RESEARCH ARTICLE



Idelalisib inhibits osteoclast differentiation and pre-osteoclast migration by blocking the PI3K δ -Akt-c-Fos/NFATc1 signaling cascade

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Abstract Since increased number of osteoclasts could lead to impaired bone structure and low bone mass, which are common characteristics of bone disorders including osteoporosis, the pharmacological inhibition of osteoclast differentiation is one of therapeutic strategies for preventing and/or treating bone disorders and related facture. However, little data are available regarding the functional relevance of phosphoinositide 3-kinase (PI3K) isoforms in the osteoclast differentiation process. To elucidate the functional involvement of PI3K\delta in osteoclastogenesis, here we investigated how osteoclast differentiation was influenced by idelalisib (also called CAL-101), which is p1108-selective inhibitor approved for the treatment of specific human B cell malignancies. Here, we found that receptor activator of nuclear factor kappa B ligand (RANKL) induced PI3K\delta protein expression, and idelalisib

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inhibited RANKL-induced osteoclast differentiation. Next, the inhibitory effect of idelalisib on RANKL-induced activation of the Akt-c-Fos/NFATc1 signaling cascade was confirmed by western blot analysis and real-time PCR. Finally, idelalisib inhibited pre-osteoclast migration in the last stage of osteoclast differentiation through down-regulation of the Akt-c-Fos/NFATc1 signaling cascade. It may be possible to expand the clinical use of idelalisib for controlling osteoclast differentiation. Together, the present results contribute to our understanding of the clinical value of PI3K δ as a druggable target and the efficacy of related therapeutics including osteoclastogenesis.

Keywords Phosphoinositide 3-kinases · Idelalisib · Osteoclast differentiation

Introduction

Bone homeostasis requires a tightly maintained balance between osteoblast-mediated bone formation and osteoclast-mediated bone resorption (Takayanagi 2005). Skeletal disorders, such as osteoporosis, Paget's disease, rheumatoid arthritis, and periodontal disease, are characterized by low bone density, which is mainly caused by excessive activity and/or increased numbers of osteoclasts rather than impaired osteoblastic bone formation (Rodan and Martin 2000; Khosla and Riggs 2005; Manolagas and Parfitt 2010). Therefore, most presently available therapeutics have been developed to mitigate the extent of bone loss and reduce bone loss-related fractures through the inhibition of osteoclast differentiation and/or osteoclastic activity (Marie and Kassem 2011).

Bone resorbing osteoclasts are multinucleated giant cells—also called multinucleated osteoclast cells

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(MNCs)—derived from hematopoietic stem cells. Osteoclast development is a complex multi-step process involving differentiation, migration and fusion, which is triggered by two critical factors: macrophage/monocyte colony-forming factor (M-CSF) and receptor activator of nuclear factor kappa B ligand (RANKL) (Feng 2005). Notably, RANKL induces the expression of osteoclastogenesis-related genes by activating several essential signaling molecules and transcription factors, including c-Fos and nuclear factor of activated T cells, cytoplasmic 1 (NFATc1) (Feng 2005; Takayanagi 2005; Takayanagi 2007a). The resulting MNC formation is essentially responsible for mineralized matrix degradation (Nakamura et al. 2012).

Phosphoinositide 3-kinases (PI3Ks) belong to a large family of lipid signaling kinases that are considered potential drug targets. PI3Ks are divided into three classes (I, II, and III) based on their sequence homology and substrate specificity (Vanhaesebroeck et al. 2001; Cantley 2002; Foster et al. 2003). Moreover, class I PI3Ks are subcategorized into two groups: class IA, which comprises three isoforms, PI3K α (p110 α), PI3K β (p110 β), and PI3K δ $(p110\delta)$; and class IB, which includes only PI3K γ $(p110\gamma)$ (Foster et al. 2003). Class I PI3Ks are implicated in the maintenance of various diseases-including cancer, inflammation, and autoimmunity-prompting pharmaceutical companies to focus on the development of class I PI3K inhibitors (Stark et al. 2015; Vanhaesebroeck et al. 2016). Interestingly, the PI3K δ pathway is overactivated in types of B-cell malignancies (Herman et al. 2010; Ikeda et al. 2010; Lannutti et al. 2011; Meadows et al. 2012; Pauls et al. 2012). These findings have directed research focus towards the development of p1108-selective inhibitors (Fruman and Rommel 2011; Norman 2011).

