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# Disruption of LRP6 in osteoblasts blunts the bone anabolic activity of PTH

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#### Abstract

Mutations in Low-density lipoprotein receptor-related protein 6 (LRP6) are associated with human skeletal disorders. LRP6 is required for parathyroid hormone (PTH)-stimulated signaling pathways in osteoblasts. We investigated whether LRP6 in osteoblasts directly regulates bone remodeling and mediates the bone anabolic effects of PTH by specifically deleting LRP6 in mature osteoblasts in mice (LRP6 KO). Three month-old LRP6 KO mice had a significant reduction in bone mass in the femora secondary spongiosa relative to their wild type littermates, whereas marginal changes were seen in femoral tissue of 1 month-old LRP6 KO mice. The remodeling area of the 3 month-old LRP6 KO mice showed a decreased bone formation rate as detected by Goldner's Trichrome staining and calcein double labeling. Bone histomorphometric and immumohistochemical analysis revealed a reduction in osteoblasts but little change in the numbers of osteoclasts and osteoprogenitors/osteoblast precursors in LRP6 KO mice compared to wild type littermates. In addition, the percentage of the apoptotic osteoblasts on the bone surface was higher in LRP6 KO mice compared to wild type littermates. Intermittent injection of PTH had no effect on bone mass nor osteoblastic bone formation in either trabecular and cortical bone in LRP6 KO mice, whereas all were enhanced in wild type littermates. Additionally, the antiapoptotic effect of PTH on osteoblasts in LRP6 KO mice was less significant compared to wild type mice. Therefore, our findings demonstrate that LRP6 in osteoblasts is essential for osteoblastic differentiation during bone remodeling and the anabolic effects of PTH.

#### Keywords

LRP6; PTH; bone remodeling; osteoblast differentiation; osteoblast apoptosis

#### Introduction

The regulation of bone mass, or bone remodeling in adults, is a dynamic process orchestrated by bone-forming osteoblasts and bone-resorbing osteoclasts. Growth factors

#### Disclosure

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such as active TGF and IGF-1 are released during osteoclastic bone resorption to recruit mesenchymal stem cells (MSCs) and promote their differentiation to osteoblasts to couple new bone formation to bone resorption (1–3). An imbalance of osteoblastic bone formation and osteoclastic bone resorption leads to bone disorders such as osteoporosis. Parathyroid hormone (PTH) stimulates bone remodeling and is the only FDA-approved anabolic therapy for osteoporosis as intermittent administration of PTH selectively stimulates bone formation (4–6). The bone anabolic effect of PTH also has been demonstrated extensively in mice and rats (7). Histologic studies have shown that the increase in bone formation is largely due to an increase in the number of matrix-synthesizing osteoblasts (7;8). However, the signaling mechanisms responsible for PTH-mediated effects on bone are still incompletely understood as the traditional signaling pathway of cyclic AMP (cAMP) production and protein kinase A (PKA) activation does not provide a satisfactory explanation for all the anabolic effects of PTH.

Low-density lipoprotein receptor-related protein 5 and 6 (LRP5 and 6) are primarily known for their key role in mediating the transduction of signals from secreted Wnt proteins to catenin (9–13). Although they have similar structures, several lines of evidence suggest that LRP5 and LRP6 may have unique roles regarding skeletal growth and bone remodeling. Mutations in *LRP5* or *LRP6* are associated with distinct phenotypes of skeletal diseases. Inactivation of LRP5 was identified as the causative genetic alteration underlying Osteoporosis-Pseudoglioma Syndrome (OPPG) (14), a rare syndrome associated with premature, generalized osteoporosis leading to bone fracturing and progressive blindness. A gain-of-function mutation in LRP5 (G171V) has been identified in individuals displaying high-bone-mass (15–17). Individuals with a LRP6 mutation show early coronary disease and severe osteoporosis (18). A common protein variant of LRP6 (Ile1062Val) is associated with fracture risk in elderly men (19;20). Mouse genetic studies also reveal distinct roles LRP5 and LRP6 in bone. Lrp5 knockout mice are viable but suffer from osteoporosis in adulthood (21). In contrast, Lrp6 knockout mice are perinatal lethal due to developmental abnormalities like truncations of the axial skeleton, limb defects, and loss of the paraxial mesoderm (10;22). The analysis of the phenotypes of mice carrying heterozygous mutations of Lrp6 and either heterozygous or homozygous mutations of Lrp5 show that Lrp6 participates in the control of bone mass accrual in a manner that suggests more than a simple redundancy with Lrp5 (23). These studies suggest that LRP6 is involved in both bone development and bone remodeling in adults.

We have previously found that PTH orchestrates the signaling pathways of different growth factors, such as Wnts, TGF and BMPs, that directly regulate osteoblast differentiation and function (24–26). Particularly, PTH stabilizes -catenin through LRP6 (24). LRP6 is also required for PTH-mediated cAMP production (27). PTH promotes the interaction of the cytoplasmic domain of LRP6, but not LRP5, with G s to set up a functional PTH1R-G s-adenylate cyclase complex for the rapid production of cAMP and subsequent PKA activation. Furthermore, we recently found that LRP6 in MSCs acts as a negative regulator for BMP-induced MSC commitment to the osteoblastic lineage. PTH disrupts the LRP6-organized extracellular antagonist network by inducing endocytosis of a PTH1R-LRP6-antagonist complex, resulting in increased commitment and differentiation of MSCs to osteoblasts (26). Notably, PTH-stimulated bone formation can still occur in LRP5-deficient mice (28;29), indicating that LRP5 is not required for PTH anabolic actions in bone. Collectively, these findings suggest that LRP6 specifically is a key element in PTH-mediated signaling pathways enhancing osteoblastic numbers and function.

It is known that PTH enhances the number and the activation of osteoblasts through increasing osteoblast proliferation and differentiation, and decreasing osteoblast apoptosis (7;8;30;31). As LRP6 is a key element in PTH-elicited cAMP/PKA and -catenin signaling,

it may mediate the effects of PTH on osteoblasts and its positive effect on bone formation. Here we systemically analyzed the function of LRP6 in bone remodeling in osteoblastspecific LRP6-deficient mice. We found that LRP6 in osteoblasts is required for osteoblast differentiation and survival during bone remodeling in adult mice. Importantly, PTH anabolic effects were blunted in the LRP6 KO mice.

