

# Lifelong accumulation of bone in mice lacking *Pten* in osteoblasts

Ximeng Liu\*, Katia J. Bruxvoort<sup>†</sup>, Cassandra R. Zylstra<sup>†</sup>, Jiarong Liu\*, Rachel Cichowski<sup>†</sup>, Marie-Claude Faugere<sup>‡</sup>, Mary L. Bouxsein<sup>§</sup>, Chao Wan\*, Bart O. Williams<sup>†¶</sup>, and Thomas L. Clemens<sup>\*¶</sup>

\*Department of Pathology, University of Alabama at Birmingham, Birmingham, AL 35294; <sup>†</sup>Laboratory of Cell Signaling and Carcinogenesis, Van Andel Research Institute, Grand Rapids, MI 49503; <sup>‡</sup>Department of Medicine, University of Kentucky, Lexington, KY 40536; and <sup>§</sup>Orthopedic Biomechanics Laboratory, Beth Israel Deaconess Medical Center, Boston, MA 02215

Edited by Richard P. Lifton, Yale University School of Medicine, New Haven, CT, and approved November 29, 2006 (received for review May 25, 2006)

**Bone formation is carried out by the osteoblast, a mesenchymal cell whose lifespan and activity are regulated by growth factor signaling networks. Growth factors activate phosphatidylinositol 3-kinase (PI3K), which enhances cell survival and antagonizes apoptosis through activation of Akt/PKB. This process is negatively regulated by the *Pten* phosphatase, which inhibits the activity of PI3K. In this study, we investigated the effects of Akt activation in bone *in vivo* by conditionally disrupting the *Pten* gene in osteoblasts by using Cre-mediated recombination. Mice deficient in *Pten* in osteoblasts were of normal size but demonstrated a dramatic and progressively increasing bone mineral density throughout life. *In vitro* osteoblasts lacking *Pten* differentiated more rapidly than controls and exhibited greatly reduced apoptosis in association with markedly increased levels of phosphorylated Akt and activation of signaling pathways downstream of activated Akt. These findings support a critical role for this tumor-suppressor gene in regulating osteoblast lifespan and likely explain the skeletal abnormalities in patients carrying germ-line mutations of *PTEN*.**

Akt | bone acquisition | osteoblast survival | high bone mass

The development and maintenance of the mammalian skeleton are controlled by actions of morphogens and growth factors on bone cells. Bone formation is carried out by the osteoblast, a mesenchymal cell whose lifespan and activity are regulated by growth factor signaling networks (1, 2). Skeletal growth factors such as insulin-like growth factor-1 (IGF-1) affect osteoblast proliferation and lifespan by activating anti-apoptotic pathways, increasing cell proliferation, and influencing differentiation (3). A key control point in many anti-apoptotic pathways is the lipid kinase phosphatidylinositol (PI) 3-kinase (PI3K), which is activated in response to various extracellular signals (4, 5). On activation of growth factor receptor tyrosine kinases, a p85 regulatory subunit of PI3K is recruited to phosphorylated tyrosine residues. This action engages and activates a catalytic subunit (p110) and induces phosphorylation of the inositol ring of PI(4)P or PI(4,5)P<sub>2</sub> at the D position to generate the second messengers PI(3,4)P<sub>2</sub> and PI(3,4,5)P<sub>3</sub> (4, 5). A key downstream target of PI3K and PIP<sub>3</sub> is the serine–threonine Akt kinase family (also known as protein kinase B). PIP<sub>3</sub> generated in the plasma membrane recruits Akt and PI-dependent kinase 1 (PDK1) through an interaction between the PI and the Akt or PDK1 pleckstrin homology (PH) domains. Once recruited to the plasma membrane, Akt is phosphorylated and activated by PDK1. Activated Akt promotes both cell growth and cell survival by regulating numerous downstream pathways including those controlled by GSK3, BAD/Bcl-XL, p53, and TSC2/mTOR (6).

The phosphatase and tensin homologue deleted on human chromosome 10 (*PTEN*) gene encodes a PIP phosphatase, which negatively regulates PI3K by dephosphorylating PI(3,4,5)P<sub>3</sub>. *PTEN* was originally cloned as a tumor suppressor for gliomas (7–9), but is deleted or inactivated in many other tumor types (10). Loss of *PTEN* function in either embryonic stem cells or human cancer cell lines results in an accumulation of PI(3,4,5)P<sub>3</sub> and persistent activation of Akt, leading to increases in cell proliferation, survival,

and migration (11–13). In addition, germ-line *PTEN* mutations are associated with Cowden disease, Bannayan–Zonana syndrome, and Lhermitte–Duclos disease, disorders characterized by hamartomas involving multiple organs (14, 15).

To directly investigate the role of *Pten* in osteoblasts *in vivo*, we disrupted the gene encoding *Pten* by using Cre-mediated gene replacement techniques. Mice carrying an osteoblast-specific deletion of *Pten* had normal body size but demonstrated progressive increases in bone volume and density throughout life. Osteoblasts from the mutant mice demonstrated a striking decreased susceptibility to apoptosis and accelerated differentiation capacity. These findings provide *in vivo* evidence that signaling via the PI3K/*Pten* pathway promotes osteoblast survival and function.