While several pharmacological studies have reported the functional importance of PI3Ks in the function of mature osteoclasts, relatively little data are available regarding the functional relevance of PI3K isoforms in the process of osteoclast differentiation. Therefore, in our present study, we investigated the effect of idelalisib on osteoclast differentiation with the aim of elucidating the functional involvement of PI3K δ in the osteoclastogenesis.

Materials and methods

Reagents

Mouse soluble RANKL and M-CSF were purchased from R&D Systems (Minneapolis, MN, USA). Penicillin, streptomycin, cell culture medium, and fetal bovine serum (FBS) were purchased from Invitrogen (Thermo Fisher Scientific, Waltham, MA, USA). Idelalisib was purchased from Selleckchem (Houston, TX, USA). Antibodies against c-Fos, NFATc1, actin, PI3K- β and PI3K- δ were from Santa Cruz Biotechnology (Dallas, TX, USA). Antibodies against p-Akt (Ser 473), p-Akt (Thr 308), Akt, p-ERK, ERK, p-p38, p38, p-JNK, JNK, PI3K- α , and PI3K- γ were obtained from Cell Signaling Technology (Danvers, MA, USA).

Osteoclast differentiation

Isolation of bone marrow cells (BMCs) from 5-week-old male ICR mice (Damool Science, Daejeon, Korea) was carried out in strict accordance with the recommendations in the Standard Protocol for Animal Study of Korea Research Institute of Chemical Technology (KRICT; Permit No. 2012-7D-02-01). The protocol (ID No. 7D-M1) was approved by the Institutional Animal Care and Use Committee of KRICT (IACUC-KRICT). All efforts were made to minimize suffering, animal number, and stress/ discomfort. Mice were euthanized by cervical dislocation, and then BMCs were obtained by flushing isolated femurs and tibias with α -MEM supplemented with antibiotics (100 units/mL penicillin and 100 µg/mL streptomycin). BMCs were cultured for 1 day on a culture dish in α -MEM supplemented with 10% FBS and 10 ng/mL of M-CSF. Nonadherent BMCs were plated on a Petri dish and cultured in humidified 5% CO₂ at 37 °C for 3 days in the presence of M-CSF (30 ng/mL). After non-adherent cells were washed out, adherent cells were used as bone marrow-derived macrophages (BMMs). When BMMs were cultured with M-CSF (30 ng/mL) and RANKL (10 ng/mL) for 3 days, most of cells differentiated into tartrate-resistant acid phosphatase-positive (TRAP⁺)-mononuclear osteoclasts, and TRAP⁺-MNCs generated by the fusion between each mononuclear cells were observed in the differentiation day 4. Therefore, for the complete formation of TRAP⁺-MNCs, BMMs (1 \times 10⁴ cells/well in a 96-well plate or 3 \times 10⁵ cells/well in a 6-well plate) were cultured with M-CSF and RANKL for 4 days, and most of small round mononuclear TRAP⁺-cells generated from BMMs incubated with M-CSF and RANKL for 3 days were considered to preosteoclasts (Takeshita et al. 2000).

TRAP staining and activity assay

Mature osteoclasts were visualized by staining for TRAP, a biomarker of osteoclast differentiation. Briefly, BMMs-derived MNCs were fixed with 3.7% formaldehyde for 5 min, permeabilized with 0.1% Triton X-100 for 5 min, and stained with the Leukocyte Acid Phosphatase Kit 387-A (Sigma-Aldrich, St. Louis, MO, USA). TRAP⁺-MNCs with three or more nuclei were counted as mature osteoclasts. To measure TRAP activity, the permeabilized

cells were incubated with TRAP buffer (100 mM sodium citrate, pH 5.0, 50 mM sodium tartrate) including 3 mM *p*-nitrophenyl phosphate (Sigma-Aldrich) at 37 °C for 5 min. Reaction mixtures were transferred into a new plate containing an equal volume of 0.1 N NaOH, and optical density values were determined at 405 nm in Wallac EnVision microplate reader (PerkinElmer, Turku, Finland).

Cytotoxicity assay

Cytotoxicity was evaluated by quantitatively measuring lactate dehydrogenase (LDH). Briefly, BMMs $(1 \times 10^4$ cells/well) were seeded in a 96-well plate and incubated for 24 h. Then, cells were incubated with idelalisib in the presence of M-CSF (30 ng/mL) for 3 days, and released LDH in culture supernatants was detected using CytoTox 96 Non-Radioactive Cytotoxicity Assay kit (Promega, Madison, WI, USA) according to the manufacturer's protocol. The absorbance was measured at 492 nm using Wallac EnVision microplate reader.