#### **Materials and Methods**

#### Antibodies and reagents

Primary antibodies including goat anti-LRP6 (Abcam), rabbit anti-osterix (Abcam), rabbit anti-osteocalcin (Takara), mouse anti -catenin (BD Biosciences), and rat anti-BrdU (Abcam) were used for immunohistochemical analysis. Secondary antibodies for immunohistochemistry were from Jackson ImmunoResearch. Human PTH(1–34) was purchased from Bachem Bioscience Inc. Adenovirus-GFP (Ad-GFP) and Adenovirus-Cre (Ad-Cre) recombinant adenovirus were purchased from Vector Laboratories.

#### LRP6 conditional null mice and PTH treatment

*Lrp6<sup>f/f</sup>* mice were obtained from Van Andel Research Institute (32;33). Transgenic mice expressing the Cre recombinase under the control of a 3.9-kb fragment of the human osteocalcin promoter (*Oc-Cre*), created in a Friend leukemia virus strain B genetic background, were obtained from T. Clemens (Baltimore, MD) (34). LRP6<sup>f/f</sup> mice were crossed with *Oc-Cre* mice to generate *Oc-Cre*; *Lrp6<sup>f/f</sup>* progeny, which were used in subsequent mating. The animals that had a genotype of  $Cre^{-/-}$ ;*LRP6<sup>f/f</sup>* were considered control (WT), and  $Cre^{+/-}$ ;*LRP6<sup>f/f</sup>* were considered LRP6 deletion (KO). All animals were maintained in the Animal Facility of the Johns Hopkins University School of Medicine. The experimental protocol was reviewed and approved by the Institutional Animal Care and Use Committee of the Johns Hopkins University, Baltimore, MD, USA. Genomic DNA was extracted from tail snips with phenol/chloroform. Genotyping of the animals was achieved by PCR for Cre recombinase (5'-CAAATAGCCCTGGCAGAT-3' and 5'-TGATACAAGGGACATCTTCC-3') and the loxP sites (5'-

GGGGTTCTACTTTTGTGTGTGG -3' and 5'- CCGTCTGTTTGCATAAAGCAACA -3 '). For PTH treatment, two month-old male Cre<sup>-/-</sup>;LRP6<sup>*ff*</sup>(WT) and Cre<sup>+/-</sup>;LRP6<sup>*ff*</sup>(KO) mice were randomized into four groups: WT-Vehicle, WT-PTH, KO-vehicle, and KO-PTH. Ten mice of each treatment group were used. Mice were treated daily, five days per week, with either vehicle (1mM acetic acid in PBS) or human PTH1-34 at a dose of 80 µg/kg with subcutaneous injection for 4 weeks.

#### Analysis of skeletal phenotypes

Mice were anesthetized by inhalation of 2.5% isoflurane (Abbott Laboratories, Abbott Park, IL, USA) mixed with O<sub>2</sub> (1.5 liters/minute). For  $\mu$ CT analysis, femora obtained from mice were dissected free of soft tissue, fixed overnight in 70% ethanol and analyzed by a high resolution  $\mu$ CT (SkyScan1076 in-vivo CT, SKYSCAN company). Image Reconstruction software (NRecon v1.6), data analysis software (CTAn v1.9), and three-dimensional model visualization software (CTVol v2.0) were used to analyze parameters of the trabecular bone in the metaphysis and mid-diaphyseal cortical bone. The scanner was set at a voltage of 50 kVp, a current of 201  $\mu$ A and a resolution of 12.63679  $\mu$ m per pixel. Cross-sectional images of the distal femur were used to perform three-dimensional histomorphometric analysis of trabecular bone. For 1 month-old mice, the sample area selected for scanning was a 2.0-mm length of the metaphyseal trabecular bone immediately subjacent to the growth plate. For 3 month-old mice, the sample area selected for scanning was a 2.0-mm length of the metaphyseal secondary spongiosa, originating 1.0 mm below the lowest point of growth plate (illustrated in Fig. 1E). Cortical morphometry was analyzed within a 600 µm long

section at mid-diaphysis of the femur and included measurements of average thickness and cross-sectional area.

For histochemistry, immunohistochemistry, and histomorphometric analysis, the femora were resected and fixed in phosphate-buffered saline (pH 7.4) containing 4% paraformaldehyde for 48 hours, decalcified in 10% ethylenediamine tetraacetic acid (EDTA) (pH 7.0) for 14 days and embedded in paraffin. Four-µm-thick longitudinally oriented sections of bone including the metaphysis and diaphysis were processed for hematoxylineosin and immunohistochemical staining. For static histomorphometry, measurements of two-dimensional parameters of the trabecular bone were performed with OsteoMeasureXP Software (OsteoMetrics, Inc.). For 1 month-old mice, the sample area selected for calculation was a 1 mm<sup>2</sup> area within the metaphyseal trabecular bone. For 3 month-old mice, the sample area selected for scanning was a 1 mm<sup>2</sup> area within the metaphyseal secondary spongiosa. All sections were observed on Olympus BX51 microscope. For dynamic histomorphometry, two sequential doses of calcein (8 mg/10 ml sterile saline) were injected intraperitoneally at 3 and 10 days prior to sacrifice. Bone histomorphometry analysis in undecalcified femoral sections was performed by OsteoMeasure system (OsteoMetrics, Inc.).

Immunohistochemistry analysis was performed using standard protocol as the manufacturer recommended (EnVision<sup>™</sup> System, Dako, USA). Briefly, the bone sections were processed for antigen retrieval by digestion in 0.05% trypsin (pH 7.8) for 15 minutes at 37°C, and then incubated with antibodies against LRP6 (diluted 1:50), osteocalcin (Ser463/465) (diluted 1:100), osterix (diluted 1:400), BrdU (diluted 1:100), and -catenin (diluted 1:100) overnight at 4 °C. An HRP-streptavidin detection system (Dako) was subsequently used to detect the immunoactivity followed by counterstaining with hematoxylin (Sigma). Sections incubated with 1% nonimmune serum PBS solution served as negative controls.

TUNEL assays were performed using parafin-embedded bone sections. Apoptotic cells were detected using DeadEnd<sup>TM</sup> Colorimetric Apoptosis Detection System (Promega, Madison, WI) according to manufacturer instructions.

#### Measurements of serum osteocalcin and cross-linked C terminal telopeptide of type 1 collagen (CTX-I)

Serum bone formation marker osteocalcin and bone resorption marker CTX-I, were measured using commercial kits: Mouse Osteocalcin EIA Kit (Biomedical technologies, Inc. Stoughton, MA) and Mouse CTX-I ELISA Kit (MyBioSource, Inc., San Diego, CA) according to the manufacturer's instructions.

