors (Fig. 3b), TNAP expression in Fig. 5 appeared not to change. In explanation of this data, intensity of the RT-PCR bands obtained from cells in control medium and that from cells treated with phosphates is almost similar at 5 days after addition of phosphates [23].

Luciferase assay

The dual luciferase assay showed that transcription activity over at least 4.5 kb of the region upstream of the human TNAP gene was not increased by addition of either β -glycerophosphate or NaH_2PO_4 (Fig. 7), suggesting that the phosphates do not directly transactivate the promoter region of the TNAP gene.

mRNA stability

Decay curves of the TNAP mRNA levels after addition of actinomycin D revealed that neither β -glycerophosphate

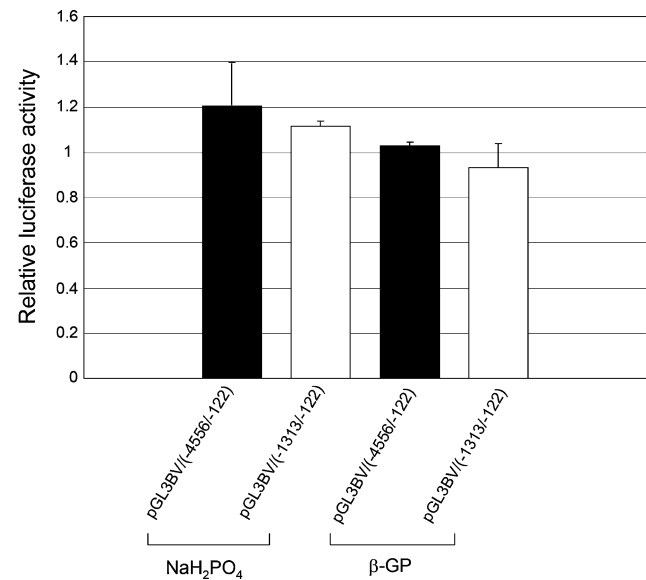


Fig. 7 Effects of the phosphates β -glycerophosphate and NaH_2PO_4 on relative luciferase activity driven by the promoter region of the TNAP gene. Two reporter plasmids containing regions $-4,556$ to -122 and $-1,313$ to -122 of the TNAP promoter region were transfected for 24 h into SaOS-2 cells, then the cells were cultured for 48 h with β -glycerophosphate or NaH_2PO_4 . Luciferase activity was measured 48 h after addition of the phosphates. Luciferase activity is expressed as a ratio relative to activity in the lysates of cells cultured without β -glycerophosphate or NaH_2PO_4

nor NaH_2PO_4 had any effect on TNAP mRNA stability (data not shown).

Effects of glycerol on ALP and NPP1 activity

SaOS-2 cells were cultured for 5 days with 0.5–30 mM glycerol to evaluate the influence of glycerol on ALP and NPP1 activity. Glycerol did not induce ALP and NPP1 enzymatic activity (Fig. 8).

Discussion

In the present study, we showed that a TNAP inhibitor, levamisole, inhibited β -glycerophosphate-induced mineralization and TNAP expression, whereas a Pi transporter inhibitor, PFA, inhibited Pi-induced mineralization but not TNAP expression. Induction of PHEX by β -glycerophosphate was inhibited by levamisole. These results suggest that the products of the extracellular TNAP-induced hydrolysis of β -glycerophosphate induced TNAP expression

and mineralization and that PHEX expression may be linked to TNAP expression. Since one of the products of TNAP-induced hydrolysis of β -glycerophosphate, glycerol, was found to not affect TNAP expression, the factor that induced TNAP expression and mineralization is phosphate. However, phosphates did not directly transactivate the promoter region of the TNAP gene and did not affect TNAP mRNA stability.

We used human osteoblast-like cells, SaOS-2, derived from osteosarcoma in this study. SaOS-2 cells have several osteoblastic properties such as elevated ALP activity, release of matrix-vesicle-like structures with the capacity to mineralize, and production of mineralized matrix in chambers implanted in nude mice [31, 32], suggesting that they mimic primary osteoblast to a certain extent. SaOS-2 cells show extremely high ALP enzymatic activity without abnormal TNAP gene structure [33], and deposit around them an extensive collagenous matrix that is able to mineralize in the presence of an exogenous phosphate donor, β -glycerophosphate [34]. We previously demonstrated that SaOS-2 cells show either β -glycerophosphate-induced or retinoic acid-induced TNAP expression [23, 30]. β -Glycerophosphate was used as an artificial substrate for TNAP in the present study as well as in many of the in vitro studies, which increases the local phosphate concentration. When β -glycerophosphate supplements are kept at levels that produce Pi concentration in agreement with physiologic levels, its use may reflect mineralization per se rather than ALP expression [35]. β -Glycerophosphate and elevated levels of Pi are required for the initiation of mineralization, but once the process is initiated, mineralization will continue at non-elevated levels of phosphates [36]. A study using Fourier transform infrared imaging spectroscopy revealed that hydrolysis of AMP or β -glycerophosphate yields Pi which leads to the formation of mature crystalline, apatite mineral, while the hydrolysis of ATP inhibits the formation of mature hydroxyapatite [37]. As regards concentration of phosphates, we used equimolar concentration of β -glycerophosphate and inorganic NaH_2PO_4 . In a study using mouse MC3T3-E1 cells, supplementation of 10 mM β -glycerophosphate yields 4.8 mM Pi after 48 h culture [38], suggesting that degraded products of β -glycerophosphate stimulate TNAP expression during the time lag and that actual concentration of β -glycerophosphate and Pi may not be equivalent in a cell culture system. Accordingly, further analysis of quantitative comparison of effects caused by β -glycerophosphate with those caused by NaH_2PO_4 may be necessary.