Western blot analysis

Western blot analysis was performed as described previously (Yeon et al. 2014). Briefly, cells were washed, lysed, and centrifuged at $10,000 \times g$ for 15 min. After protein quantification, proteins were denatured, separated on SDS-PAGE gels, and transferred onto PVDF membranes (EMD Millipore, Burlington, MA, USA). After probing with antibody, the membranes were developed using SuperSignal West Femto Maximum Sensitivity Substrate (Pierce, Thermo Fisher Scientific) and visualized with LAS-3000 luminescent image analyzer (Fuji Photo Film Co., Ltd., Tokyo, Japan). Actin was used for the loading control. Densitometric analysis was performed using ImageJ software (https://imagej.nih.gov/ij/download.html).

Real-time PCR

Real-time PCR was performed as described previously (Yeon et al. 2014). Primers were chosen with the online Primer3 design program (Rozen and Skaletsky 2000). The

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primer sets used in this study were listed in Table 1. Briefly, total RNA was isolated with TRIzol reagent (Thermo Fisher Scientific), and the first-strand cDNA was synthesized with the Omniscript RT kit (Qiagen, Germantown, MD, USA) according to the manufacturer's protocol. SYBR green-based QPCR was performed in Stratagene Mx3000P Real-Time PCR system (Thermo Fisher Scientific) by using Brilliant SYBR Green Master Mix (Thermo Fisher Scientific). All reactions were run in triplicate, and data were analyzed by the $2^{-\Delta\Delta CT}$ method (Livak and Schmittgen 2001). Gene encoding hypoxanthine phosphoribosyltransferase 1 (HPRT1) was used as the internal standard, and the statistical significance was determined with *HPRT1*-normalized $2^{-\Delta\Delta CT}$ values.

Cell migration assay

The migratory ability of pre-osteoclasts was measured in Boyden chamber with modifications (Yeon et al. 2014). Briefly, after generating pre-osteoclasts, cells were resuspended with α -MEM medium containing 0.1% FBS, M-CSF (30 ng/mL), RANKL (10 ng/mL), and/or idelalisib. α-MEM medium (30 µL) containing 10% FBS and/or idelalisib was added into the bottom chamber, and after placing over the gelatin-coated membrane filter, the silicone gasket, and the top chamber, cell suspension (2×10^4) cells/50 µL) was added into the top chamber, followed by culture in humidified 5% CO2 at 37 °C for 12 h. Then, cells in the upper surface of the membrane were carefully removed with a cotton swab, and pre-osteoclasts that had migrated across the membrane to the lower surface of the membrane were fixed and stained with Diff-Quik stain kit (Siemens Healthcare, Erlangen, Germany). The number of migrated cells were counted in random areas of membrane.

Statistical analysis

All experiments were performed in triplicate and all quantitative values were presented as mean \pm SD. Statistical differences were analyzed using Student's t-test or ANOVA with post hoc analysis using GraphPad Prism 5

Table 1	Primer sequences used
in this study	

Target gene	Forward Primer $(5'-3')$	Reverse primer $(5'-3')$
NFATc1	GGGTCAGTGTGACCGAAGAT	GGAAGTCAGAAGTGGGTGGA
TRAP	GATGACTTTGCCAGTCAGCA	ACATAGCCCACACCGTTCTC
OSCAR	AGGGAAACCTCATCCGTTTG	GAGCCGGAAATAAGGCACAG
DC-STAMP	CCAAGGAGTCGTCCATGATT	GGCTGCTTTGATCGTTTCTC
ATP6vOd2	AGACCACGGACTATGGCAAC	CGATGGGTGACACTTGGCTA
Cathepsin K	GGCCAACTCAAGAAGAAAAAC	GTGCTTGCTTCCCTTCTGG
HPRT1	TGCTCGAGATGTCATGAAGG	AGAGGTCCTTTTCACCAGCA

(GraphPad Software, San Diego, CA, USA), and a value of P < 0.05 was considered significant.

Results

PI3K isoforms are increased during osteoclast differentiation

To investigate how PI3K isoforms are involved in osteoclast differentiation, we evaluated their protein expression levels during RANKL-induced commitment of BMMs into osteoclasts. As shown in Fig. 1, western blot analysis confirmed expression of all isoforms in BMMs, and revealed that the expression levels of PI3K α , γ , and δ were temporally increased by RANKL treatment. In addition, PI3K α , β , and γ were strongly induced during the late stage of osteoclast differentiation, while PI3K δ was strongly induced in the early stage compared to the other isoforms.