#### Primary osteoblast culture and adenovirus infection

Calvaria (frontal and parietal bones) was removed from newborn (5~6 day-old) *LRP6*<sup>*I*/*T*</sup> mice aseptically. After removal of sutures, the calvariae were subjected to serial digestions in 1.0 mg/ml of collagenase type IV (Sigma). Briefly, calvaria were digested in 10 ml of digestion solution for 15 minutes at 37°C under constant agitation. The digestion solution was then collected, and the digestion was repeated an additional 4 times. Digestion solutions 3–5, which contained the osteoblasts, were pooled together. After centrifugation at 600g for 10 min, the fractions were resuspended in -Minimum Essential Medium ( -MEM) containing 10% Fetal Calf Serum (FCS), and filtered through a 70-mm cell strainer. Osteoblasts were obtained and cultured in -MEM containing 10% FCS and 1% penicillin/ streptomycin at 37°C in a humidified incubator supplied with 5% CO<sub>2</sub>. To eliminate LRP6 from the cells, monolayer osteoblasts were infected with control adenovirus (Ad-GFP) or

Cre recombinase virus M1 (Ad-CreM1) (Vector Laboratories) at a multiplicity of infection (MOI) of 100 for most experiments.

#### Quantitative Real-Time PCR (qRT-PCR)

Total RNA for qRT-PCR was extracted from bone tissue and cultured cells using Trizol reagent (Invitrogen) according to the manufacturer's protocol. For qRT-PCR, cDNA was prepared with random primers using the SuperScript First-Strand Synthesis System (Invitrogen) and analyzed with SYBR GreenMaster Mix (QIAGEN) in the thermal cycler with two sets of primers specific for each targeted gene. Relative expression was calculated for each gene by 2-CT method with GAPDH for normalization. Primers used for qRT-PCR were: Col1A (5'-GCGAAGGCAACAGTCGCT-3') and (5'-CTTGGTGGTTTTGTATTCGATGAC-3'); Osteocalcin (5'-CTGACCTCACAGATCCCAAGC-3') and (5'-TGGTCTGATAGCTCGTCACAAG-3'); bone sialoprotein (BSP) (5'-TACCGGCCACGCTACTTTCTTTAT-3') and (5'-GACCGCCAGCTCGTTTTCATCC-3'); dentin matrx protein-1 (DMP1) (5'-GGCTGTCCTGTGCTCTCCCAG-3') and (5'-GGTCACTATTTGCCTGTGCCTC-3'); Axin2 (5'-ATGAGTAGCGCCGTGTTAGTG-3') and (5'-GGGCATAGGTTTGGTGGACT-3'); Naked2 (5'-GGGACGACAAGGGTTCCCTA-3') and (5'-AGTGCGTCAATGTTCAAGTGC-3'); bone morphogenetic protein-4 (BMP4) (5'-TTGATACCTGAGACCGGGAAG-3') and (5'-ACATCTGTAGAAGTGTCGCCTC-3'); osteoprotegrin (OPG) (5'-GTGAAGCAGGAGTGCAAC-3') and (5'-GCAAACTGTGTTTCGCTC-3'); receptor activator of nuclear factor kappa-B ligand (RANKL) (5'-TGTACTTTCGAGCGCAGATG-3') and (5'-ACATCCAACCATGAGCCTTC-3'); matrix metalloprotein 13 (MMP13) (5'-GGTCCCAAACGAACTTAACTTAC-3') and (5'-CCTTGAACGTCATCATCAGGAAG-3'); c-JUN (5'-GCCAACATGCTCAGGGAACAGGT-3') and (5'-GCCCCTCAGCCCTGACAGTCTG-3'), c-fos (5'-AATGGTGAAGACCGTGTCAGGA-3') and (5'-CCCTTCGGATTCTCCGTT TCT-3'); and GAPDH (5'-AGGTCGGTGTGAACGGATTTG-3') and (5'-GGGGTCGTTGATGGCAACA-3').

#### Statistical analysis

All data were presented as mean  $\pm$  SEM. For comparison of histomorphometric parameters in WT and KO mice, Student's *t*-test was used. For quantitative analysis of immunostaining data, Student's *t*-test was performed followed by Chi-square Test. Significant level was defined as P < 0.05.

#### Results

#### Bone mass is decreased in adult mice with Oc-Cre-mediated conditional LRP6 deletion

To determine the role of LRP6 in osteoblastic bone formation, we generated mice lacking Lrp6 specifically in mature osteoblasts (Oc-Cre;  $Lrp6^{f/f}$ , named "KO" hereafter) using *Cre*mediated recombination. Transgenic mice expressing the *Cre* recombinase driven by the human osteocalcin promoter were crossed with homozygous mice that expressed *loxP*flanked Lrp6 ( $Lrp6^{f/f}$ ). To confirm that LRP6 deletion in KO mice was specific to osteoblasts, we performed PCR analysis using a combination of primers that specifically detected floxed Lrp6 alleles ( $Lrp6^{f}$ ) and null alleles (Lrp6). Genomic DNA was extracted from skeletal tissues including calvariae, femur, and spine as well as several non-skeletal tissues such as brain, kidney, lung, liver, heart, and muscle from the KO mice. *Cre*-mediated recombination occurred exclusively in tissues that contained osteoblastic cells, whereas nonskeletal tissues retained the intact Lrp6-floxed alleles ( $Lrp6^{f}$ ) (Fig. 1A). LRP6-positive

osteoblasts in femur tissue were also detected by immunohistochemical analysis of tissue sections using an antibody against LRP6. While 58.6% of the bone surface osteoblasts expressed LRP6 in LRP6<sup> $\ell/f$ </sup> mice (named "WT" hereafter), almost none of the osteoblasts expressed LRP6 in KO mice (Fig. 1B and Supplemental Fig. S1). No significant reduction of LRP6-positive cell numbers in bone marrow was observed in KO mice compared to WT mice (Fig. 1C and Supplemental Fig. S1). Thus, LRP6 deficiency was limited to mature osteoblasts on the bone surface. KO mice were born at the expected Mendelian frequency, and all exhibited survival indistinguishable from that of WT mice. Mice expressing cre only did not exhibit any skeletal abnormality relative to wild type mice in deferent studies (34;35) and no abnormal phenotype was observed in our analysis (data not shown). The body weight and femoral length were not decreased in either male or female KO mice compared with WT controls at 1 and 3 months of age (Table 1).