## Results

**Osteoblast-Specific Disruption of the *Pten* Gene Increases Postnatal Bone Acquisition.** OC-Cre<sup>TG/+</sup>; *Pten*<sup>flx/flx</sup> mice, which were viable and fertile, were crossed with *Pten*<sup>flx/flx</sup> mice to generate litters in which half of the progeny were of the OC-Cre<sup>TG/+</sup>; *Pten*<sup>flx/flx</sup> genotype (referred to as  $\Delta Pten$ ), whereas the other half were control littermates (lacking the OC-Cre transgene and, therefore, wild type for *Pten* gene function). PCR analysis by using DNA templates from tissues of  $\Delta Pten$  offspring confirmed that Cre-mediated recombination occurred exclusively in bone (Fig. 1C).

We generated cohorts of these animals and performed a longitudinal analysis of their whole-body bone mineral density (BMD) by using dual-energy x-ray absorptiometry (Fig. 2A). At 3 months of age, mice of both sexes displayed significant increases in BMD relative to their wild-type cohorts ( $P = 0.002$  in both cases). BMD increased in mutant mice relative to controls ( $P = 0.002$ ) at 6 weeks (data not shown) and increased progressively as the animals aged. By 15 months of age, female  $\Delta Pten$  mice had 71% higher whole-body BMD than controls ( $P = 2 \times 10^{-6}$ ), whereas BMD in males increased by 60% ( $P = 5 \times 10^{-5}$ ). BMD was similarly increased in both axial and peripheral skeletal sites at 12 months (Fig. 2B and C).

Three-dimensional micro computed tomography (microCT) scanning and histological analysis of mutant long bones showed pronounced increases in both cortical and trabecular bone. Trabeculae from mutant femurs were thicker and less separated than those of controls (Fig. 3). Cortical thickness in the femurs was

Author contributions: X.L. and K.J.B. contributed equally to this work; B.O.W. and T.L.C. designed research; X.L., K.J.B., C.R.Z., J.L., R.C., M.-C.F., M.L.B., C.W., and B.O.W. performed research; X.L., K.J.B., C.R.Z., M.-C.F., M.L.B., B.O.W., and T.L.C. analyzed data; and X.L., K.J.B., B.O.W., and T.L.C. wrote the paper.

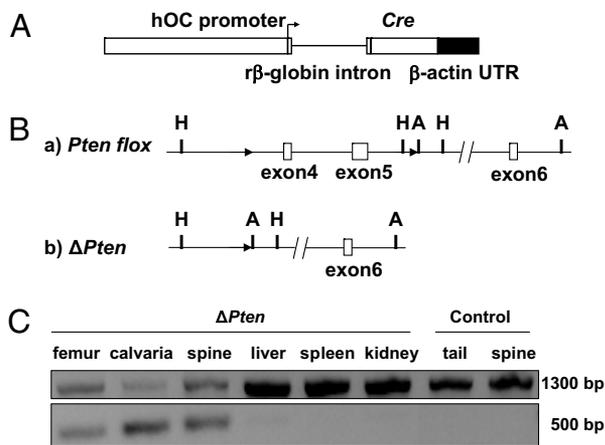
The authors declare no conflict of interest.

This article is a PNAS direct submission.

Abbreviations: 7-AAD, 7-aminoactinomycin D; BMD, bone mineral density; IGF-1, insulin-like growth factor-1; microCT, micro computed tomography; moi, multiplicity of infection; PI, phosphatidylinositol; PDK1, PI-dependent kinase 1; PI3K, PI 3-kinase.

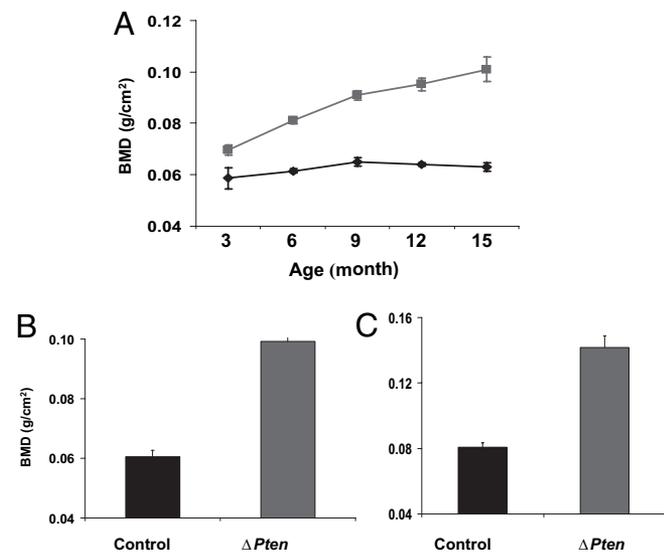
<sup>¶</sup>To whom correspondence may be addressed. E-mail: bart.williams@vai.org or tclemens@path.uab.edu.