The RANKL-mediated induction of c-Fos has been known to be required for the auto-amplification of NFATc1, enabling the robust induction of NFATc1 during osteoclastogenesis (Asagiri and Takayanagi 2007). To confirm the RANKL-induced commitment of BMMs into osteoclasts, we evaluated transcription factors related to osteoclast differentiation, c-Fos and NFATc1, and found that they were temporally induced by RANKL, as

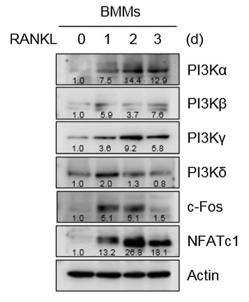


Fig. 1 Increased protein expression patterns of PI3K isoforms during osteoclast differentiation. In the presence of M-CSF (30 ng/mL), BMMs were cultured with RANKL (10 ng/mL) for the indicated number of days, and then the protein expression levels of the PI3K isoforms were evaluated by western blot analysis. The relative ratios of PI3K isoforms/actin, c-Fos/actin and NFATc1/actin were presented. The numbers are the intensity compared with that of the control

expected; RANKL strongly induced the protein expression of c-Fos 1 \sim 2 days after its treatment, and then the sub-sequential induction of NFATc1 was observed.

$\label{eq:pharmacological PI3K\delta inhibition by idelalisib inhibits RANKL-induced osteoclast differentiation$

Idelalisib (Fig. 2a) is one of p110δ-selective inhibitors. Although PI3K α , β , γ , and δ were all induced during RANKL-induced osteoclast differentiation of BMMs, the following experiments mainly focused on the use of idelalisib to investigate the biological relevance of PI3K δ inhibition in osteoclast differentiation. Interestingly, idelalisib dose-dependently inhibited TRAP⁺-MNC formation (Fig. 2b). We confirmed its inhibitory effect on osteoclast differentiation by counting the number of TRAP⁺-MNCs (Fig. 2c) and measuring TRAP activity (Fig. 2d).

We could not exclude the possibility that the RANKLinduced commitment of BMMs into osteoclasts might be inhibited by cytotoxicity of idelalisib towards BMMs. Therefore, we further evaluated idelalisib's effect on BMM survival by measuring the activity of LDH, a stable cytosolic enzyme that is released upon cell lysis. As shown in Fig. 2e, idelalisib exhibited no cytotoxicity in BMMs, indicating that its anti-osteoclastogenic activity was not due to cytotoxicity.

Idelalisib inhibits RANKL-induced activation of the Akt-c-Fos/NFATc1 signaling cascade

To elucidate idelalisib's anti-osteoclastogenic mechanism, we performed western blot analysis to evaluate how idelalisib affected the activation of osteoclastogenesis-related signaling molecules, including Akt, ERK, p38, and JNK. As shown in Fig. 3a, idelalisib attenuated the RANKLinduced phosphorylation of Akt at serine 473, while the other molecules were not changed.

We further examined idelalisib's effects on the RANKL-induced protein expressions of c-Fos and NFATc1, which are master transcriptional regulators of complete osteoclastogenesis. The IC₅₀ of idelalisib on TRAP activity was approximately 3 μ M (Fig. 2d); therefore, the following experiments were performed using 3 μ M idelalisib. In BMMs treated with idelalisib during their differentiation into osteoclasts, we observed inhibition of the RANKL-mediated induction of c-Fos and NFATc1 at 24 and 48 h after RANKL treatment, respectively (Fig. 3b). Real-time PCR analysis confirmed that idelalisib strongly inhibited NFATc1 expression (Fig. 3c).