To dissect the role of LRP6 in osteoblast differentiation in bone growth and remodeling, we analyzed the femora bone phenotypes of young (1 month-old) and adult (3 month-old) mice. One month-old KO mice did not exhibit significant differences in trabecular bone volume, number, thickness or separation relative to their WT littermates as measured by  $\mu$ CT (Fig. 1D–1H). However, 3 month-old KO mice showed reduced trabecular bone volume and number, and greater trabecular bone separation in the secondary spongiosa area compard to WT littermates (Fig. 1D–1H). No significant differences were observed for the cortical bone parameters including bone area, bone area/total area, and cortical bone thickness in both 1 and 3 month-old KO mice as compared to their WT littermates (Fig. 1I–1M). The results suggest that LRP6 deficiency in mature osteoblasts primarily affected trabecular bone remodeling in adults.

# Bone formation, but not bone resportion, is impaired in mice with Oc-Cre-mediated conditional LRP6 deletion

Bone histomorphometric analyses of femora revealed no changes in the number of osteoblasts and osteoclasts in 1 month-old KO mice (Fig. 2A-2D). A reduced number of osteoblasts but no change in the numbers of osteoclasts was observed in 3 month-old KO mice as compared to their WT littermates (Fig. 2E-2H). Reduced bone formation in 3 month-old KO mice was also confirmed by Goldner's Trichrome staining and double calcein labeling analysis (Fig. 2I-K). Distal femur showed a decrease in bone volume (green/ *blue*) and newly formed bone (*red*) (Fig. 2I). We analyzed dynamic changes in bone remodeling by injecting calcein twice at 7-day intervals before euthanizing the animals. The distance between two consecutive labels in the trabecular bone of the femur was less in KO mice than in the WT controls (Fig. 2J). Quantification of these parameters demonstrated that KO mice had reduced BRF/BS when compared to WT littermates (Fig. 2K). To investigate if osteoclast bone resorption activities were affected, we performed TRAP staining in the femoral sections of 3 month-old mice. The number of TRAP-positive osteoclasts on the trabecular bone surface of KO mice was unchanged compared to WT mice (Fig. 2L and 2M). We then examined whether -catenin signaling was affected by LRP6 deficiency in osteoblasts. We analyzed -catenin target gene expression in femora bone tissue. Axin2 (36-38), Naked2 (39;40), BMP4 (41) and OPG (42;43) were significantly down-regulated by 30-40% in KO mice relative to WT mice (Fig. 2N-2Q). Of note, RANKL expression was also decreased in KO mice (Fig. 2R). Overall, the ratio of OPG/RANKL was maintained (Fig. 2S) in KO mice compared to WT littermates, which may be the main reason why osteoclastic bone resorption was not affected. Taken together, these results demonstrate that deletion of LRP6 in osteoblasts impaired bone formation without affecting osteoclastic bone resorption during bone remodeling in adult mice.

# Oc-Cre-mediated conditional LRP6 deletion leads to decreased expression of osteoblast differentiation genes and accelerated osteoblast apoptosis during bone remodeling

To explore the cellular mechanism underlying the reduced osteoblastic bone formation observed in the KO mice, we examined osteogenic gene expression in the femoral trabecular bone tissue of 3 month-old LRP6 KO mice. Bone surface osteocalcin<sup>+</sup>- and osterix<sup>+</sup> osteoblast numbers in the second spongiosa of KO mice were significantly reduced relative to WT mice (Fig. 3A, 3B, Supplemental Fig. S2A, and S2B). Consistently, the number of osteocytes, representing terminal differentiation of the osteoblast lineage, was also significantly reduced in the trabecular bone matrix of KO mice relative to their WT littermates (Fig. 3C and Supplemental Fig. S2C). The mRNA expression levels as assessed by qRT-PCR of bone matrix proteins produced by mature osteoblasts (Col1A, osteocalcin, and BSP) and osteocytes (DMP1) were also decreased in bone tissue from 3 month-old KO mice (Fig. 3E). The expression levels of these genes were not changed in 1 month-old mice compared to their age-matched WT controls (Fig. 3D). We next sought to determine whether deletion of LRP6 in osteoblasts affected proliferation and/or apoptosis of osteoblasts in the secondary spongiosa area during bone remodeling. Analysis of BrdU labeling revealed that the proliferation of bone surface osteoblasts and bone matrix osteocytes in the second spongiosa of KO mice was not different from that of WT mice (Fig. 3F and Supplemental Fig. S2D). However, the percentage of apoptotic osteoblasts on the bone surface was greater in LRP6 KO mice relative to their WT littermates as detected by TUNEL assay (Fig. 3G and Supplemental Fig. S2E). In the bone marrow of the secondary spongiosa, TUNEL<sup>+</sup> cells (Fig. 3H and Supplemental Fig. S2E) and Osterix<sup>+</sup> osteoprogenitors (Fig. 3I and Supplemental Fig. S2B) were not changed in KO mice relative to their WT littermates. These results indicate that LRP6 deficiency in osteoblasts resulted from increased apoptosis, decreasing the number of bone surface osteoblasts, but did not affect the pool of osteoblastsderiving osteoprogenitors during bone remodeling.

# Oc-Cre-mediated LRP6 deficient mice exhibit a blunted response in bone formation to PTH treatment

To determine whether LRP6 in osteoblasts is required for PTH anabolic effects on bone, 2 month-old male KO mice and littermate WT controls were treated with PTH1-34 for 4 weeks. Femora trabecular and cortical bone was analyzed by  $\mu$ CT and histomorphormetry. In WT mice, PTH increased bone mass in trabecular and cortical bone (Fig. 4A and Supplemental Fig. S3). Particularly, PTH1-34 increased BV/TV (Fig. 4B), BMD (Fig. 4C), Tb.N (Fig. 4D), Tb.Th (Fig. 4F), and decreased Tb.Sp (Fig. 4E) in WT mice compared to the vehicle-treated WT mice. However, these PTH-induced changes were not seen in KO mice and actually resulted in a decreased BMD compared to the vehicle-treated KO mice. Goldner's Trichrome staining revealed increased newly formed bone (red) with PTH1-34 treatment only in WT mice, not in KO mice compared to their respective vehicle-treated controls (Fig. 4G). The capacity of PTH1-34 treatment to increase the amount of bone deposited between two timed administrations of calcein labeling, an indicator of bone mineralization in WT mice, was diminished in KO mice (Fig. 4H and 4I). Quantification of these calcein intervals revealed that the in absence of LRP6, PTH1-34 treatment did not change the mineral apposition rate (MAR) in KO mice, whereas the MAR was increased in the PTH1–34-treated WT mice compared to their respective vehicle-treated controls (Fig. 4J), suggesting decreased activity of individual osteoblasts. Analyses of dynamic parameters of bone formation further indicated that PTH1-34 increased the bone formation rate over the bone surface (BFR/BS), which reflects the quantity of osteoblasts forming the bone matrix, in WT mice but not in KO mice (Fig. 4K). Thus, the PTH anabolic effects were abrogated by LRP6 deletion in osteoblasts.