© 2007 by The National Academy of Sciences of the USA



**Fig. 1.** Strategy for osteoblast-specific deletion of *Pten*. (A) Human osteocalcin (hOC) construct used to generate mouse lines expressing a *Cre* transgene. Cloning of this construct is described in ref. 37. The arrow indicates the transcriptional orientation. (B) Strategy for generation of a *loxP*-flanked allele of the *Pten* gene. (a) *Pten*-floxed allele. Exons 4 and 5 were flanked by two *loxP* sequences, shown as black arrowheads (A, Apal; H, HindIII). (b) *Cre*-mediated deletion produces the  $\Delta Pten$  allele. (C) Allele-specific PCR detection of the  $\Delta Pten$  allele in mutant and control mice. (Upper) PCR fragments generated for the *Pten*-floxed allele. (Lower) *Pten* deletion ( $\Delta Pten$ ) allele. Deletion of exons 4 and 5 was detected only in skeletal tissues.

increased by 43% at 3 months of age (data not shown) and by 250% ( $256 \pm 7$  vs.  $676 \pm 9$ , mean  $\pm$  SE) at 12 months of age (Fig. 3). The thickness of calvarial bone was similarly increased in the mutants, suggesting that loss of *Pten* influences the development of bone formed through intramembranous processes (Fig. 4A). Calvaria from mice heterozygous for the conditional *Pten* allele (*OC-Cre*<sup>TG/+</sup>; *Pten*<sup>lox/+</sup>) were slightly but consistently thicker than those

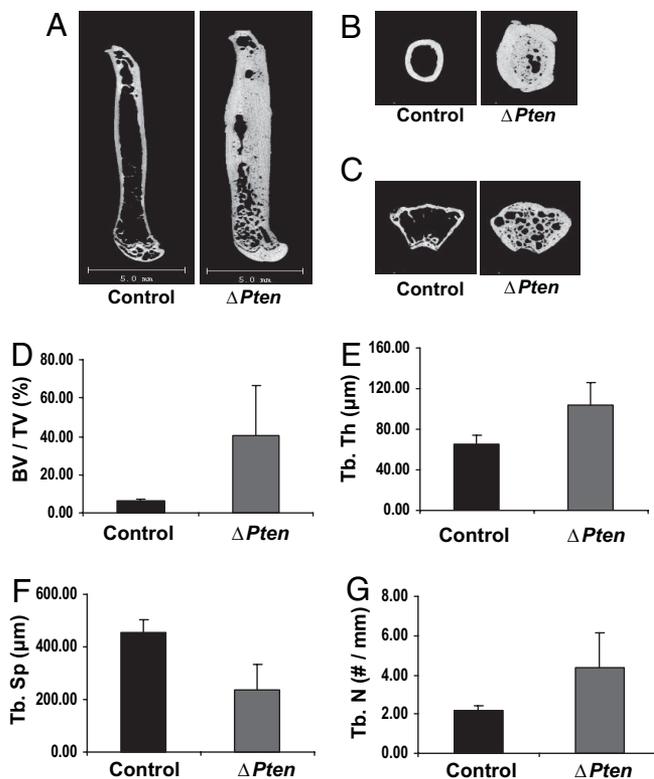


**Fig. 2.** Deletion of *Pten* leads to cumulative increases in BMD. (A) Mice carrying an osteoblast-specific deletion of *Pten* exhibit progressive increases in BMD with age. ■,  $\Delta Pten$ ; ◆, wild-type cohorts (male mice shown). The error bars indicate SEM. At least five mice in each category were examined. The increase in BMD was already statistically significant ( $P < 0.002$ ) at 3 months of age for both sexes (females not shown), and the differences steadily increased with age. (B and C) BMD was statistically elevated ( $P < 10^{-5}$ ) for both the lumbar spine (B) and femur (C) in 12-month-old male (shown) and female (data not shown)  $\Delta Pten$  mice relative to control littermates (at least five mice were evaluated for each category).

of controls (Fig. 4A). In addition, serum alkaline phosphatase levels in the heterozygous mice were intermediate between the wild-type and homozygous mutant animals (Fig. 4B). These results indicate that loss of a single *Pten* allele results in increased bone mass.

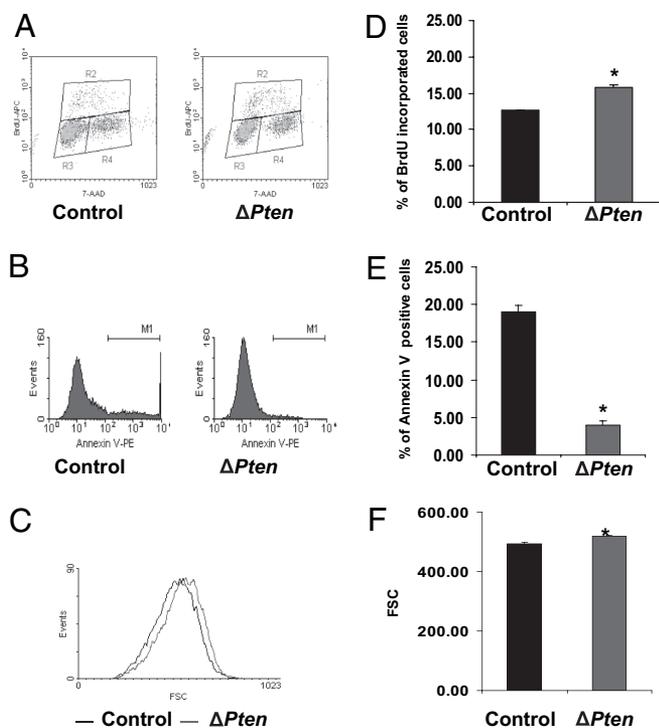
To determine the impact of the *Pten* mutation on bone-formation rates, dynamic histomorphometric measurements were performed on 6-week-old mice doubly labeled with sequential doses of calcein. Cancellous bone formation and mineral apposition rates were significantly increased in the  $\Delta Pten$  mice compared with controls (Fig. 5). In addition, the number of osteoblasts lining bone surfaces was increased, and serum osteocalcin, a marker of osteoblastic activity, was also elevated, although not to a statistically significant level (data not shown). There was no evidence of defective mineralization; osteoid volume and mineralization lag time were similar to control values (data not shown). Finally, we found no evidence for a defect in bone resorption; osteoclast numbers were increased, suggesting appropriate coupling of bone formation with resorption (Fig. 5E), and circulating osteoprotegerin levels were similar to those measured in control serum (data not shown).