To additionally confirm that idelalisib inhibited osteoclastogenesis-related transcription factors, we evaluated the mRNA expression levels of the osteoclastogenesis-related genes encoding TRAP, osteoclast-associated receptor

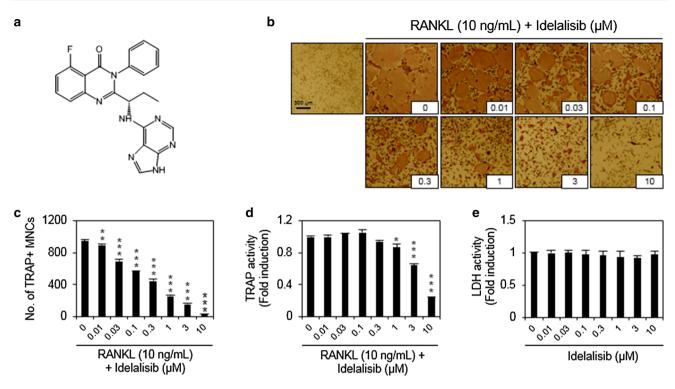


Fig. 2 Idelalisib inhibits the osteoclast differentiation of BMMs. **a** Chemical structures of idelalisib. **b**–**d** In the presence of M-CSF (30 ng/mL), BMMs were pretreated with the vehicle control (0.1% DMSO) or idelalisib for 5 min and then cultured with RANKL (10 ng/mL) for 4 days. Then, the effect of idelalisib on the RANKL-induced differentiation of BMMs into osteoclasts was evaluated by TRAP staining (b; magnification, $100 \times$), counting the TRAP⁺-MNCs with \geq 3 nuclei (**c**), and measuring TRAP activity (**d**). TRAP⁺-MNCs were photographed under a light microscope with $100 \times$ magnification. **e** The cytotoxicity of idelalisib towards BMMs was evaluated by measuring LDH activity. **P* < 0.05; ***P* < 0.01; ****P* < 0.001

(OSCAR), dendritic cell-specific transmembrane protein (DC-STAMP), d2 isoform of vacuolar (H⁺) ATPase V₀ domain (ATP6v0d2), and cathepsin K. As shown in Fig. 3d, idelalisib significantly inhibited the mRNA expression levels of each of these genes. Overall, these results suggested that idelalisib's anti-osteoclastogenic activity could be due to its specific inhibition of PI3K δ , which subsequently downregulates the expression of osteoclastogenesis-related genes by inhibiting the RANKL-induced activation of the Akt-c-Fos/NFATc1 signaling axis.

Idelalisib inhibits the last stage of osteoclast differentiation

To determine what stages of osteoclast differentiation were affected by idelalisib, we reevaluated idelalisib's effects on RANKL-induced osteoclast differentiation by treating cells at eight time-points, (b)-(i), as shown in Fig. 4a. Interestingly, idelalisib strongly inhibited TRAP⁺-MNC formation when BMMs were incubated with idelalisib from differentiation day 3 to 4, (b–e) in Fig. 4b, c. Additionally, idelalisib moderately inhibited TRAP⁺-MNC formation when BMMs were continuously incubated with idelalisib from day 0 to day 3, (i) in Fig. 4b, c. Anti-osteoclastogenic activity of idelalisib was clarified by visualizing the accumulation of mononuclear cells in the microscopic images, (b–e) and (i) in Fig. 4b, as well as measuring the number of TRAP + -MNCs (Fig. 4c). On the other hand, osteoclast differentiation was not impacted by incubation with idelalisib for only 1 day within days 0 to 3, (f–h) in Fig. 4b, c.

Idelalisib inhibits pre-osteoclast migration by blocking activation of the Akt-c-Fos/NFATc1 signaling cascade

At the last stage of osteoclast differentiation, pre-osteoclasts migrate and fuse to each other to generate MNCs. Thus, our results suggested that the pharmacological inhibition of PI3K δ by idelalisib might affect pre-osteoclast migration during osteoclast differentiation. To investigate this possibility, we used the Boyden chamber migration assay to examine the effect of idelalisib on the migration of pre-osteoclasts. As shown in Fig. 5a, b, the migration of pre-osteoclasts was significantly and dose-dependently inhibited by idelalisib.