#### Oc-Cre-mediated LRP6 deficient mice exhibit blunted osteoblast differentiation and antiapoptotic responses to PTH

Given that the anabolic effect of PTH results from a tight balance between osteoblastic bone formation and osteoclastic bone resorption (7;44), we then investigated whether LRP6 deficiency impairs PTH-stimulated bone formation only or both osteoblastic bone formation and osteoclastic bone resorption. Analysis of the secondary spongiosa by bone histomorphometry revealed that the two static indices of bone formation stimulated by PTH, the number of osteoblasts per bone surface (N.Ob/B.Pm) (Fig. 5A) and the percentage of surfaces covered by osteoblasts (Ob.S/BS) (Fig. 5B), were only observed in WT mice but not in KO mice. On the contrary, the two indices of trabecular bone resportion, the number of osteoclasts per bone surface (N. Oc/B. Pm) (Fig. 5C) and the percentage of surfaces covered by osteoclasts (Oc.S/BS) (Fig. 5D), increased by a similar extent in WT and KO mice in response to PTH1-34 treatment compared to their respective vehicle-treated controls. Consistently, Oc-Cre-mediated LRP6 deficiency blunted the increase in number of osteocalcin<sup>+</sup> osteoblasts on the bone surface (Fig. 5E and 5F) but did not change the effect of increased number of TRAP<sup>+</sup> osteoclasts stimulated by PTH1-34 in KO mice (Fig. 5G and Fig. 5H). We also measured serum concentrations of osteocalcin and CTX-I in response to PTH1-34 treatment. The levels of both serum osteocalcin and CTX-I in WT and KO mice increased significantly in the PTH-treated groups compared to the vehicle-treated groups. However, the serum osteocalcin concentration was not stimulated in PTH-treated KO mice as significantly as in PTH-treated WT mice, whereas the PTH-stimulated increase in serum CTX-I concentration was similar in the KO and WT mice (Fig. 5I and 5J). Thus, LRP6mediated signaling in mature osteoblasts is important for osteoblast differentiation during bone remodeling and PTH-stimulated bone formation.

We next examined whether LRP6 deletion in osteoblasts affects the number of osterixpositive cells, which are osteoblast-deriving osteoprogenitors, by immunohistochemical analysis. In WT mice, PTH significantly increased osterix<sup>+</sup> osteoprogenitors in both bone marrow and on the bone surface (Fig. 6A-6C). On the contrary, in KO mice, although PTH was able to increase the number of osterix<sup>+</sup> osteoprogenitors in the bone marrow, PTH failed to increase the number of bone surface osterix<sup>+</sup> cells compared to vehicle treatment (Fig. 6A-6C), indicating that deletion of LRP6 in osteoblasts does not affect the pool of osteoblast-deriving bone marrow osteoprogenitors but likely impairs the recruitment, terminal differentiation and/or survival. As an increased number of apoptotic osteoblasts on the bone surface was observed in KO mice (Fig. 3G and Supplemental Fig. S2E) and PTH has previously been shown to have anti-apoptotic effects, we examined whether LRP6 deletion in osteoblasts affects PTH-inhibition of osteoblast apoptosis. The percentage of apoptotic osteoblasts on bone surface was reduced by PTH treatment in WT mice, consistent with previous reports (7;45;46). However, the anti-apoptotic effect of PTH was abrogated by LRP6 deletion (Fig. 6D and 6E). These results suggest that LRP6 is required for osteoblast survival in PTH-stimulated bone remodeling.

#### PTH failed to stimulate β-catenin and Gα<sub>s</sub>-PKA signaling in LRP6-deficient osteoblasts

Finally, we sought to determine whether PTH-activated signaling pathways are affected by the deletion of LRP6 in osteoblasts. We previously found that LRP6 is essential for PTH stimulated -catenin signaling, a positive regulator in bone formation, in osteoblasts (24). We tested whether LRP6 deficiency would affect -catenin signaling activation induced by PTH in osteoblasts. Consistent with our previous report (24), PTH1–34 significantly increased -catenin<sup>+</sup> osteoblasts on bone surface in secondary spongiosa of WT mice (Fig. 7A and 7B). -catenin<sup>+</sup> osteoblasts on bone surface in the same area in KO mice did not increase in response to PTH1–34 treatment compared to vehicle-treated controls (Fig. 7A and 7B). In addition, the PTH-stimulated increased expression of -catenin target genes

Naked2 and Axin1 was reduced in LRP6-deficient PTH-treated calvarial osteoblasts compared with those in WT calvarial osteoblasts (Fig. 7C). We previously also demonstrated that LRP6 mediates PTH-activated G  $_{\rm s}$ -PKA signaling activation in osteoblasts (27). We detected the mRNA expression of c-JUN, c-fos (47–49), and MMP13 (50;51), genes stimulated by PTH through G  $_{\rm s}$ -PKA-dependent pathway. LRP6-deficiency in osteoblasts reduced the relative abundance of PTH-stimulated c-fos, RANKL, and MMP13 gene expression in isolated cavarial preosteoblasts compared to PTH-treated WT osteoblasts (Fig. 7C). The results suggest that Oc-Cre mediated conditional LRP6 deletion alleviated both -catenin signaling and G  $_{\rm s}$ -PKA signaling activation stimulated by PTH in osteoblasts.

#### Discussion

Mutations in both LRP5 and LRP6 have been linked to changes in bone mass in humans and mice, but the role of LRP6 in bone metabolism has not been studied as extensively relative to that of LRP5. In addition to its role in wnt-canonical pathway signaling, LRP6 is also required for PTH-stimulated signaling pathways in osteoblasts (24;27). Whether or not LRP6 mediates PTH anabolic bone effects *in vivo* has not been investigated. Here we demonstrate that LRP6 deficiency in mature osteoblasts decreased bone formation in the secondary spongiosa area in 3 month-old adult mice that have active bone remodeling, but had marginal effects on bone acquisition in young mice. Further, the body weight and femur length in 1 month-old LRP6-deficient mice of both sexes were not reduced in comparison with those of wild type littermates. Thus, osteoblast-specific LRP6 is required for bone formation specifically during bone remodeling, and is most likely not critical in bone development and postnatal growth. In addition, the anabolic effect of intermittent PTH administration during bone remodeling was abolished in osteoblast-specific LRP6-deficient mice. These in vivo evidence and the previous finding that intermittent PTH treatment was fully anabolic in LRP5-deficient mice (13;14) consistently support the concept that LRP6, but not LRP5, is an essential mediator for PTH-elicited bone anabolic effect during bone remodeling.