**Constitutive Activation of Akt in Osteoblasts Deficient in *Pten*.** To examine the effect of *Pten* loss on signaling pathways downstream of Akt, we analyzed the expression of several key Akt targets after *Cre*-mediated disruption of the *Pten* gene in calvarial osteoblasts *in vitro*. Cells derived from mice carrying the floxed alleles were infected with adenovirus expressing the *Cre* recombinase (Adeno-*Cre*) or a control adenovirus directing the expression of green fluorescent protein (Adeno-GFP) and then cultured in the presence of  $\beta$ -glycerol phosphate and ascorbate (mineralizing medium). Infection with a virus concentration of  $15 \times 10^6$  pfu/ml was sufficient to decrease *Pten* mRNA and protein to 10% of the levels



**Fig. 3.** Disruption of the *Pten* gene increases bone volume. MicroCT analysis was performed on femur (A), midshaft femur (B), and distal femur (C) of  $\Delta Pten$  and control mice at 12 months of age. Bar graphs show the trabecular thickness (Tb. th) ( $P < 0.006$ ) (D), ratio of bone volume to tissue volume (BV/TV) (E), trabecular separation (Tb. Sp) (F), and trabecular number (Tb. N) (G), respectively. Error bars represent SEM.



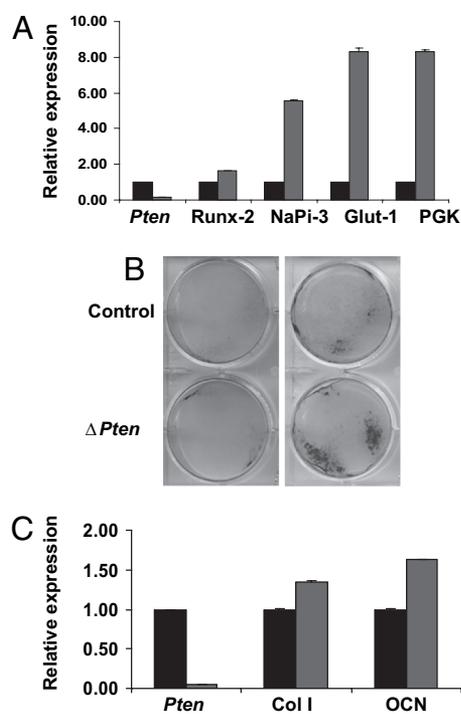


**Fig. 7.** Loss of *Pten* in osteoblasts *in vitro* reduces apoptosis. (A and D) BrdU incorporation. Primary osteoblasts were infected with Adeno-Cre or Adeno-GFP virus at a multiplicity of infection (moi) of 100 for 48 h. Cells were then cultured in  $\alpha$ MEM with 1% serum for 24 h and then  $\alpha$ MEM with 10% serum for an additional 12 h before harvesting cells for BrdU and 7-aminoactinomycin C (7-AAD) staining. BrdU was added to the medium 1 h before harvesting cells. (A) Density plot of the cell population in  $G_0/G_1$  (region R3),  $G_2/M$  (region R4), and S phase (region R2). (D) Percentage of BrdU incorporation (mean  $\pm$  SD) in  $\Delta Pten$  (with Adeno-Cre) and control (with Adeno-GFP) osteoblasts ( $n = 3$ ,  $P = 0.000315$ ). (B and E) Annexin V staining. Primary osteoblasts were infected with Adeno-Cre or Adeno-GFP virus at a moi of 100 for 48 h. Cells were then exposed to  $\alpha$ MEM with 1% serum for 24 h and serum-free  $\alpha$ MEM for an additional 12 h to induce apoptosis before harvesting cells for annexin V staining. Samples were analyzed by FACSCalibur immediately after staining. (B) Representative histograms of annexin V-positive cells ( $M_1$ ). PE, phycoerythrin. (E) Percentage of annexin V-positive cells in  $\Delta Pten$  (with Adeno-Cre) and control (with Adeno-GFP) osteoblasts ( $n = 3$ ,  $P = 1.21 \times 10^{-5}$ ). (C) *Pten* deletion increases forward scatter (FSC) of osteoblast cells. *Pten*-deficient cells are represented by the trace shifted to the right. (F) Mean FSC of control and  $\Delta Pten$  osteoblasts. \*, Significantly different from control at  $P < 0.05$ .

