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Authors

Dole, Neha S

Yee, Cristal S

Mazur, Courtney M

et al.

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TGF β regulation of perilacunar/canalicular remodeling is sexually dimorphic

Neha S. Dole¹, Cristal S. Yee¹, Courtney M. Mazur^{1,2}, Claire Acevedo³, Tamara Alliston¹

¹Department of Orthopaedic Surgery, University of California San Francisco, CA

²UC Berkeley-UCSF Graduate Program in Bioengineering, San Francisco, CA

³Department of Mechanical Engineering, University of Utah, Salt Lake City, UT

Abstract

Bone fragility is the product of defects in bone mass and bone quality, both of which show sex-specific differences. Despite this, the cellular and molecular mechanisms underpinning the sexually dimorphic control of bone quality remain unclear, limiting our ability to effectively prevent fractures, especially in postmenopausal osteoporosis. Recently, using male mice, we found that systemic or osteocyte-intrinsic inhibition of TGF β signaling, achieved using the 9.6-kb DMP1 promoter-driven Cre recombinase (T β RII^{ocy-/-} mice), suppresses osteocyte perilacunar/canalicular remodeling (PLR) and compromises bone quality. Since systemic TGF β inhibition more robustly increases bone mass in female than male mice, we postulated that sex-specific differences in bone quality could likewise result, in part, from dimorphic regulation of PLR by TGF β . Moreover, since lactation induces PLR, we examined the effect of TGF β inhibition on the female skeleton during lactation. In contrast to males, female mice that possess an osteocyte-intrinsic defect in TGF β signaling were protected from TGF β -dependent defects in PLR and bone quality. The expression of requisite PLR enzymes, the lacuno- canalicular network, and the flexural strength of female T β RII^{ocy-/-} bone was intact. With lactation, however, bone loss, and induction in PLR and osteocytic parathyroid hormone type I receptor (PTH1R) expression, were suppressed in T β RII^{ocy-/-} bone, relative to the control littermates. Indeed, differential control of PTH1R expression, by TGF β and other factors, may contribute to dimorphism in PLR regulation in male and female T β RII^{ocy-/-} mice. These findings provide key insights into the sex-based differences in osteocyte PLR that underlie bone quality and highlight TGF β signaling as a crucial regulator of lactation-induced PLR.

2. Introduction

Evolution selects for skeletal adaptations that improve survival and reproductive fitness. A prime example is the ability of female mammals to release stored calcium from bone

Corresponding Author: Tamara Alliston, PhD, University of California, San Francisco, Department of Orthopaedic Surgery, 513 Parnassus Avenue, S1155 San Francisco, CA 949041, Phone number-415-502-6523, Tamara.Alliston@ucsf.edu.

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to support lactation. In spite of a sharp drop in bone mass during lactation, the female skeleton mechanically adapts to accommodate these metabolic demands⁽¹⁻⁵⁾. Changes in bone quality, including trabecular microarchitecture and cortical bone geometry, compensate for the loss of bone mass. Other aspects of bone quality, including the material properties of bone extracellular matrix (ECM), show sexual dimorphism^(6,7). Through mechanisms that remain unclear, the protective advantage of such skeletal adaptations for reproductive females is lost later in life. Women over 50 have a 4-fold higher rate of osteoporosis relative to men of the same age^(8,9). Furthermore, about half of osteoporotic fractures cannot be explained by clinically low bone mass^(10,11). Therefore, understanding the sexually dimorphic regulation of bone quality will elucidate mechanisms of bone fragility in post-menopausal osteoporosis, which affects over 200 million people worldwide^(12,13).

Skeletal variation in men and women arises in part from endocrine differences in hormones such as androgens, estrogens, and inhibin⁽¹⁴⁾. These hormones act in concert with paracrine growth factors to control bone cell function^(15,16). Osteoblasts and osteoclasts also exhibit sexual dimorphism at the cell-intrinsic level⁽¹⁷⁻²⁰⁾. Osteocytes respond to high levels of PTH/PTHrP during lactation by inducing perilacunar/canalicular remodeling (PLR)^(21,22). In PLR, osteocytes resorb, and later replace, bone matrix surrounding their intricate lacunocanalicular network^(21,23-25). Originally termed osteocyte osteolysis, PLR supports the metabolic demand for calcium during lactation through coordinated induction of several key genes, including matrix metalloproteinases (Mmps; namely Mmp2, Mmp13, and Mmp14), cathepsin K (Ctsk), carbonic anhydrase 2, and tartrate-resistant acid phosphatase (Acp5/TRAP) to mediate resorption^(23,24,26-28). After weaning, osteocytes replace the bone lost during lactation⁽²⁹⁾, although the underlying molecular mechanism remains elusive. It is possible that genes such as soluble cytosolic phosphatase (Phospho1) and dentin matrix protein (Dmp1)⁽³⁰⁻³²⁾, both of which are implicated in PLR, participate in replenishing the mineralized bone. Until recently, PLR was primarily considered a response to metabolic stress or disease. Work from our group and others reveals PLR to be crucial in homeostasis as well, where it regulates bone quality through TGF β signaling⁽³³⁾. However, the extent to which the sexually dimorphic control of PLR extends dimorphism in bone quality remains unknown.

TGF β is a key driver of bone mass and quality that dictates activities of all of the cells of the bone remodeling unit. Indeed, dysregulated TGF β signaling plays a causal role in several skeletal diseases, including Camurati Engelman disease and osteogenesis imperfecta^(34,35). This key cytokine, which is largely produced by osteoblasts and osteocytes, is embedded within the bone ECM in a latent form⁽³⁶⁻³⁸⁾. Upon osteoclastic resorption, TGF β is released and activated to promote recruitment and commitment of osteoblast progenitors to the resorption site, while inhibiting their terminal differentiation. TGF β also promotes osteoclast formation and survival and couples bone resorption with formation⁽³⁹⁾. In osteocytes, we found that TGF β signaling is indispensable for PLR^(33,40). Using pharmacologic TGF β antagonists and a novel mouse model, we demonstrated that osteocyte-intrinsic ablation of TGF β signaling is sufficient to suppress PLR and cause bone fragility due to severely compromised bone quality⁽³³⁾. Although female mice accrue more bone mass in response to pharmacologic TGF β antagonists than males⁽³⁷⁾, the sexually dimorphic regulation of PLR or bone quality by TGF β has not been evaluated.

Therefore, we used mice with an osteocyte-specific deletion of TGF β receptor II (T β RII) to investigate the sexually dimorphic regulation of PLR in mice at homeostasis and during the metabolic stress of lactation. Our findings reveal a dual role for TGF β in the control of PLR in female bone. While TGF β is required for lactation-induced PLR, at homeostasis, female bone is insensitive to loss of osteocytic TGF β signaling. Here we elucidate a cellular mechanism that contributes to the dimorphic control of PLR in males and females, a mechanism that protects bone quality in reproductive age females.