Bone remodeling is mediated by tight coupling of osteoclastic bone resorption on the bone surface and osteoblastic new bone formation at the same sites. In the coupling process, local factors released and activated during osteoclastic bone resorption induce the migration of MSCs/osteoprogenitors to the remodeling bone surface where they differentiate into mature osteoblasts to form new bone (1-3). Our results show that LRP6 is required for osteoblast differentiation and survival during the recruitment of osteoblasts in the remodeling process. Indeed, mRNA levels of the osteoblast differentation markers were reduced in bone tissue of 3 month-old LRP6-deficient mice, whereas the levels of these genes were maintained in 1 month-old LRP6-deficient mice. Further analysis of the cellular changes in the secondary spongiosa of the femur of LRP6 KO mice revealed a dramatically reduced number of osteoblasts and increased number of apoptotic osteoblasts on the bone surface, indicating that LRP6 is necessary for the survival of osteoblastic cells during bone remodeling. PTH produces osteoanabolic activity via anti-apoptotic mechanisms in osteoblasts, and increased cell survival is one of the major contributors to the increased number of osteoblasts caused by PTH (7). Our demonstration that deletion of LRP6 abolished the anti-apoptotic effect of PTH on osteoblasts suggests that LRP6 is involved in PTH anti-apoptotic effect in osteoblasts. While the molecular mechanisms by which LRP6 regulates osteoblast apoptosis remain to be determined, it has been reported that the attenuation of osteoblast apoptosis by PTH requires PKA-mediated phosphorylation (7;45;46). Our results showed that the expression of PKA downstream target genes stimulated by PTH such as c-fos, RANKL, and MMP13 were abolished in LRP6-deficient osteoblasts, indicating that LRP6-deficiency may promote osteoblast apoptosis through disruption of PTH-stimulated PKA signaling.

Interestingly, LRP6 deletion did not affect osteoblast differentiation and bone formation during bone development and growth. LRP6, containing a large extracellular domain, has been proposed to act as a central organizer for the extracellular antagonist network for the regulation of signaling pathways of different local growth factors (26). Different types and levels of extracellular factors produced may differentially influence the effect of LRP6 in osteoblasts. The extracellular bone marrow microenvironment in bone growth is different from that of bone remodeling, which involves osteoclast bone resorption and release of matrix factors. Mice with LRP6 deletion in early mesenchyme exhibited relatively normal skeleton (33), providing additional evidence for the dispensable role of LRP6 in bone formation during development.

LRP6 deletion in osteoblasts did not affect osteoclastogenesis as indicated by normal osteoclast numbers, normal TRAP+ cell numbers on bone surface, and normal serum CTX-I concentrations relative to wild type mice. On the other hand, our results from the current study strongly support our previous finding that LRP6 is a key mediator for PTH activated--catenin signaling in osteoblasts (24) as deletion of LRP6 resulted in reduced expression of downstream target genes of -catenin signaling in bone tissue. It is known that deletion of catenin within mature osteoblasts causes a decrease in OPG production and an increase in osteoclast activity (42;43;52). Consistently, we found LRP6 KO mice to have decreased expression of OPG. It is also known that RANKL is a direct target gene of PKA signaling in osteoblasts (53-55). We found that PTH lost its ability to stimulate PKA signaling in osteoblasts in LRP6 KO mice, which may be a major contributor to our observation of reduced RANKL production in osteoblasts. Thus, downregulation of OPG and RANKL separately caused by suppressed -catenin and PKA signaling, may reach a balance to maintain the osteoclast activity in LRP6 KO mice and explain the normal ratio we observed of RANKL/OPG compared to WT controls. This finding provides one possible explanation for the different bone phenotypes derived from the -catenin-deficiency (42;43;52) and our findings with LRP6-deficiency. Our results also suggest that LRP6 is required for both PTHstimulated -catenin and PKA signaling, which most likely play different roles in regulating osteoblast and osteoclast activities. PTH directly stimulates PKA signaling activation, which subsequently phosphorylates transcription factors, such as Cbfa1 and cAMP-response element-binding protein (CREB), regulating the transcription of many genes and multiple cellular activities including osteoblast differentiation, survival/apoptosis, cell cycle, proliferation, and osteoclast activity (7;56;57). PTH can also activate -catenin signaling in osteoblasts and thereby regulate the activity of osteoclasts. Additionally, PTH stimulates many other signaling pathways such as PKC, MAPK, etc. It remains to be determined whether LRP6 is involved in the activation of these signaling pathways in osteoblasts. In addition to PTH signaling, LRP6 may also be involved in other growth factor/cytokinestimulated RANKL production in osteoblasts, which is another interesting topic for future study. Nevertheless, we have found that LRP6 in osteoblasts is not only a mediator for catenin signaling activation, but regulates the functions of osteoblasts and osteoclasts through distinct PTH-signaling mechanisms.

One of the functions of PTH is to promote the commitment and differentiation of MSCs or osteoprogenitors into mature osteoblasts (7;58;59). BMP-induced MSC commitment to osteoblastic lineage was suppressed by LRP6 via extracellular antagonist network, which can be disrupted by PTH-induced endocytosis of the PTH1R-LRP6 complex, resulting in increased commitment of MSCs to the osteoblastic lineage (26). Further characterization of the role of MSC- or osteoprogenitor-specific LRP6 mediated signaling in bone remodeling using nestin-Cre or osterix-Cre mouse models are needed. Whether PTH still stimulates bone formation in these mouse models is also worth testing. MSCs are one of the major constituents of the bone marrow microenvironment that supports hematopoietic stem cells (58–60). It is known that intermittent PTH treatment expands hematopoietic stem and

progenitor cells (61;62). MSCs have been shown to express PTH receptors and to increase in number and undergo redistribution in bone marrow upon PTH treatment. Therefore, examining whether LRP6 plays a role in PTH-regulated stem cell behavior in bone marrow is another interesting topic for future study.