We have previously demonstrated that β -glycerophosphate and NaH_2PO_4 induced TNAP mRNA expression and enzymatic activity [23]. In the present study, we confirmed these results and exhibited that β -glycerophosphate and NaH_2PO_4 reduced cell viability (Fig. 2). Inhibition of cell

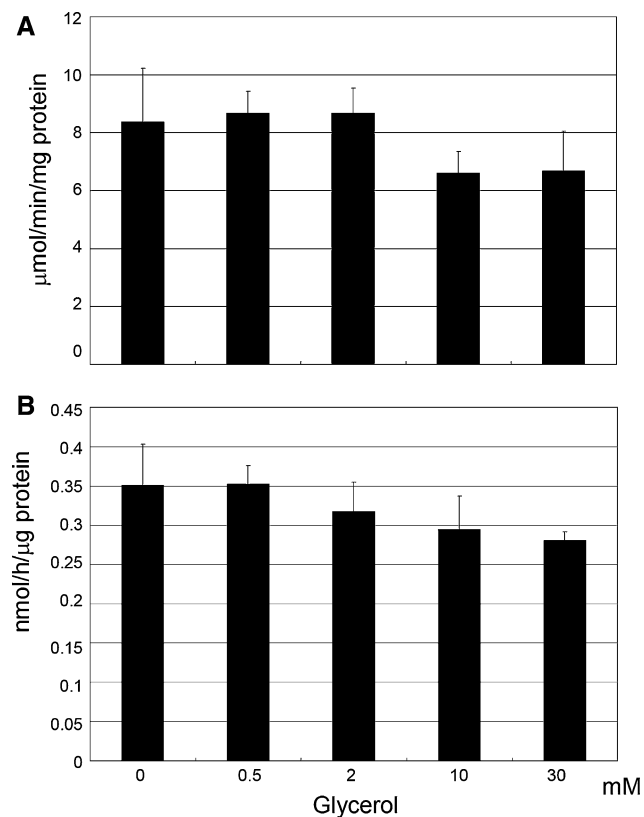


Fig. 8 Effects of glycerol on ALP and NPP1 activity. SaOS-2 cells were cultured for 5 days with 0.5–30 mM glycerol to evaluate the influence of glycerol on ALP (a) and NPP1 (b) activity. Data are expressed as mean \pm SD for an experiment carried out in triplicate. Differences among the groups were evaluated using one way ANOVA. No statistical differences among the groups were observed

proliferation occurs by apoptosis due to incorporation of Pi into the cells through Pi transporters in osteoblast-like cells [39]. In addition, PFA or levamisole further contributed to reduction of cell viability when they were added to the media containing β -glycerophosphate or NaH_2PO_4 , respectively. Levamisole reduces thymidine incorporation into DNA of chick osteoblastic cells [40]. Inhibition of TNAP by levamisole resulted in complete inhibition of β -glycerophosphate-induced mineralization (Fig. 1), consistent with the previous report that levamisole and PPI inhibit β -glycerophosphate-induced mineralization [40]. Furthermore, β -glycerophosphate-induced ALP mRNA expression was inhibited by levamisole, in parallel with inhibition of mineralization and TNAP enzymatic activity (Fig. 3), which indicates that levamisole acts at the transcription step. Levamisole also inhibits expression of the type X collagen $\alpha 1$ chain mRNA in the chick chondrocytes, though the mechanism is not clearly defined [41]. Nevertheless, because levamisole is a low-molecular-weight uncompetitive inhibitor of TNAP enzymatic activity [42], the products of TNAP-induced hydrolysis of β -glycerophosphate may induce transcription of TNAP, given that inhibition of TNAP activity caused inhibition of TNAP mRNA expression. However, although levamisole inhibited NaH_2PO_4 -induced TNAP mRNA, TNAP enzymatic activity of the cells cultured with NaH_2PO_4 and levamisole was not affected. This inconsistency may be caused by seeming high specific activity due to low protein concentration in NaH_2PO_4 plus levamisole that may exert slightly toxic effect on the cells.