3. Material and Methods

3.1 Mice.

The generation of viable osteocyte specific T β RII^{ocy-/-} mice was described previously⁽³³⁾. In brief, homozygous TGF β type II receptor (T β RII)-floxed mice were bred with the hemizygous 9.6-kb DMP1-Cre mice^(41,42). Half of the mice resulting from the cross were DMP1-Cre Tg+; T β RII^{flox/flox} (labeled T β RII^{ocy-/-} mice) and half were littermates lacking the DMP1-Cre transgene (T β RII^{flox/flox}) served as controls (labeled control mice). All the transgenic mouse lines were backcrossed for 8 generations into a congenic C57BL/6 background. In all the experiments 15-week old male and female T β RII^{ocy-/-} and control mice were used. For lactation studies, T β RII^{ocy-/-} mice and control mice were time-mated at 10 weeks of age. Females that did not become pregnant after 1 week were removed from the study to minimize age-related differences in the bone phenotype. Following delivery, for each lactating mother the litter size was adjusted to 8 pups in order to normalize for breast milk requirements. Mice were sacrificed after 7 days of lactation. For all experiments, we used control mice that were true littermates for the mutant mice. All studies were conducted with the approval of the Institutional Animal Care and Use Committee of the University of California San Francisco. Additional details on mice are provided in the Supplemental data.

3.2 Micro-computed tomography (micro-CT).

For skeletal phenotyping, 15-week-old male and female mouse bones were harvested by cleaning off the surrounding soft tissue. Cleaned bones were subsequently fixed in 10% neutral buffered formalin and stored in 70% ethanol. A 2 mm region of the femoral metaphysis and a 1 mm region of the femoral mid-diaphysis were scanned using a Scanco μ CT50 specimen scanner to assess trabecular and cortical parameters, respectively. Scanning was performed with an X-ray potential of 55 kVp and current of 109 μ A and at a resolution of 10 μ m. Table 1 shows standard parameters for n = 8 mice per group⁽⁴³⁾. Additional details of micro-CT procedure are provided in the Supplemental data.

3.3 Quantitative RT-PCR analysis.

As previously described, RNA was extracted and purified from bones devoid of marrow using the miRNeasy mini kit (Qiagen, Valencia, CA)⁽³³⁾. Real-time quantitative PCR (qPCR) was performed using specific primers (iQ SYBR Green Supermix, BioRad) and Taqman probes (Applied Biosystems Taqman Assays, Thermo Fisher Scientific) following manufacturer's protocol. Results are shown as relative gene expression, using expression of the 18s ribosomal RNA as an internal reference gene. Additional details on the RNA extraction and primers are provided in the Supplemental data (Table S1).

3.4 Histology.

For immunohistochemistry and immunofluorescence, paraffin embedded bone sections (7 μ m thickness) were incubated with rabbit polyclonal anti-MMP13 antibody (Abcam ab39012); rabbit polyclonal monoclonal anti-MMP14 antibody (Abcam ab53712); rabbit polyclonal anti-Cathepsin K (CTSK) antibody (Abcam ab19027), or rabbit monoclonal anti-T β RII antibody (Abcam ab186838) and mouse monoclonal anti-PTHr1 antibody (NBP1-40067: Novus Biologicals). Corresponding rabbit IgG negative control antibody was used to verify the specificity of primary antibodies. Ploton silver staining was performed to visualize the osteocyte lacuno-canalicular network in bones and the osteocyte network area was determined and normalized to total bone area, as previously described^(33,44). Images were acquired using a Nikon Eclipse E800 bright-field microscope and analyzed with Image J. Sections were evaluated for one femur from each of n = 3 mice per group, as indicated in the figure legends. Quantitative averages represent data collected from 4–5 high-powered fields for each bone. Additional details on staining, image analysis, and quantification are provided in the Supplemental data.

3.5 Synchrotron radiation Micro-Tomography (SR μ CT).

SR μ CT studies were used to assess the degree of mineralization of the bone as well as the volume and degree of anisotropy of the lacunae. The mid-diaphysis of 15-week-old virgin and lactating female mouse femora were scanned with a 20 keV x-ray energy, a 300 ms exposure time and a 5x magnifying lens for a spatial resolution of 1.3 μ m (n=4 bones/group). Additional details of SR μ CT procedure are described in the Supplemental data.

3.6 Flexural strength tests.

Whole bone flexural strength of femora was determined by three-point bending of intact, hydrated femurs isolated from T β RII^{ocy-/-} and control mice. 15-week old males and female virgin and lactating mice of each genotype were used for macromechanical testing as previously described⁽³³⁾. Details on the flexural strength tests are described in Supplemental data.

3.7 Serum Analysis.

Blood was isolated from male and virgin and lactating female mice using a cardiac puncture protocol and processed for serum. Serum calcium and phosphorus were measured with colorimetric assays (Abcam). PTH in serum was analyzed by Enzyme-linked Immunosorbent Assay (ELISA) (Quidel Corporation, Catalog no. 60-2305).

3.8 Statistical Analysis.

Sample size was determined based on a power calculation that provides an 80% chance of detecting a significant difference (p<0.05). Technical replicates and biological replicates (n) used for all experiments are described in the figure legends. Where technical replicates are used, data are expressed as mean \pm S.E.M. Otherwise, data are reported as mean \pm S.D. Prism 5.0 (GraphPad Software, Inc., San Diego, CA, USA) was used for statistical analysis. Comparisons between two groups were evaluated by an unpaired two-tailed Student's t

test. For comparing between more than two groups, we used one-way ANOVA followed by Tukey's test for multiple comparisons.

4. Results

4.1. Loss of osteocyte-intrinsic TGF β signaling impacts bone mass and quality in male but not female mice.

Recent studies from our group implicate osteocyte-intrinsic TGF β signaling to be key for controlling perilacunar/canalicular remodeling (PLR) and bone quality, however it is unclear if this regulatory function of TGF β is subject to sex-specific differences. Indeed, we previously observed sexual dimorphism in the effect of TGF β inhibition on bone mass and osteoblast and osteoclast numbers⁽³⁷⁾. To address this question, we used a previously developed mouse model for osteocyte-intrinsic inhibition of TGF β signaling to assess the bone phenotype in both sexes. Using the 9.6-kb *Dmp1-Cre* transgene, the floxed T β R $\text{II}^{\text{ocy-/-}}$ allele was preferentially excised from odontoblasts and osteocytes of T β R $\text{II}^{\text{ocy-/-}}$ mice^(33,41,42). The efficiency of this mouse model was confirmed by observing reduction in T β R II mRNA and protein levels in the long bone osteocytes of T β R $\text{II}^{\text{ocy-/-}}$ male and female mice (Figure 1A-D, Fig S1A-C and Fig 3 C-F).

To assess the skeletal phenotype of T β R $\text{II}^{\text{ocy-/-}}$ male and female mice, micro-computed tomography (μ CT) was used. These analyses revealed an increase in trabecular thickness and trabecular bone volume in male T β R $\text{II}^{\text{ocy-/-}}$ mice compared to the control littermates (Figure 1E-G, table 1). Although cortical bone mass was unchanged, the cortical bone mineralization of T β R $\text{II}^{\text{ocy-/-}}$ male mice was reduced by 3.5% relative to littermate controls (Figure 1E, H-I). Flexural strength tests showed that both yield stress and bending modulus of T β R $\text{II}^{\text{ocy-/-}}$ male bones was decreased by 20%, thereby indicating a reduction in the bone material quality of T β R $\text{II}^{\text{ocy-/-}}$ males compared to control mice (Figure 1J-K). As expected, based on our prior work on 8-week old T β R $\text{II}^{\text{ocy-/-}}$ males, the current findings confirm the critical role of osteocyte-intrinsic TGF β signaling in maintaining bone quality in 15-week old male mice.