In conclusion, our studies demonstrate that LRP6 in osteoblasts is a positive regulator for osteoblastic bone formation during bone remodeling and plays a critical role in the osteoanabolic action of PTH.

#### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Fig. 1. Osteoblast-specific deletion of LRP6 causes bone loss in trabecular bone in adult but not in young mice

(A) PCR analysis was performed using different tissues harvested from LRP6 KO mice to examine specific deletion of *Lrp6*. Bands represent *Lrp6* gene alleles that are either flanked with a loxP site (*Lrp6<sup>f</sup>*) or deleted (*Lrp*). (B and C) Number of LRP6<sup>+</sup> cells on bone surface (B) and in bone marrow (C). N. LRP6<sup>+</sup> cells/B.Pm, number of LRP6-positive cells per bone perimeter. N. LRP6<sup>+</sup> cells/T.Ar, number of LRP6-positive cells per tissue area. n = 7; \*p < 0.001. Data are presented as mean  $\pm$  SEM. (D–H) Representative  $\mu$ CT images and quantitative analysis of the trabecular bone area of distal femur from 1 and 3 month-old male mice. Yellow squares indicate the area chosen for analysis. Scale bar: 500 $\mu$ M.

Trabecular bone volume fraction (BV/TV) (**E**), trabecular number (Tb.N) (**F**), trabecular thickness (Tb.Th) (**G**), and trabecular separation (Tb. Sp) (**H**). Light blue column, WT; Dark blue column, KO. n = 10; \*p < 0.05. (**I–M**) Representative  $\mu$ CT images and quantitative analysis of cross-sections of femoral mid-diaphyses from 1 and 3 month-old male mice. Scale bar: 500 $\mu$ M. Cortical bone area (BA) (**J**), total area with the periosteal circumference (TA) (**K**), cortical bone area as a percentage of total area within the periosteal circumference (BA/TA) (**L**), and cortical bone thickness (Co.Th) (**M**). n = 10. Light blue column, WT mice; Dark blue column, KO mice. Data are presented as mean ± SEM.

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Fig. 2. Osteoblast-specific deletion of LRP6 causes decreased osteoblastic bone formation without affecting osteoclastic bone resorption in adult mice femora

(A–D) Bone histomorphometric analysis of trabecular bone from 1 month-old male mice. Number of osteoblasts per bone perimeter (N.Ob/B. Pm) (A), osteoblast surface per bone surface (Ob.S/BS) (B), number of osteoclasts per bone perimeter (N.Oc/B. Pm) (C), and osteoclast surface per bone surface (Oc.S/BS) (D). n = 8. (E–H) Bone histomorphometric analysis of trabecular bone of the secondary spongiosa from 3-month-old male mice. Number of osteoblasts per bone perimeter (N.Ob/ B. Pm) (E), osteoblast surface per bone surface (Ob.S/BS) (F), number of osteoclasts per bone perimeter (N.Oc/B. Pm) (G), and osteoclast surface per bone surface (Oc.S/BS) (H). n = 8. \*p < 0.001. (I) Trichrome staining

of the metaphyseal trabecular bone at distal femora of 3 month-old mice. Mineralized bone stains green/blue and osteoid stains red. Upper panels are lower power image with boxes outlining the area of higher power in bottom panels. Scale bars:  $100\mu$ M. (**J**–**K**) Representative calcein double labeling and quantification of bone formation rate of the metaphyseal trabecular bone at distal femora. Scale bar:  $100\mu$ M (**J**). Bone formation rate per bone surface (BFR/BS (**K**)). n = 6. \*p < 0.01. (**L**–**M**) Representative tartrate-resistant acid phophatase (TRAP) staining and quantitative analysis of trabecular bone sections from distal femur. Scale bars:  $100\mu$ M (**L**). Number of TRAP-positive cells per mm<sup>2</sup> tissue area (N.TRAP<sup>+</sup> cells/T.Ar) (**M**). n = 7. (**N**) Quantitative RT-PCR analysis of -catenin target genes (Axin 2, Naked2, BMP4, and OPG) and RANKL expression in bone tissue extracts from 3 month-old mice. Light blue column, WT; Dark blue column, KO. (**O**) Ratio of RANKL/OPG in bone tissue extracts from 3 month-old mice. n = 8; \*p < 0.05. Data are presented as mean  $\pm$  SEM.





(A–B) Quantification of immunohistochemical staining for osteocalcin (Ocn) (A) and osterix (Osx) (B) in femur sections of 3 month-old male mice. Number of Ocn-positive cells per mm<sup>2</sup> tissue area (N.Ocn<sup>+</sup> cells/T.Ar), number of Osx-positive cells per mm<sup>2</sup> tissue area (N.Osx<sup>+</sup> cells/T.Ar). n = 8; \*p < 0.01, \*\*p < 0.001.. Data are presented as mean ± SEM. (C) Number of osteocytes in bone matrix. Number of osteocytes per mm<sup>2</sup> bone area (N.osteocytes/B.Ar), n = 8; \*p < 0.05. (D–E) Quantitative RT-PCR analysis of osteoblast marker genes in bone tissue extract from 1 month- (D) and 3 month-old (E) WT and KO mice. Collagen 1A (Col1A), osteocalcin (Ocn), bone sialoprotein (BSP), dentin matrix

protein-1 (DMP1). n = 7; \*p < 0.05. (F) Percentage of BrdU<sup>+</sup> osteoblasts out of total bone surface osteoblasts in the secondary spongiosa area in femur sections of 3-month-old mice. n = 8 (G–H) Analysis of apoptotic cells in trabecular bone (G) and bone marrow (H) of femur from 3 month-old WT and KO mice by TUNEL assay. Data presented as percentage of apoptotic osteoblasts out of total osteoblasts on bone surface (G) and number of TUNEL-positive cells per mm<sup>2</sup> tissue area (N.TUNEL<sup>+</sup>/T.Ar) (H). n = 6. \*p < 0.001. (I) Quantitative analysis of immunohistochemical staining for osterix (Osx) in secondary spongiosa bone marrow of femur sections in 3 month-old male mice. Number of Osx-positive cells per mm<sup>2</sup> tissue area (N.Osx<sup>+</sup> cells/T.Ar), n = 8; Data are presented as mean ± SEM.