To determine the effect of loss of *Pten* on osteoblast differentiation, cells were differentiated in the presence of  $\beta$ -glycerol phosphate and ascorbate. Osteoblasts deficient in *Pten* showed increased expression of several genes, including *Runx-2*, *NaPi3*, *Glut-1*, and *PGK*, which in our hands are expressed early in the differentiation of primary mouse osteoblasts (Fig. 8A). At later times, disruption of *Pten* was associated with increased alkaline phosphatase expression (Fig. 8B) and augmented expression of gene markers for the differentiated osteoblast, including collagen I and osteocalcin (Fig. 8C). Overall, these results are compatible with the increased numbers and activity of osteoblasts seen in mice lacking *Pten* *in vivo*.

## Discussion

In this study, we investigated the role of the *Pten* phosphatase in bone by using Cre-mediated recombination strategies to conditionally disrupt this gene in osteoblasts. We show that loss of *Pten* profoundly increases bone mass at all skeletal sites by increasing the number of osteoblasts, most likely through retarding apoptosis.



**Fig. 8.** Loss of *Pten* accelerates osteoblast differentiation *in vitro*. (A and C) Effect of *Pten* deletion on osteoblast gene expression. Primary osteoblasts were infected with Adeno-Cre or Adeno-GFP virus at a moi of 100 for 48 h (A) or 21 days (C). Relative expression of the indicated mRNAs in osteoblasts expressing (black bars) or lacking (gray bars) *Pten* was determined by real-time PCR. Col I, collagen I; OCN, osteocalcin. (B) Effect of *Pten* deletion on alkaline phosphatase expression. Primary osteoblasts were infected with Adeno-Cre or Adeno-GFP virus at a moi of 100 and then grown in medium containing ascorbate and  $\beta$ -glycerol phosphate. Monolayers were stained for alkaline phosphatase at 10 (Left) and 21 days (Right) of culture. (C) Expression of Collagen I (Col I) and osteocalcin (OCN) was also enhanced in the absence of *Pten*.

The role of PTEN as a tumor suppressor is illustrated by the fact that it is frequently mutated or lost in human cancers (19–21). Homozygosity for the null mutation (*Pten*<sup>-/-</sup>) in mice results in early embryonic lethality, indicating that the function of *Pten* is critical for normal development (22–24). *Pten* heterozygous mice develop a spectrum of neoplasias in organs such as the breast, thyroid, endometrium, and prostate, which closely resembles the situation in humans heterozygous for a *PTEN* mutation (22, 25–27). The bone mass in such individuals has not been reported.

Growth factors such as insulin and IGF-1 signal through the PI3K pathway, which promotes proliferation and inhibits apoptosis via activation of Akt/PKB in a variety of cell types (28). Previous studies also support a role for this pathway in bone progenitor cell differentiation. For example, in mesenchymal stromal cells, blockade of insulin/IGF-1-activated PI3K/AKT pathways decreased bone morphogenetic protein-induced alkaline phosphatase and osteopontin expression in human mesenchymal stem cells (6). In addition, enhanced osteoblastic differentiation by forced expression of the transcription factor Runx-2 in C3H/10T1/2 mouse embryo fibroblasts was blocked by treatment with IGF-1 antibody, inhibiting Akt signaling with LY294002, or through adenoviral introduction of dominant-negative AKT. The current observations provide *in vivo* validation of the importance of *Pten* in regulating PI3K signaling in osteoblasts and suggest that sustained activation of the PI3K pathway in mature osteoblasts lacking *Pten* enables these cells to extend their lifespan.

The precise downstream targets of PI3K/Akt in osteoblasts are unknown. In a number of tissues, the biological response to the

activation of the PI3K/Akt pathway appears to manifest in a tissue-specific fashion. For example, *Pten* deficiency in the prostate (29, 30) leads to the activation of FOXO3a and p70S6K pathways and enhanced proliferation, whereas in the bladder, which is less sensitive to tumor formation, *Pten* deficiency leads to decreased levels of p27, increased levels of p21, and decreased proliferation rates (31). Previous studies using genetic methods similar to those used here have shown that selective deletion of *Pten* in the liver (32) and skin (33) resulted in increased organ size. Surprisingly, however, disruption of *Pten* in adipocytes (34, 35) and muscle (36) increased sensitivity to insulin, but did not change the number of fat or muscle cells or alter the overall size of these organs. In the case of the adipocyte, a reduction in the level of resistin, a secretory protein that plays a key role in the development of insulin resistance, may mediate the enhanced sensitivity to insulin (36). Thus, it appears that signals generated by activation of the PI3K pathway have an impact on different functional gene pathways even among cells of the mesenchymal lineage.

In conclusion, we demonstrate that loss of *Pten* function in murine osteoblasts positively influences postnatal bone acquisition through constitutive activation of the PI3K pathway. We attribute this anabolic effect to a decreased rate of apoptosis, which would prolong the time each osteoblast spends in synthesizing and mineralizing bone matrix.