PFA inhibited both β -glycerophosphate-induced and NaH_2PO_4 -induced mineralization, but did not affect TNAP expression (Figs. 1 and 3). PFA is a competitive inhibitor of the type III Na/Pi cotransporter that does not inhibit ALP activity [43]. These results indicate that NaH_2PO_4 -induced mineralization requires an influx of Pi through the type III Na/Pi cotransporter. The type III Na/Pi cotransporter is a widely expressed transmembrane protein and transports Pi into cells [27]. SaOS-2 cells express the type III Na/Pi cotransporter, and an osteotropic factor, insulin-like growth factor (IGF) I, stimulates expression of the Na/Pi cotransporter [44]. The type III Na/Pi cotransporter expressed in isolated matrix vesicles plays a role in matrix mineralization of ATDC5 chondrogenic cells [28], and a study using PFA showed that the type III Na/Pi cotransporter also plays an important role in bone mineralization in rat [45]. It is likely that β -glycerophosphate-induced mineralization requires both TNAP activity and an influx of Pi through the Pi transporter.

Levamisole inhibited β -glycerophosphate-induced NPP1 activity and PFA inhibited NaH_2PO_4 -induced NPP1 activity (Fig. 4). NPP1 forms PPI from ATP, which inhibits hydroxyapatite formation [9, 40]. ENPP1 mRNA

expression, however, was neither enhanced by addition of β -glycerophosphate or NaH_2PO_4 nor inhibited by levamisole or PFA, suggesting that the changes in NPP1 activity were not caused by changes in transcription, but were posttranscriptionally responded to the changes in TNAP activity and Pi influx. Addition of exogenous TNAP and transfection of TNAP cDNA into mouse osteoblastic cells increase ENPP1 expression and NPP1 activity accompanied by enhancement of mineralization [46, 47]. The increase in NPP1 activity when β -glycerophosphate was added (Fig. 4a) may be a response to the increase in TNAP activity, which may account for why mineralization proceeded under this condition. Although regulation of NPP1 by Pi has not been investigated, inhibition of Pi influx by PFA resulted in the decrease in NPP1 activity.

PHEX is a transmembrane endopeptidase that is predominantly expressed in bone and teeth [48]. It increases Pi reabsorption in the renal proximal tubules by clearance of a phosphaturic factor or activation of a phosphate-conserving factor, resulting in retention of Pi [48]. Mutations in the human PHEX gene cause X-linked hypophosphatemic rickets, which is characterized by hypophosphatemia and hypomineralization of bone [49]. β -Glycerophosphate upregulates the PHEX gene in primary mouse osteoblasts [50] and in SaOS-2 cells after induction of TNAP [23]. In the present study, levamisole completely inhibited and PFA partly inhibited β -glycerophosphate-induced PHEX expression (Fig. 6). This pattern of inhibition is similar to that observed for inhibition of mineralization, that is, levamisole completely inhibited and PFA partly inhibited mineralization. As regards the relationships between PHEX and mineralization, although the PHEX induction and the increased mineralization do not seem to be related directly each other in the *in vitro* experiment, patients with X-linked hypophosphatemic rickets show skeletal hypomineralization [49]. In addition, U₂OS osteoblast-like cells transfected with active TNAP cDNA exhibit increased levels of PHEX mRNA accompanied by mineralization when cultured with β -glycerophosphate, whereas U₂OS cells transfected with mutant TNAP cDNA do not respond to β -glycerophosphate [51]. Therefore, although the mechanism by which this occurs is unknown at present, TNAP may be involved in PHEX induction. In contrast, NaH_2PO_4 did not affect PHEX expression.

The mechanism by which phosphates induce TNAP is unknown. A luciferase assay using reporter plasmids containing the promoter region of the human TNAP gene showed no direct transactivation of the gene by the phosphates (Fig. 7). Thus, the induction mechanism may be indirect or a region responsive to the phosphates may be situated in a region far from the coding region. We identified a retinoic acid response element in the promoter region of the human TNAP gene [30]. In contrast, we

found that there was no direct transactivation of the TNAP gene by the active form of vitamin D [52]. In fact, vitamin D modulated TNAP gene expression posttranscriptionally by increasing mRNA stability [52]. In the present study, therefore, we determined whether addition of the phosphates altered mRNA stability, but the phosphates had no effect on mRNA stability.

These results raise the question of what stimulated the expression of TNAP. One of the products of TNAP-induced hydrolysis of β -glycerophosphate is glycerol; however, we found that glycerol has no effect on ALP and NPP1 activity (Fig. 8). Glycerol induces osmotic stress [53] but has not been shown to be involved in bone metabolism. Thus, expression of TNAP is induced by signals obtaining from the phosphates derived from degraded β -glycerophosphate and transported by the Na/Pi cotransporter. Pi is a specific signal for induction of osteopontin gene expression in mouse MC3T3-E1 cells through a member of the MAPK (mitogen activated protein kinases) ERK1/2 (extracellular signal-regulated kinase), protein kinase C, and proteasome [54]. Pi also stimulates matrix Gla protein expression in mouse chondrocytes through ERK1/2 signaling pathway [55]. Further analysis whether MAPK and other kinase cascades involve in phosphate-stimulated TNAP expression is needed.

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