Fig. 4. Intermittent PTH treatment failed to increase bone formation in mice with osteoblast-specific deletion of *LRP6* 

(A–F) Representative  $\mu$ CT images and quantitative analysis of distal femur from 3 monthold male mice treated with vehicle or PTH1–34 (80 $\mu$ g/kg. b.w, five days a week for four weeks). Scale bar: 500 $\mu$ M (A). Trabecular bone volume fraction (BV/TV) (B), bone mineral density (BMD) (C), trabecular number (Tb.N) (D), trabecular separation (Tb. Sp) (E), and trabecular thickness (Tb.Th) (F). n = 10 for each treatment group; \*p < 0.05. \*\*p < 0.01. (G) Trichrome staining of metaphyseal trabecular bone at distal femora from 3 month-old male mice treated with vehicle or PTH1–34 (80 $\mu$ g/kg. b.w, five days a week for four weeks). Mineralized bone stains green/blue and osteoid that lines bone stains red. Scale bar: 100 $\mu$ M.

(**H** and **I**) Calcein double labeling of cortical (**H**) and metaphyseal trabecular bone (**I**) in distal femora. Scale bars: 10 $\mu$ M for (**H**), and 100 $\mu$ M for (**I**). (**J**–**K**) Mineral apposition rate (MAR) (**J**) and bone formation rate per bone surface (BFR/BS) (**K**). n = 6; \*p < 0.05, \*\*p < 0.001. (**L**–**M**) Serum levels of osteocalcin (Ocn) (**L**) and c-terminal telopeptide of type-1 collagen (CTX-I) (**M**) in vehicle- or PTH-treated WT and KO mice. n = 7; \*p < 0.001. Data are presented as mean ± SEM.



Fig. 5. LRP6 in osteoblasts is essential for PTH-stimulated osteoblastic bone formation but not osteoclastic bone resorption

(A–D) Bone histomorphometric analysis of trabecular bone in the secondary spongiosa of 3month-old male mice treated with vehicle or PTH1–34 (80µg/kg. b.w, five days a week for four weeks). Number of osteoblasts per bone surface (N.Ob/B. Pm) (A), osteoblast surface per bone surface (Ob.S/BS) (B), number of osteoclasts per bone surface (N.Oc/ B. Pm) (C), and osteoclast surface per bone surface (Oc.S/BS) (D). n = 8; \*p < 0.001. (E–H) Representative immunohistochemical staining and quantitative analysis of osteocalcin (Ocn) (E–F) and tartrate resistant acid phosphatase (TRAP) (H-I) in femur sections of 3 month-old male mice treated with vehicle or PTH(1–34) (80µg/kg. b.w, five days a week for four

weeks). Scale bars:  $100\mu$ M (**E** and **H**). Number of osteocalcin-positive cells per mm<sup>2</sup> tissue area (N.Ocn<sup>+</sup> cells/T.Ar.) (**F**), number of TRAP-positive cells per mm<sup>2</sup> tissue area (N.TRAP<sup>+</sup> cells/T.Ar.) (**H**). n = 7; \*\*p < 0.05, \*p < 0.001. (**I-J**) Serum levels of osteocalcin (Ocn) (**I**) and c-terminal telopeptide of type-1 collagen (CTX-I) (**J**) in vehicle- or PTH-treated WT and KO mice. n = 7; \*p < 0.001. Data are presented as mean ± SEM.

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### Fig. 6. LRP6 deficiency in osteoblasts impairs PTH-stimulated anti-apoptotic effects on osteoblasts without affecting PTH-increased bone marrow osteoprogenitors

(A–C) Representative immunohistochemical staining and quantitative analysis of osterix (Osx) in femur sections of 3 month-old male mice treated with vehicle or PTH1–34 (80µg/kg. b.w, five days a week for four weeks). Scale bars:  $100\mu$ M (A). Number of Osx-positive cells per mm<sup>2</sup> tissue area. (N.Osx<sup>+</sup> cells/T.Ar) in bone marrow (B) and on bone surface (C). (D–E) Representative immunohistochemical staining and quantitative analysis of TUNEL<sup>+</sup> cells in femur sections of 3 month-old male mice treated with vehicle or PTH1–34 (80µg/kg. b.w, five days a week for four weeks). Black arrow: TUNEL<sup>+</sup> osteoblasts, red arrow: TUNEL osteoblasts. Scale bars: 100µM (D). Percentage of TUNEL<sup>+</sup> osteoblasts out of the total osteoblasts on bone surface (E). Red boxes in upper panels of A and D represent higher

magnification shown in lower panels, respectively. n = 7 for each treatment group; \*p < 0.05, \*\*p < 0.001. Data are presented as mean  $\pm$  SEM.



**Fig. 7. PTH** failed to stimulate -catenin and G s-PKA signaling in LRP6-deficient osteoblasts (**A–B**) Representative immunohistochemical staining and quantitative analysis of -catenin in femur sections of 3 month-old male mice treated with vehicle or PTH1–34 (80µg/kg. b.w, five days a week for four weeks). Arrows indicate -catenin<sup>+</sup> osteoblasts on bone surface. Scale bars: 100µM (**A**). Number of -catenin-positive cells per mm<sup>2</sup> tissue area (N. - catenin<sup>+</sup> cells/T.Ar) (**B**). n = 7 for each treatment group; \*p < 0.001. (**C**) Quantitative RT-PCR analysis of G s-PKA target genes (c-JUN, c-Fos, and MMP13) and -catenin target genes (Naked2 and Axin2) in calvarial osteoblasts isolated from  $Lrp6^{f/f}$  mice infected with adenovirus carrying either GFP (WT) or Cre (KO). Cells were treated with osteogenic medium in combination with either vehicle or PTH. n = 8; \*p < 0.05. Data are presented as mean ±SEM.

# Table 1

# Baseline skeletal phenotype in 1 month-old and 3 month-old mice

Values presented are mean±SEM

		Male			Female	
1 month-old mice	WT (n=10)	KO (n=10)	P value	WT (n=10)	KO (n=10)	P value
Femoral length (mm)	13.41±0.90	$13.79 \pm 0.99$	0.34	$11.56\pm0.84$	$11.89 \pm 0.88$	0.48
Body mass (g)	18.24±3.33	20.24±3.27	0.15	14.58±1.68	15.39±2.28	0.41
3 month-old mice		Male			Female	
	WT (n=10)	KO (n=10)	P value	WT (n=10)	KO (n=10)	P value
Femoral length (mm)	15.95±0.75	15.62±0.75	0.5	$14.80 \pm 0.28$	$15.00 \pm 0.50$	0.50
Body mass (g)	28.32±1.77	29.83±1.96	0.25	19.60±1.82	20.20±0.97	0.61