None of these bone quality defects in T β R $\text{II}^{\text{ocy-/-}}$ males were present in female T β R $\text{II}^{\text{ocy-/-}}$ mice. In particular, the trabecular and cortical bone mass and geometry of T β R $\text{II}^{\text{ocy-/-}}$ females remained intact, and the flexural strength of T β R $\text{II}^{\text{ocy-/-}}$ female bones was identical to that of the control group (Figure 1E-I, J-K, Table 2). These observations suggest that female mouse bones do not depend on osteocytic TGF β signaling for maintaining bone mass and quality, and that osteocyte-intrinsic TGF β impacts bone in a sexually dimorphic manner.

4.2 TGF β signaling regulates osteocyte-mediated PLR in sex-specific manner.

Bone quality is in part regulated by PLR, and our prior evidence from 8-week-old T β R $\text{II}^{\text{ocy-/-}}$ male mice ascribes their defects in bone quality to PLR suppression⁽³³⁾. Therefore, we hypothesized that the distinct bone quality phenotype of male and female T β R $\text{II}^{\text{ocy-/-}}$ mice is driven by differences in osteocytic PLR. Since an intact lacuno-canalicular network (LCN) is a hallmark of functional PLR, we first examined the LCN in femora from 15-week old male and female T β R $\text{II}^{\text{ocy-/-}}$ mice. Compared to control male

mouse bones, the LCN of 15-week old $T\beta RII^{ocy-/-}$ bones was severely deteriorated, with a 50% reduction in LCN area and blunted canalicular length (Figure 2A-B, D). It is possible that the deregulation of TGF β signaling may affect the integration of osteocytes into the bone matrix and an inducible model would be needed to conclusively address this question. Nonetheless, our previous observation of similar lacuno-canalicular defects in response to pharmacologic inhibition of TGF β signaling (SD208) suggests that this is not solely a developmental defect⁽³³⁾. In addition, the male $T\beta RII^{ocy-/-}$ bones also displayed a strong coordinated decrease in the expression of genes encoding factors involved in bone resorption during PLR, namely, *Mmp2*, *Mmp14* (matrix metalloproteinases), *Ctsk* (Cathepsin K), *Acp5* (tartrate-resistant acid phosphatase), *Atp6v0d2* (*AtpD2*), and *Atp6v1g1* (*AtpG1*) (vacuolar ATPase pump encoding genes) (Figure 2F and H). Moreover, osteocyte-intrinsic protein expression of MMP13, MMP14 and CTSK was reduced in $T\beta RII^{ocy-/-}$ male mouse bones (Figure 2J and Figure S1).

Female $T\beta RII^{ocy-/-}$ mice responded differently to osteocyte-intrinsic TGF β ablation. In the female $T\beta RII^{ocy-/-}$ mice, the osteocyte LCN was intact with no changes observed in LCN area or canalicular length (Figure 2A, C and E). Consistent with these results, the expression of PLR genes in the bone of female $T\beta RII^{ocy-/-}$ mice did not differ from that in control mice (Figure 2G, I and K). Thus, the homeostatic regulation of PLR and bone quality in female mice occurs independently of TGF β signaling. Since the absence of osteocytic TGF β signaling in males adversely affects these outcomes, our data show a sexually dimorphic role for TGF β in the control of PLR and bone quality.

4.3 Osteocytic deletion of $T\beta RII$ prevents lactation induced PLR.

Although the homeostatic regulation of PLR in females is TGF β -independent, we hypothesized that intact TGF β signaling may be required to induce PLR in response to the metabolic stress of lactation. Using an established murine model of lactation, we assessed PLR outcomes in 15-week old lactating $T\beta RII^{ocy-/-}$ and lactating control mice. These results were compared to those in the 15-week old female $T\beta RII^{ocy-/-}$ and control virgin mice, described above. Upon validation of targeted $T\beta RII$ deletion at the mRNA and protein levels in the lactating $T\beta RII^{ocy-/-}$ mice (Figure 3A, C-F), we monitored the effect of lactation on the expression of PLR genes. During lactation, control mice showed a 2.5 to 4-fold induction in the expression of the *Mmp14*, *Ctsk*, *Acp5* and *Atp6v0d2* genes (Figure 3A-B). The increased expression of PLR enzymes was also apparent at the protein level, with a 3 to 5-fold increase in the percentage of MMP13, MMP14 and CTSK-stained cortical bone area in the lactating control group (Figure 3C-D, E-G). Similar to resorptive genes, the lactation-induced expression of bone formation genes, *Dmp1* and *Runx2*, was absent in $T\beta RII^{ocy-/-}$ mice (Figure S2A-B).

Consistent with these observations, lactating control mice undergo a 50% increase in the LCN area, evident from the Ploton silver staining (Figure 4A-B). Although no changes in the canalicular length were observed in either of the virgin or lactating $T\beta RII^{ocy-/-}$ mice compared to control group, changes in osteocyte lacunar features were detected with lactation. Using synchrotron X-ray tomography (SR μ T), we quantified the lacunar volume of thousands of osteocyte lacunae, visualized in 3D. By conducting a tertile analysis of

small, medium, and large- sized osteocytes, we observed that the lactating controls have significantly fewer small-sized lacunae, but more medium and large lacunae, compared to virgin controls. On the other hand, this lactation-induced shift from small to larger osteocyte lacunae is impaired in $T\beta RII^{ocy-/-}$ mice, where lactation-dependent differences are only apparent in the largest osteocytes. This suggests that osteocyte-intrinsic $TGF\beta$ signaling is required for maximal induction of PLR by lactation (Figure 4C).

Overall, lactation failed to induce PLR in $T\beta RII^{ocy-/-}$ mice. Lactating $T\beta RII^{ocy-/-}$ mice did not exhibit a significant induction in PLR genes, LCN area, lacunar volume and canalicular length (Figure 3 and 4). Therefore, distinct from the $TGF\beta$ -independent regulation of PLR in female bone at homeostasis, osteocyte-intrinsic $TGF\beta$ signaling is required for the induction of PLR during lactation.

4.4 Targeted deletion of osteocytic $T\beta RII$ mitigates lactation-induced bone loss.

μCT analyses revealed that during lactation, control mice lose more than 52% of trabecular bone volume fraction (BV/TV, Table 1). This decrease in BV/TV is reflected by the corresponding reduction in trabecular thickness (32%) and increase in trabecular spacing (45%) during lactation in control mice (Table 1, Figure 5A-E). The trabecular bone mineralization is also reduced by 9% in the lactating control mice. A decrease in cortical bone thickness (27% reduction) and mass (25% reduction) was observed in lactating controls but the cortical mineralization remained intact (Table 1, Figure 5F-G). In contrast to control group, $T\beta RII^{ocy-/-}$ mice did not lose trabecular bone mass during lactation, and their trabecular number remained unaltered, while their trabecular thickness and spacing decreased only slightly with lactation. $T\beta RII^{ocy-/-}$ mice lost some cortical thickness and bone mass during lactation, however the magnitude of cortical bone loss was significantly lower in lactating $T\beta RII^{ocy-/-}$ mice than control mice (i.e. 14% vs. 25%, $p < 0.05$) (Figure 5A-G).