## Materials and Methods

**Generation of OC-Cre<sup>TG/+</sup>; *Pten*<sup>flox/flox</sup> Mice.** Mice lacking *Pten* in osteoblasts were generated by crossing OC-Cre mice (37, 38) with homozygous conditional mutants carrying modified *Pten* alleles (with *loxP* sites flanking exons 4 and 5) (39) to generate OC-Cre<sup>TG/+</sup>; *Pten*<sup>flox/flox</sup> progeny and control littermates (lacking the OC-Cre gene; Fig. 1). Mice homozygous for floxed *Pten* alleles were crossed to animals homozygous for the osteocalcin-Cre transgene to generate OC-Cre<sup>TG/+</sup>; *Pten*<sup>flox/+</sup> mice. These mice were then crossed with *Pten*<sup>flox/flox</sup> mice to generate litters that contained approximately 1/4 OC-Cre<sup>TG/+</sup>; *Pten*<sup>flox/flox</sup> mice, which were then used for subsequent crosses. The specificity and efficiency of recombination at several loci were reported previously by using these animals (37). All experiments performed were in compliance with the guiding principles of the *Guide for the Care and Use of Laboratory Animals* (ref. 40, available at [www.nap.edu/books/0309053773/html](http://www.nap.edu/books/0309053773/html)) and approved before use by the Institutional Animal Care and Use Committee of the Van Andel Research Institute.

**Genotype Analysis.** DNA was prepared from tail biopsies by using an AutoGenprep 960 automated DNA isolation system (AutoGenM, Holliston, MA). PCR-based strategies were then used to genotype these mice (details available on request). The Cre transgene was detected by PCR (1 min at 94°C, 1 min at 53°C, and 1 min at 72°C, for 30 cycles) using the primers 5'-CAAATAGCCCTGCGAGATTC-3' (forward) and 5'-TGATACAAGGGACATCTTCC-3' (reverse) to generate a 260-bp product corresponding to a portion of the OC promoter and the rabbit  $\beta$ -globin intron. The status of the *Pten* locus was determined by PCR with the primers 5'-GTCACCAGGATGCTTCTGAC-3' (forward) and 5'-GAAACGGCCTTAACGACGTAG-3' (reverse).

**Deminerlized Bone Histology.** Tissue samples were fixed in formalin overnight, decalcified in Immunocal decalcifying agent (Decal, Baltimore, MD) overnight, and then dehydrated through a graded alcohol series in a Ventana Renaissance processor (Ventana Medical Systems, Tucson, AZ). Tissues were paraffin-embedded, and 5- $\mu$ m sections were adhered to glass slides. Slides were deparaffinized for immunohistochemistry.

**Mineralized Bone Histology.** Femurs were fixed in ethanol at room temperature, dehydrated, and embedded in methyl methacrylate. Three-micrometer sections were cut with a Microm microtome

(Thornwood, NY) and stained with modified Mason–Goldner trichrome stain. The number of osteoblasts and osteoclasts per bone perimeter was measured at standardized sites under the growth plate by using a semiautomatic method (Osteoplan II; Kontron, Munich, Germany) at a magnification of  $\times 200$ . These parameters comply with guidelines of the nomenclature committee of the American Society of Bone and Mineral Research (38).

**Dual Energy X-Ray Absorptiometry.** Mice were anesthetized through inhalation of 2% isoflurane with oxygen (1.0 liter/min) before and during the procedure ( $\leq 5$  min). The mice were placed on a specimen tray in a Lunar PIXImus II bone densitometer (GE Healthcare, Piscataway, NJ) for analysis. BMD was calculated by the PIXImus software based on the active bone area in the subcranial region within the total body image, specifically in the femur and spine. Calibrations were performed with a phantom by using known values, and quality assurance measurements were performed daily with this same phantom.

**MicroCT.** High-resolution images of the femur were acquired by using a desktop microtomographic imaging system (MicroCT40; Scanco Medical, Basserdorf, Switzerland). The femur was scanned at 45 keV with an isotropic voxel size of 6  $\mu$ m, and the resulting 2-dimensional cross-sectional images are shown in gray scale. Scanning was started in the mid-epiphysis and extended proximally for  $\approx 3.6$  mm (600 microCT slices).

**Osteoblast Isolation and Culture.** Osteoblasts were isolated from calvaria of newborn *Pten*<sup>flox/flox</sup> mice by serial digestion in  $\alpha$ MEM (Mediatech, Herndon, VA) containing 10% BSA, 25 mM Hepes at pH 7.4, 0.2 mg/ml collagenase type I (Worthington, Lakewood, NJ), 0.7 mg/ml collagenase type 2 (Worthington), and 1 mM CaCl<sub>2</sub>. Calvaria were digested for 15 min at 37°C with constant agitation. The digestion solution was collected, washed with fresh medium, and digested an additional five times. Digestions 3–6 (containing the osteoblasts) were centrifuged, washed with  $\alpha$ MEM containing 10% FBS and 1% penicillin/streptomycin, and cultured for 48 h at 37°C.

**Adenovirus Infection.** Monolayer osteoblasts were infected with control adenovirus (Adeno-GFP) or Cre-recombinase virus (Adeno-Cre; Microbix Biosystems, Toronto, ON, Canada) at a moi of 100 for most experiments. Osteoblasts were harvested after 48 h. mRNA was extracted from osteoblasts for *Pten* mRNA quantitation by real-time PCR.

**Osteoblast Proliferation.** After exposure to adenovirus, osteoblasts were plated on 100-mm plates ( $1.0 \times 10^6$  per plate) and cultured in  $\alpha$ MEM with 1% FBS for 48 h and  $\alpha$ MEM with 10% serum for 12 h before harvesting cells for BrdU and 7-AAD staining. BrdU was added to the medium 1 h before harvesting cells. After staining with anti-BrdU-APC and 7-AAD (BD PharMingen, San Diego, CA), cells were analyzed by FACSCalibur (Becton Dickinson, Franklin Lakes, CA), and 20,000 events were collected. Results were analyzed with WinMDI version 2.8 (The Scripps Research Institute, La Jolla, CA).