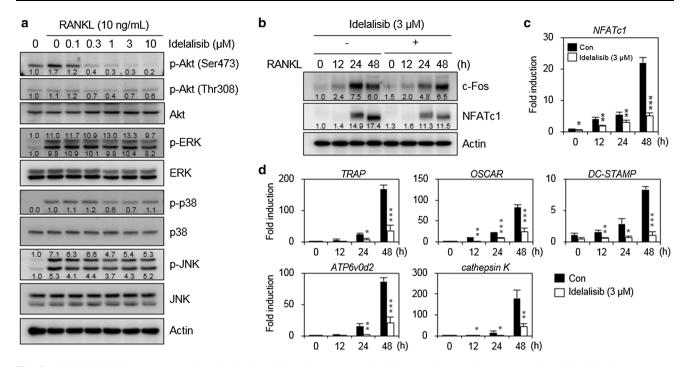


Fig. 3 Idelalisib inhibits RANKL-mediated activation of the Akt-NFATc1 signaling cascade. **a** In the presence of M-CSF (30 ng/mL), BMMs were pretreated with idelalisib for 5 min, and then incubated with RANKL (10 ng/mL) for 5 min. Next, western blot analysis was performed to evaluate the effect of idelalisib on the RANKL-induced activation of signaling molecules. The relative ratios of p-Akt/Akt, p-ERKs/ERK, p-p38/p38 and p-JNKs/JNKs were presented. The numbers are the intensity compared with that of the control. **b–d** In the presence of M-CSF (30 ng/mL), BMMs were pretreated with idelalisib (3 μ M) for 5 min, and then incubated with RANKL (10 ng/mL) for the indicated time. Then, western blot analysis was performed to evaluate the effect of idelalisib on the RANKL-mediated induction of c-Fos and NFATc1. The relative ratios of c-Fos/actin and NFATc1/actin were presented. The numbers are the intensity compared with that of the control (**b**). Real-time PCR analysis was also performed to evaluate the effects of idelalisib on the RANKL-mediated mRNA inductions of NFATc1 (**c**) and osteoclastogenesis-related genes (**d**). **P* < 0.05; ***P* < 0.01; ****P* < 0.001

We additionally performed western blot analysis to confirm the inhibitory effect of idelalisib on downstream molecules of PI3K δ , such as Akt, c-Fos, and NFATc1, in pre-osteoclasts. As shown in Fig. 5c, idelalisib inhibited the RANKL-induced phosphorylation of Akt at serine 473, but not at threonine 308, in pre-osteoclasts. Moreover, idelalisib inhibited the RANKL-mediated induction of c-Fos and NFATc1 protein expression in pre-osteoclasts.

Discussion

Several pharmacological studies have investigated the contributions of specific PI3K isoforms to mature osteoclast's function. Wortmannin is a pan-PI3K inhibitor that shows little selectivity within the PI3K family (IC₅₀:-PI3K α , 1 nM; IC₅₀:PI3K β , 10 nM; IC₅₀:PI3K γ , 9 nM; and IC₅₀:PI3K δ , 5 nM) and reports have described its inhibitory effects on bone resorptive activity, attachment, spreading, and chemotaxis of osteoclasts (Nakamura et al. 1995; Lakkakorpi et al. 1997; Pilkington et al. 1998; Juss et al. 2012). On the other hand, GS-9820 is a selective PI3K δ inhibitor with an IC₅₀:PI3K α of 5.44 µM, IC₅₀:PI3Kβ of 3.38 μM, IC₅₀:PI3Kγ of 1.40 μM, and IC₅₀:PI3Kδ of 12.7 nM. A recent pharmacological inhibition study used wortmannin and GS-9820 to verify that PI3Kδ plays a critical role in regulating the osteoclast cytoskeleton and the resorptive activity of mature osteoclasts (Shugg et al. 2013).

Research groups have also examined how specific PI3K isoforms contribute to osteoclast differentiation by assessing the direct actions of pharmacological inhibitors on osteoclast precursors. Interestingly, one report demonstrated that osteoclastogenesis was significantly decreased by treatment with PI3K α inhibitors, but not with PI3K β inhibitors or PI3Ko inhibitors such as IC87114 (Grey et al. 2010). IC87114 shows an IC_{50} :PI3K α of > 100 μ M, IC₅₀:PI3Kβ of 5 μM, IC₅₀:PI3Kγ of 1 μM, and IC₅₀:PI3Kδ of 100 nM, and is thus certainly more selective for PI3K δ than for other PI3K class I enzymes (Hawkins et al. 2015). However, it is possible that its IC₅₀:PI3Kδ value of 100 nM may be insufficient for it to show specificity to PI3Kδ. Thus, in our present study, we reexamined the functional relevance of PI3K δ in osteoclast differentiation by using idelalisib, which is the only p110 δ -selective inhibitor approved by the FDA and EMA. A previous in vitro cell-