The high-resolution quantitative imaging available through synchrotron SR μCT further revealed a comparable trend towards increased mineralization around osteocyte lacunae with lactation that was apparent in both genotypes (Figure S3). However, μCT data did not show significant differences in overall cortical bone mineralization during lactation in either genotype (Table 1). As expected, stiffness and yield load were significantly reduced in lactating control mice relative to their virgin counterparts (Table 2), consistent with the loss of cortical bone. However, these parameters were unaffected by lactation in the $T\beta RII^{ocy-/-}$ mice. When normalized for structural parameters, tissue properties such as bone bending modulus, yield stress, and ultimate stress were unaffected by lactation in either genotype (Table 2). Collectively, these data establish the role of osteocyte-intrinsic $TGF\beta$ signaling in lactation, such that absence of this mechanism alleviates lactation-induced bone remodeling by osteocytes and surface cells.

4.5 PTH involvement in the sexually dimorphic effects of osteocytic $TGF\beta$.

As we previously reported, the elevated bone mass in 8-week old $T\beta RII^{ocy-/-}$ male mice is accompanied by reduced Rankl (*Tnfsf11*) expression and decreased osteoclast numbers⁽³³⁾. Therefore, we hypothesized that the sex-specific differences in bone mass in $T\beta RII^{ocy-/-}$

male and female mice were driven by dimorphic regulation of Rankl. Here we find that 15-week old male $T\beta RII^{ocy-/-}$ mice also express low levels of Rankl mRNA, with a stark reduction in the Rankl/Opg ratio, compared to control males. Female $T\beta RII^{ocy-/-}$ mice, on the other hand, showed intact Rankl and Rankl/Opg ratios during homeostasis (Figure 6A-B). During lactation, the high demand for calcium drives bone resorption, in part, through robust Rankl expression⁽⁴⁵⁾. Indeed, lactating control mice increase Rankl mRNA expression with a resulting 7-fold increase in the Rankl/Opg ratio (Figure 6C and S2C). Lactating $T\beta RII^{ocy-/-}$ mice, however, fail to stimulate Rankl mRNA expression. Instead Opg mRNA levels are suppressed in the lactating $T\beta RII^{ocy-/-}$ mice, through an apparent compensatory mechanism to increase the Rankl/Opg ratio 2-fold to release calcium for lactation (Figure 6C and S2C).

These differences in Rankl/Opg expression led us to focus on the role of PTH signaling in the sexually dimorphic effects of TGF β in osteocytes. Several lines of evidence motivated our focus: First, PTH signaling directly regulates Rankl gene expression in osteocytes^(46,47). Second, PTH and parathyroid hormone related peptide (PTHrP), key regulators of calcium and phosphate homeostasis, induce PLR during lactation^(5,21,48-50). Third, PTH and PTHrP signal through a common parathyroid hormone receptor (PTHr1) in osteocytes; and PTHr1 ablation in osteocytes blocks lactation-inducible PLR⁽²¹⁾. Fourth, as in $PTHr1^{ocy-/-}$ mice, $T\beta RII^{ocy-/-}$ mice are unable to induce PLR or bone loss during lactation⁽²¹⁾. Fifth, $T\beta RII$ and PTHr1 interact in osteoblasts to augment PTH signaling and increase bone formation⁽⁵¹⁾. Based on these findings, we hypothesized that TGF β and PTH signaling utilize a common molecular mechanism to regulate PLR.

To test this hypothesis, we first examined the effect of $T\beta RII$ ablation on the expression of PTHr1 in the osteocytes. Both immunohistochemistry and quantitative PCR revealed a stark reduction (>70%) in the expression of PTHr1 in osteocytes of $T\beta RII^{ocy-/-}$ male mice compared to controls (Figure 6D, G, H). This was consistent with the low Rankl mRNA and Rankl/Opg ratios seen in the $T\beta RII^{ocy-/-}$ males. In females during homeostasis, no changes were detected in the expression of PTHr1 at the mRNA or protein level between control and $T\beta RII^{ocy-/-}$ mice, which agrees with the lack of changes in Rankl/Opg in $T\beta RII^{ocy-/-}$ females (Figure 6E, G, I). During lactation, PTHr1 expression is robustly increased in control females, but loss of osteocytic $T\beta RII$ mitigates this response in lactating $T\beta RII^{ocy-/-}$ mice (Figure 6F, G, J), consistent with the lower overall Rankl/Opg ratio in lactating $T\beta RII^{ocy-/-}$ mice compared to lactating controls.

Lastly, we evaluated serum levels of calcium, phosphate, and PTH in each group (Table S2). Phosphate levels did not differ in any condition. The two groups that had reduced PTHr1 expression, male and lactating female $T\beta RII^{ocy-/-}$ mice, had elevated levels of serum calcium relative to control mice. In male $T\beta RII^{ocy-/-}$ mice, the hypercalcemia may result from increased serum PTH levels. Importantly, in non-lactating female mice, none of these parameters differed between $T\beta RII^{ocy-/-}$ mice and controls, which is consistent with the insensitivity of these mice to osteocyte-intrinsic TGF β ablation in all the other outcomes we analyzed. Taken together these findings suggest a key role of $T\beta RII$ in regulating PTHr1 expression in osteocytes, and that this coordination between TGF β and PTH signaling dictates the PLR response in osteocytes during homeostasis and metabolic stress.

5. Discussion

This study provides insight into sex-specific variation in bone fragility by revealing dimorphism in the regulation of osteocyte perilacunar/ canalicular remodeling (PLR). Using a genetic model of osteocyte-specific deletion of TGF β receptor II (T β RII), we show that loss of osteocyte-intrinsic TGF β signaling affects the skeleton of male, but not female, mice in homeostatic conditions. Male T β RII^{oc β -/-} mice have defective bone quality, despite normal bone mass, resulting from impaired osteocytic PLR with deteriorated canalicular networks and reduced expression of requisite PLR enzymes. On the contrary, female T β RII^{oc β -/-} mice are protected from these deficits in both PLR and bone quality. Osteocytic TGF β signaling is necessary for PLR induction in female mice during lactation. Lactation-induced PLR is mitigated in T β RII^{oc β -/-} females, as is lactation-induced bone loss. These findings highlight distinct roles for TGF β signaling in regulating PLR in males and females to maintain bone homeostasis and implicate TGF β as an essential regulator of PLR in females principally during the metabolic stress of lactation.

Gender disparity in fracture risk stems from differences in peak bone mass accrual, bone size, and geometry⁽⁵²⁾. Sex steroids that essentially drive skeletal variations in men and women integrate systemic cues such as that from vitamin D, IGF/GH, and PTH to dictate dimorphism at the cellular level^(19,53-55). Indeed, sex differences in the cellular function of osteoblasts and osteoclasts have been attributed to differences in the bone phenotype of male and female mice⁽¹⁷⁻²⁰⁾. Differences in bone quality also contribute to the gender disparities in fracture risk.