**Osteoblast Apoptosis.** Osteoblasts were plated on 100-mm plates ( $1.0 \times 10^6$  per plate) and cultured in  $\alpha$ MEM with 1% FBS for 48 h and serum-free  $\alpha$ MEM for 12 h before harvesting cells for annexin V-PE (BD PharMingen) staining. After staining with annexin V-PE, cells were analyzed immediately by FACSCalibur (Becton Dickinson), and 20,000 events were collected.

**Protein Extraction and Immunoblots.** Proteins were extracted from cultured osteoblasts by using M-PER (Pierce, Rockford, IL) with protease inhibitors (Pierce). The extracts were separated on SDS/10% polyacrylamide gels and transferred to nitrocellulose mem-

branes. After probing with primary antibodies (Cell Signaling Technologies, Danvers, MA), the membranes were incubated with horseradish peroxidase-linked secondary anti-rabbit antibodies (Cell Signaling Technologies), and bound antibodies were visualized using the Supersignal West Femto Maximum Sensitivity Substrate (Pierce).

**Alkaline Phosphatase Staining.** Alkaline phosphatase activity was measured in cell layers as previously described by using a *p*-nitrophenyl phosphate substrate and an incubation temperature of 37°C (41). Two days after adenovirus treatment, osteoblasts were replated in six-well plates with  $1.0 \times 10^5$  cells per well. Cells were cultured in  $\alpha$ MEM with 10% FBS (GIBCO, Carlsbad, CA) and supplemented with L-ascorbic acid (50  $\mu$ g/ml; Sigma, St. Louis, MO) and  $\beta$ -glycerol phosphate (10 mM; Sigma) for 21 days. Media were changed every other day.

**Real-Time PCR.** Total RNA was extracted from cells by using the TRIzol (Life Technologies, Carlsbad, CA) extraction protocol. First-strand cDNA was synthesized by using the Invitrogen kit. Real-time PCR was performed at 57°C for 30 cycles in the Opticon Continuous Fluorescent Detector by using IQ SYBR Green supermix (Bio-Rad, Hercules, CA). Samples were run in triplicate and the results were normalized to  $\beta$ -actin expression. Primer sequences used were as follows: *Pten*, F5'-CATAACCCACCACAGCTAG-3'

and R5'-GCAGACCACAACTGAGG-3'; Runx-2, F5'-CCAAATTTGCCTAACCAGAATG-3' and R5'-GAGGCTGTGGTTTCAAAGCAC-3'; NaPi-3, F5'-CACCCATATGGCTTCTGCTT-3' and R5'-CAGGAATTCATAGCCCAGGA-3'; Glut-1, F5'-CAGTTTCGAGAAGAACATGAG-3' and R5'-GC-GGAATTC AATGCTGATGAT-3'; PGK, F5'-GGAAGCGGGTTCGTGATGA-3' and R5'-GCCTTGATCCTTTGGTTGTTTT-3';  $\beta$ -actin, F5'-CTGAACCCTAAGGCCAACCGTG-3' and R5'-GGCATACAGGGACAGCACAGCC-3'.

**Statistical Analysis.** Results were expressed as mean  $\pm$  SD or SEM as indicated. All statistical tests were two sided. An assigned significance level of 0.05 was used. Comparability of the two groups of data was assessed by using Student's *t* test.

We thank Tak Mak (University of Toronto, Toronto, ON, Canada) for providing *Pten*-flox mice, Pam Swiatek and the Laboratory of Germline Modification core facility for Autogen DNA processing, Bryn Eagleson and the VARI Vivarium core facility for expert animal husbandry, and Cynthia Knight and David Nadziejka for assistance in preparing the manuscript. This work was supported by National Institutes of Health Grants AR49410, AR52746, and P30 AR046031 and a Merit Review Grant from the Department of Veterans Affairs (to T.L.C.). Additional support was provided by the Van Andel Institute. We acknowledge the Michigan Economic Development Corporation and the Michigan Technology Tri-Corridor for supporting these cores (Michigan Animals Models Consortium Grant 085P1000815).