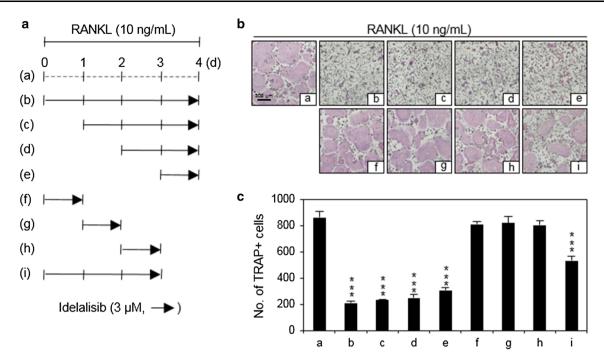


Fig. 4 Idelalisib inhibits the last stage of osteoclast differentiation. **a** Idelalisib exposure schedule. BMMs cultured in the presence of M-CSF (30 ng/mL) and RANKL (10 ng/mL) were treated with idelalisib (3 μ M) or vehicle (0.1% DMSO) for the time periods indicated by black arrows. After BMMs differentiated into osteoclasts, TRAP⁺-MNCs were photographed under a light microscope (b; magnification, 100 ×), and the number of TRAP⁺-MNCs were counted (c). *** *P* < 0.001

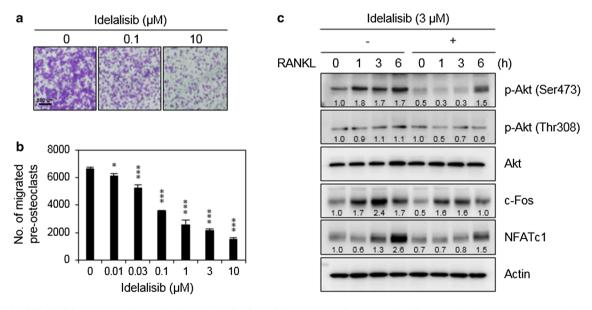


Fig. 5 Idelalisib inhibits pre-osteoclast migration and activation of the Akt-c-Fos/NFATc1 signaling cascade in pre-osteoclasts. **a** The effect of idelalisib on pre-osteoclast migration was evaluated using the Boyden chamber migration assay. The migrated pre-osteoclasts were fixed and stained, and representative images are presented (magnification, $100 \times$). **b** The number of migrated pre-osteoclasts were counted. **P* < 0.05; ****P* < 0.001. **c** Western blot analysis was performed to evaluate the effects of idelalisib on the induction of p-Akt/Akt, c-Fos and NFATc1 in pre-osteoclasts. The relative ratios of p-Akt/Akt, c-Fos/actin, and NFATc1/actin were presented. The numbers are the intensity compared with that of the control

free assay demonstrated that idelalisib was more selective for PI3K δ than for other PI3K class I enzymes (IC₅₀:-PI3 K α , 820 nM; IC₅₀:PI3 K β , 565 nM: IC₅₀:PI3K γ , 89 nM; and IC₅₀:PI3K δ , 2.5 nM) (Lannutti et al. 2011). PI3K δ is predominantly expressed in cells of hematopoietic origin (Kok et al. 2009). Therefore, at the start of this study, we confirmed the expression of PI3K δ protein in the BMMs used as precursor cells of mature osteoclasts. Triggering the commitment of BMMs into osteoclasts with RANKL resulted in increased PI3K δ protein expression, suggesting that this protein might be functionally relevant to osteoclast differentiation. All other PI3K isoforms (α , β , and γ) were also expressed in BMMs, and those proteins were expressed at increased levels following RANKL treatment, suggesting that all PI3K isoforms may play critical roles in osteoclast differentiation. In fact, the evidence showing the involvement of PI3K β to the osteoclast-mediated bone resorption in mice and humans has been reported (Győri et al. 2014), and PI3K/ Akt signaling pathway has been also suggested to play a role in RANKL-independent osteoclastogenesis (Xing et al. 2016), but in our present study, we focused on the functional relevance of PI3K\delta to the RANKL-mediated osteoclast differentiation.