Osteocyte PLR is one cellular mechanism that plays a pivotal role in maintaining bone quality. Although the contribution of PLR to sex differences in bone quality was unknown, osteocytes clearly have a dimorphic response to aging^(56,57). The osteocyte lacuno-canalicular network, which is maintained by PLR, deteriorates more rapidly in aging females than in aging males^(56,57). Here we describe sexually dimorphic regulation of osteocytes in the control of PLR and bone quality by TGF β signaling. Whereas in males, loss of osteocytic TGF β signaling suppresses PLR and increases bone fragility, female T β RII^{oc β -/-} mice demonstrate a complete lack of this regulatory mechanism. Such skeletal adaptations in females could be one of the mechanisms by which the female skeleton compensates to meet the skeletal demands of pregnancy and lactation. The extent to which this 'protective mechanism' in reproductive age females is lost following menopause, or if it contributes to the increased risk of post-menopausal bone fragility remains unknown. Nevertheless, our study shows that sex specific differences in osteocyte sensitivity to TGF β signaling contribute to the sexual dimorphism in PLR and bone quality.

TGF β signaling has previously been implicated as one of the mechanisms recruited to confer sex specific differences in bone. An intricate crosstalk exists between TGF β signaling and estrogen in bone. Estrogen promotes de novo synthesis and activation of TGF β ligand in bone. Conversely, the loss of estrogen in ovariectomized rats leads to reduced TGF β expression in bone⁽⁵⁸⁻⁶²⁾. TGF β , in turn, mediates bone protective effects of estrogen by increasing osteoclast apoptosis^(59,63). With systemic inhibition of TGF β signaling using a pharmacologic TGF β receptor I kinase inhibitor (T β RI-I), we observed increased sensitivity

of the female skeleton to TGF β inhibition, with increased osteoblast proliferation and bone mass accrual in female mice than males⁽³⁷⁾. Here, in osteocytes of female mice with ablated T β RII, we observed that loss of osteocytic TGF β signaling has no impact on PLR gene expression, the osteocyte lacuno-canalicular network, and bone quality. The extent to which these sex specific differences in osteocyte TGF β signaling are related to estrogen levels remains to be determined. The remarkably distinct regulation of TGF β signaling by sex-specific factors in osteoblasts, osteoclasts and osteocytes could afford independent, yet coordinated, regulation of bone quantity and quality to maintain mineral homeostasis and bone strength during hormonal and reproductive cycles in females.

Systemic calcium homeostasis is tightly regulated by the parathyroid gland that controls the interplay between the calciotropic hormones like, parathyroid hormone (PTH), parathyroid hormone related protein (PTHrP), calcitonin, and calciferol (1 α ,25-dihydroxycholecalciferol, vitamin D3). Together these hormones establish an equilibrium between dietary calcium absorption through the intestines, calcium reabsorption and excretion through kidneys, and its storage and release from bones⁽⁶⁴⁾. In bone, osteocytes show particularly high sensitivity to calcium fluctuations and are a target of PTH action^(21,29,65). PTHR1, the receptor that binds both PTH and PTHrP, is highly expressed in bone and kidney and mediates the PTH-dependent regulation of calcium homeostasis. Although, little is known about the crosstalk of PTH and PTHrP with TGF β signaling in bone, PTHR1 interacts directly with T β RII. In osteoblasts, PTHR1 and T β RII coordinately modulate bone formation⁽⁵¹⁾. In osteocytes, however, we observed that ablation of T β RII causes a marked reduction in PTHR1 expression and Rankl/ Opg ratios only in male mice. Given that PTHR1 is a key regulator of PLR, the PLR and bone quality defects observed in the male T β RII^{ocy-/-} may result from reduced PTHR1-dependent signaling. The reduced PTHR1-dependent signaling in T β RII^{ocy-/-} male mice appears to be compensated by increased systemic PTH levels, which in turn increase systemic calcium levels, likely by targeting tissues other than bone.

In female mice, the role of TGF β in osteocytes differs dramatically between homeostatic conditions and the metabolic stress of lactation. In homeostatic conditions, female T β RII^{ocy-/-} mice differ from T β RII^{ocy-/-} males, in that they show intact osteocytic PTHR1 expression and PLR. Other pathways that are active in females, but not in males, may converge on PTHR1 to sustain its expression in female mice at homeostasis. This capacity of female T β RII^{ocy-/-} osteocytes to normalize PTHR1 levels at homeostasis may contribute to the intact bone quality of female T β RII^{ocy-/-} mice. This capacity is challenged by the metabolic stress of lactation, which normally induces PLR to meet the high calcium demands for breastmilk production. PLR induction during lactation occurs in a hormonal milieu with low PTH and estradiol, but with high PTHrP levels. While we demonstrated reduction in circulating PTH with lactation in both genotypes, the circulating level of PTHrP was not significantly increased in either genotypes, most likely due to insufficient sensitivity of our assay. PLR induction during lactation also requires osteocytic PTHR1. Although PTHR1 mRNA is induced in lactation in both genotypes, the induction of PTHR1 protein in osteocytes is abrogated in lactating T β RII^{ocy-/-} mice relative to control littermates. Accordingly, with reduced osteocytic PTHR1 protein levels, PLR induction in female T β RII^{ocy-/-} mice during lactation is blocked. Our results point to the possibility

that the level of PTHR1 is a pivotal determinant of the effect of TGF β on PLR. Thus, the differential PLR response of osteocytes in the T β RII^{ocy-/-} mice in males vs. females, and in homeostasis vs. metabolic stress, may result from differences in the ability to stimulate PTHR1-mediated bone resorption by osteocytes. It is likely that such molecular mechanisms are in place to compensate for reproductive bone loss and protect the female skeleton from higher fracture risk. Another such compensatory mechanism that was recently suggested includes accrual of higher trabecular bone mass than mechanically necessary by female rats compared to males^(1,66).