- Mundy GR, Boyce B, Hughes D, Wright K, Bonewald L, Dallas S, Harris S, Ghosh-Choudhury N, Chen D, Dunstan C (1995) *Bone* 17:71S–75S.
- Karsenty G (2000) *Matrix Biol* 2000 19:85–89.
- Nakae J, Kido Y, Accili D (2001) *Endocr Rev* 22:818–835.
- Cantley LC (2002) *Science* 296:1655–1657.
- Cantley LC, Neel BG (1999) *Proc Natl Acad Sci USA* 96:4240–4245.
- Downward J (2004) *Semin Cell Biol* 15:177–182.
- Li DM, Sun H (1997) *Cancer Res* 57:2124–2129.
- Li J, Yen C, Liaw D, Podsypanina K, Bose S, Wang SI, Puc J, Miliarensis C, Rodgers L, McCombie R, et al. (1997) *Science* 275:1943–1947.
- Steck PA, Pershouse MA, Jasser SA, Yung WK, Lin H, Ligon AH, Langford LA, Baumgard ML, Hattier T, Davis T, et al. (1997) *Nat Genet* 15:356–362.
- Simpson L, Parsons R (2001) *Exp Cell Res* 264:29–41.
- Liliental J, Moon SY, Lesche R, Mamillapalli R, Li D, Zheng Y, Sun H, Wu H (2000) *Curr Biol* 10:401–404.
- Stambolic V, Suzuki A, de la Pompa JL, Brothers GM, Mirtsos C, Sasaki T, Ruland J, Penninger JM, Siderovski DP, Mak TW (1998) *Cell* 95:29–39.
- Sun H, Lesche R, Li DM, Liliental J, Zhang H, Gao J, Gavrilova N, Mueller B, Liu X, Wu H (1999) *Proc Natl Acad Sci USA* 96:6199–6204.
- Pilarski R, Eng C (2004) *J Med Genet* 41:323–326.
- Thiffault I, Schwartz CE, Der Kaloustian V, Foulkes WD (2004) *Am J Med Genet A* 130:123–127.
- Xing L, Boyce BF (2005) *Biochem Biophys Res Commun* 328:709–720.
- Backman S, Stambolic V, Mak T (2002) *Curr Opin Neurobiol* 12:516–522.
- Plas DR, Thompson CB (2005) *Oncogene* 24:7435–7442.
- Bose S, Chandran S, Mirocha JM, Bose N (2006) *Mod Pathol* 19:238–245.
- Yan X, Fraser M, Qiu Q, Tsang BK (2006) *Gynecol Oncol* 102:348–355.
- Davies MA, Kim SJ, Parikh NU, Dong Z, Bucana CD, Gallick GE (2002) *Clin Cancer Res* 8:1904–1914.
- Di Cristofano A, Pesce B, Cordon-Cardo C, Pandolfi PP (1998) *Nat Genet* 19:348–355.
- Suzuki H, Freije D, Nusskern DR, Okami K, Cairns P, Sidransky D, Isaacs WB, Bova GS (1998) *Cancer Res* 58:204–209.
- Suzuki A, Tsutomi Y, Akahane K, Araki T, Miura M (1998) *Oncogene* 17:931–939.
- Suzuki A, de la Pompa JL, Stambolic V, Elia AJ, Sasaki T, del Barco Barrantes I, Ho A, Wakeham A, Itie A, Khoo W, et al. (1998) *Curr Biol* 8:1169–1178.
- Stambolic V, Tsao MS, MacPherson D, Suzuki A, Chapman WB, Mak TW (2000) *Cancer Res* 60:3605–3611.
- Podsypanina K, Ellenson LH, Nemes A, Gu J, Tamura M, Yamada KM, Cordon-Cardo C, Catoretti G, Fisher PE, Parsons R (1999) *Proc Natl Acad Sci USA* 96:1563–1568.
- Levine AJ, Feng Z, Mak TW, You H, Jin S (2006) *Genes Dev* 20:267–275.
- Grunwald V, DeGraffenried L, Russel D, Friedrichs WE, Ray RB, Hidalgo M (2002) *Cancer Res* 62:6141–6145.
- Trotman LC, Alimonti A, Scaglioni PP, Koutcher JA, Cordon-Cardo C, Pandolfi PP (2006) *Nature* 441:523–527.
- Fradet Y, Lacombe L (2000) *Curr Opin Urol* 10:441–445.
- Stiles B, Wang Y, Stahl A, Bassilian S, Lee WP, Kim YJ, Sherwin R, Devaskar S, Lesche R, Magnuson MA, Wu H (2004) *Proc Natl Acad Sci USA* 101:2082–2087.
- Backman SA, Ghazarian D, So K, Sanchez O, Wagner KU, Hennighausen L, Suzuki A, Tsao MS, Chapman WB, Stambolic V, Mak TW (2004) *Proc Natl Acad Sci USA* 101:1725–1730.
- Nakashima N, Sharma PM, Imamura T, Bookstein R, Olefsky JM (2000) *J Biol Chem* 275:12889–12895.
- Tang X, Powelka AM, Soriano NA, Czech MP, Guilherme A (2005) *J Biol Chem* 280:22523–22529.
- Wijesekara N, Konrad D, Eweida M, Jefferies C, Liadis N, Giacca A, Crackower M, Suzuki A, Mak TW, Kahn CR, et al. (2005) *Mol Cell Biol* 25:1135–1145.
- Zhang M, Xuan S, Bouxsein ML, von Stechow D, Akeno N, Faugere MC, Malluche H, Zhao G, Rosen CJ, Efstratiadis A, Clemens TL (2002) *J Biol Chem* 277:44005–44012.
- Holmen SL, Zylstra CR, Mukherjee A, Sigler RE, Faugere MC, Bouxsein ML, Deng L, Clemens TL, Williams BO (2005) *J Biol Chem* 280:21162–21168.
- Suzuki A, Yamaguchi MT, Ohteki T, Sasaki T, Kaisho T, Kimura Y, Yoshida R, Wakeham A, Higuchi T, Fukumoto M, et al. (2001) *Immunity* 14:523–534.
- Institute of Laboratory Animal Resources (1996) *Guide for the Care and Use of Laboratory Animals* (Natl Acad Press, Washington, DC).
- Manolagas SC, Burton DW, Deftos LJ (1981) *J Biol Chem* 256:7115–7117.