Here, idelalisib significantly and dose-dependently inhibited TRAP⁺-MNC formation, confirming the functional relevance of PI3K δ to osteoclast differentiation. Moreover, idelalisib showed no cytotoxicity towards BMMs, but rather inhibited activation of the RANKLmediated Akt-c-Fos/NFATc1 signaling cascade. This suggested that idelalisib's anti-osteoclastogenic activity might be due to its specific potential to inhibit PI3K δ , leading to subsequent suppression of RANKL-induced activation of the Akt-c-Fos/NFATc1 signaling axis. In addition to Akt, MAP kinases (e.g., ERK, p38, and JNK) have been reported to play roles in the early stage of RANKL-induced osteoclast differentiation by controlling the activity and/or expression of c-Fos and NFATc1 (Lee et al. 2002; Huang et al. 2006; Takayanagi 2007a, b; Yamanaka et al. 2013). However, in this study, idelalisib did not affect the RANKL-induced activation of those three MAP kinases. Our data indicated that the idelalisib-mediated direct inhibition of PI3K\delta dominantly blocked the RANKL-induced phosphorylation of Akt on Ser473, but did not influence MAP kinases in the osteoclastogenesis. It has been previously demonstrated that the RANKL-induced phosphorylation of Akt on Ser473 is PI3K-dependent (Kim et al. 2003).

Akt induces osteoclast differentiation through regulation of the NFATc1 signaling cascade (Moon et al. 2012), and both c-Fos and NFATc1 are well-known transcription factors controlling the expression of osteoclastogenesisrelated genes (Takayanagi 2007a, b). In Akt1 deficiency, osteoclastogenesis is markedly inhibited, with reduced accumulation of specific osteoclast mRNAs and proteins, and impaired fusion to form MNCs (Mukherjee and Rotwein 2012). In the present study, to confirm that idelalisib's anti-osteoclastogenic effect was exerted through inhibiting the actions of c-Fos and NFATc1, we also evaluated the mRNA expression levels of osteoclastogenesis-related genes, including *TRAP*, *OSCAR*, *DC-STAMP*, *ATP6v0d2*, and *cathepsin K* (Song et al. 2009), and found that idelalisib significantly inhibited the RANKL-induced mRNA expression levels of these genes, which have been considered biomarkers for osteoclastogenesis. Notably, both *DC-STAMP* and *ATP6v0d2* contain multiple NFATc1binding sites in their promoter regions, which reportedly play critical roles in pre-osteoclast fusion (Kim et al. 2008; Song et al. 2009). In addition, the products of these genes have been shown to function in the complete fusion of the migrating pre-osteoclasts to form functionally activated MNCs.

Osteoclast precursors migrate to the bone surface and fuse with each other to form fully differentiated and functionally activated MNCs (Kikuta and Ishii 2013). Although this process is incompletely understood, it is considered that targeting cell behavior (e.g., migration) may be a potential therapeutic strategy (Millar et al. 2017). Interestingly, our present results demonstrated that idelalisib inhibited pre-osteoclast migration by blocking activation of the Akt-c-Fos/NFATc1 signaling cascade, suggesting the functional importance of PI3K δ in pre-osteoclast migration during osteoclast differentiation.

The PI3K-Akt pathway has been implicated in osteoclast precursor migration (Munugalavadla et al. 2008; Boudot et al. 2010), and PI3K δ has been reported to play an important role in controlling cell migration via Akt activity in macrophages (Vanhaesebroeck et al. 1999; Papakonstanti et al. 2008). Akt can control the transcriptional activity of NFATc1, which subsequently regulates the expression of fusion-related genes such as *DC-STAMP* and *ATP6v0d2*. Thus, the idelalisib-mediated direct inhibition of the PI3K δ -Akt signaling pathway might decrease the expression of c-Fos and NFATc1, resulting in subsequent downregulation of fusion-related genes. This suggests that the anti-osteoclastic action of idelalisib could phenotypically present in blocked pre-osteoclast migration.

In summary, our results suggest that idelalisib may inhibit pre-osteoclast migration, and that its anti-osteoclastogenic action could result from blockade of the PI3Kδ-Akt-c-Fos/NFATc1 signaling cascade. Idelalisib was recently approved for the treatment of specific human B-cell malignancies, and several clinical trials are currently investigating its possible future use in the treatment of a range of malignancies (Yap et al. 2015). Importantly, micro-osteoclast resorption has been suggested as a characteristic feature of B-cell malignancies in clinics (Rossi et al. 1990), and furthermore, significant bone erosion has been found in all clinical stages of chronic lymphocytic leukemia (Marini et al. 2017). Therefore, drug repositioning to expand the clinical use of idelalisib could be a costefficient strategy for obtaining new treatment options for a variety of diseases including skeletal disorders and diseaserelated skeletal problems (Choi et al. 2015). Finally, our present findings could improve our understanding of the clinical value of PI3K δ as a druggable target and the efficacy of related therapeutics.

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Compliance with ethical standards

Conflict of interest The authors have declared no conflict of interest.

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