In conclusion, our study highlights sexual dimorphism in TGF β 's regulation of osteocyte perilacunar/canalicular remodeling activity. With implication of PLR in both physiological and pathological conditions, including renal osteodystrophy, glucocorticoid-induced osteonecrosis, osteoarthritis, and hyperparathyroidism^(21,44,67-71), defining the sex differences in these cell-intrinsic cues and molecular networks in osteocytes will be critical to address sexual dimorphism in the prevalence and manifestation of bone fragility, as well as refining therapeutics for reducing fracture risk.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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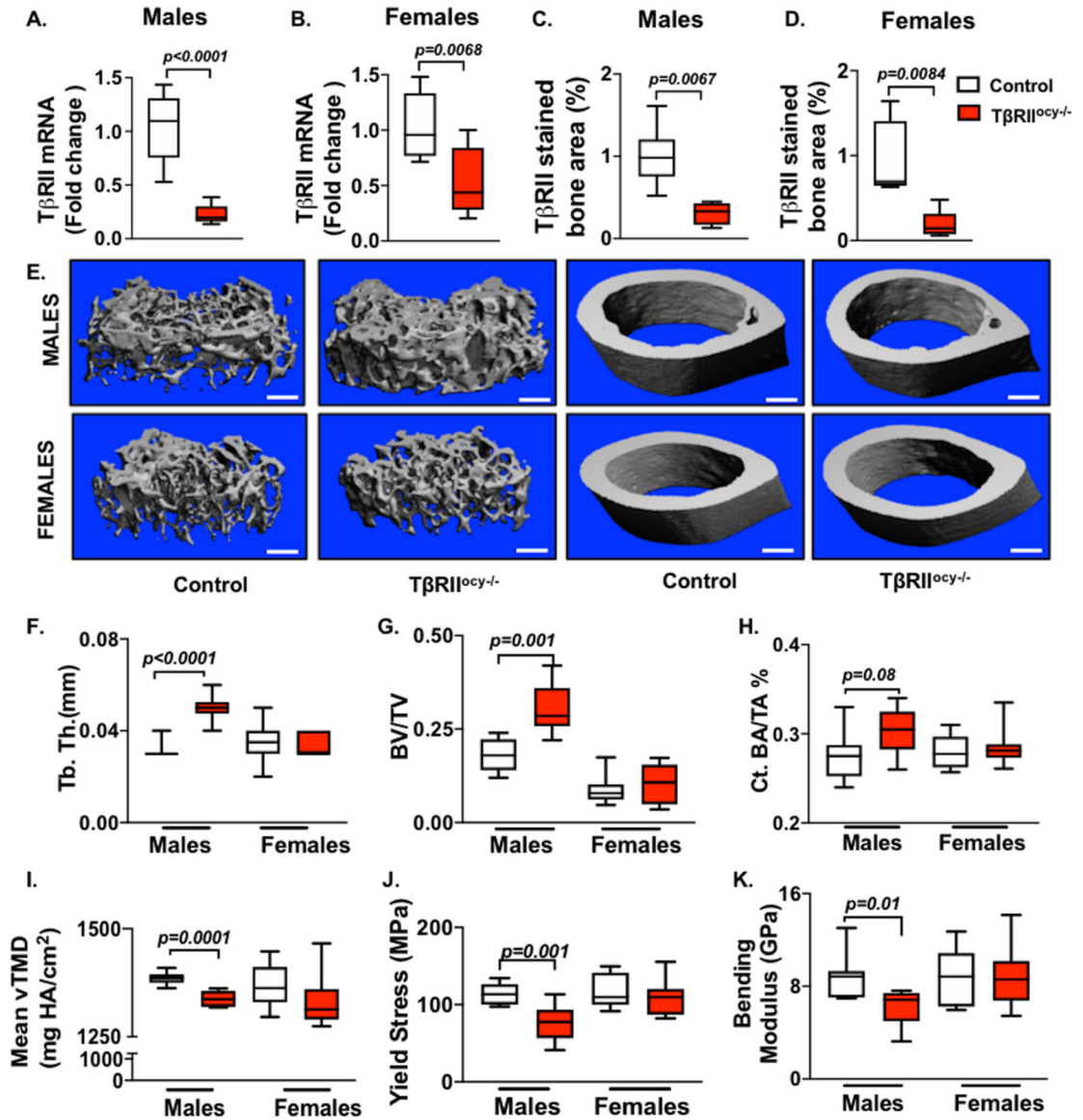


Figure 1. Loss of osteocyte-intrinsic TGF β signaling impacts bone mass and quality in male mice but does not affect female mice.

In male and female $T\beta RII^{ocy-/-}$ mice, knockdown of $T\beta RII$ at mRNA (A, B) ($n=6-9$ mice/ group) and protein (C, D) ($n=4-6$ mice/ group) level was assessed. μ CT analysis was conducted on femurs from control and $T\beta RII^{ocy-/-}$ male and female mice (15-week-old). Representative μ CT reconstructions of trabecular (left) and cortical (right) bone (E) along with the trabecular bone thickness ($Tb. Th.$) (F) and volume (BV/TV) (G), and cortical bone area (BA/TA) (H) and mineralization (Ct. Min) (I) are shown ($n=7-8$ mice/group). Flexural testing of femurs from male and female control and $T\beta RII^{ocy-/-}$ mice shows bone quality parameters yield stress (J) and bending modulus (K) ($n = 7$ mice/group). Data for A-D are presented as mean \pm SEM and as mean \pm SD for E-K and statistically significant difference were determined with Student's t test. Scale bar for E is 100 μ m ($n = 7-8$ mice/group).

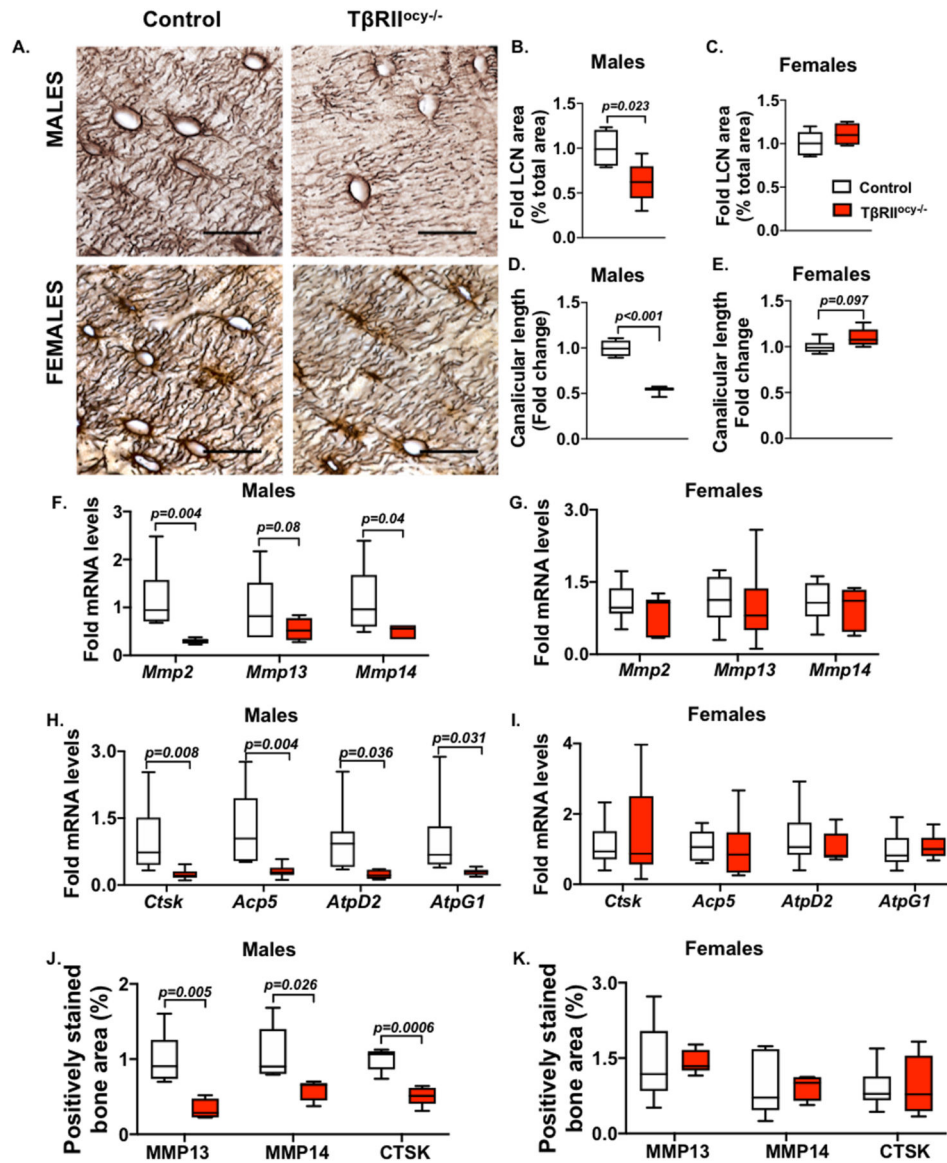


Figure 2. Osteocyte-specific disruption of TβRII reduces PLR genes in male but not female mice. 15-week-old male and female control and TβRII^{ocy-/-} mouse bones were processed for mRNA and protein. Osteocyte lacuno-canalicular network (LCN) in the femoral cortical bones was assessed by Pleton stain. Representative images (A) and quantification of LCN area (B, C) and canaliculus length (D, E) are shown. Scale bar, 20μm; n=4–6 mice/group and 4 ROI/mouse. Expression of PLR genes Mmp2, Mmp13 and Mmp14 (F, G) and Ctsk, Acp5, Atp6v0d2 (AtpD2) and Atp6v1g1 (AtpG1) are shown (H-I) (n=6–9 mice/ group). Immunohistochemistry (IHC) for MMP13, MMP14, and CTSK on femoral cortical bones of male and female control and TβRII^{ocy-/-} mice shows percent positively stained bone area (J- K) (n =3–4 mice/group and 4 ROI/mouse). Scale bar, 30μm. Error bars indicate mean ± SEM, statistically significant differences were calculated with Student's t test.

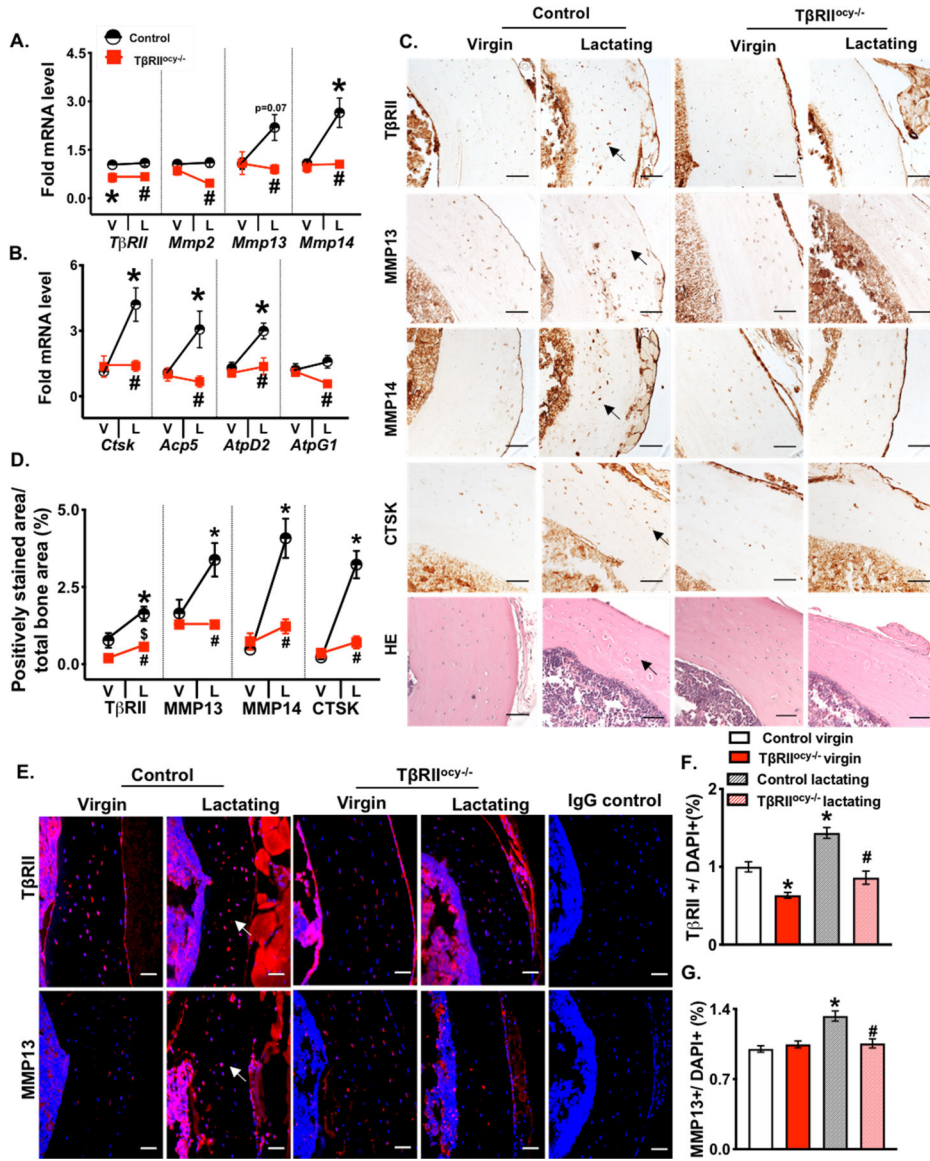


Figure 3. Lactation induced PLR is blocked by osteocyte-specific disruption of *TβRII* gene. Bones from 15-week-old virgin and lactating control and *TβRII^{ocyt-/-}* female mice were harvested and processed for gene expression (A-B) and immunohistochemistry (C-D). qPCR analysis of *TβRII* and PLR genes *Mmp2*, *Mmp13*, *Mmp14*, *Acp5*, *Ctsk*, *AtpD2* and *AtpG1* (A-B) in virgin and lactating control and *TβRII^{ocyt-/-}* mouse bones was compared (n=6–9 mice/group). Immunohistochemistry (IHC) for *MMP13*, *MMP14*, and *CTSK* was conducted on femoral cortical bones of virgin and lactating control and *TβRII^{ocyt-/-}* mice. Arrows in the representative images indicate positively stained area (D) that were quantified and normalized to total bone area (C) (n=3–4 mice/group and 4 ROI/mouse, scale bar is 50μm). Representative images of immunofluorescence of *TβRII* (top) and *MMP13* (bottom) protein and (E) quantification of cells positive for *TβRII* and DAPI, or *MMP13* and DAPI in murine cortical bone (F and G) (scale bar is 40μm) is shown. Error bars indicate mean ± SEM, *p<0.05 compared to control virgins, #p<0.05 compared to control lactating group,

and $p < 0.05$ compared to $T\beta RII^{ocy-/-}$ virgin, as calculated from one-way ANOVA with Tukey post-hoc test.

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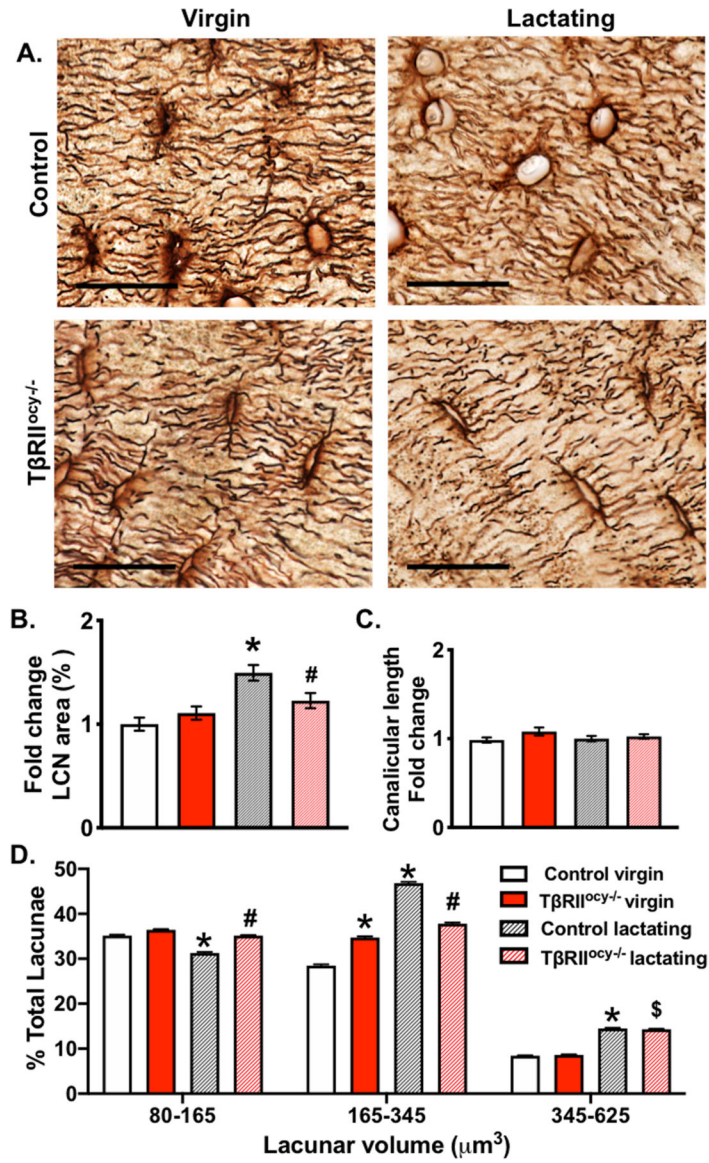


Figure 4. Osteocyte-specific disruption of TβRII gene suppresses lactation induced increase in osteocyte lacunar volume.

Representative Ploton stained histological sections of femoral cortical bones of 15-week-old virgin and lactating control and TβRII^{ocy-/-} female mice (A) show the lacuno-canalicular network integrity (scale bar = 20μm). Quantification of average percent lacuno-canalicular area (normalized to total area) (B) and canalicular length (C) after 7 days of lactation is shown. XTM was used to determine osteocyte lacunar volume and the overall lacunar size distribution in cortical bone is shown in a graph, in control mice significant reduction in the overall percentage of smaller osteocytes and increase in larger osteocytes is observed in response to lactation. Scale bar for A is 20μm, and error bars indicate mean ± SEM, *p<0.05 compared to control virgin, #p<0.05 compared to control lactating, and \$ p<0.05 compared to TβRII^{ocy-/-} virgin, as calculated from one-way ANOVA with Tukey post-hoc test (n=4–5 mice/group).

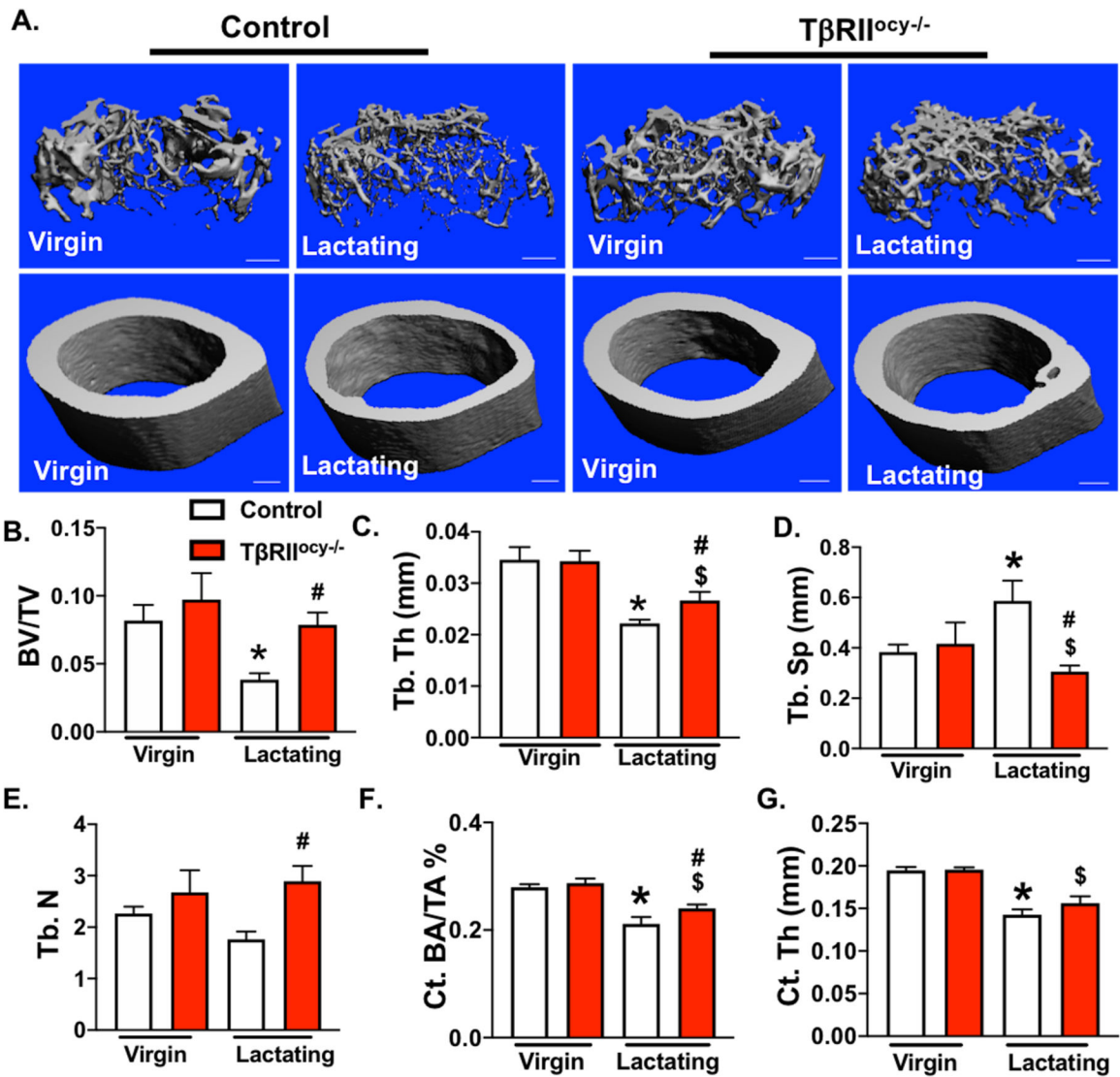


Figure 5. Lactation induced bone loss in TβRII^{ocy-/-} is mitigated.

μCT analysis of femurs from 15-week old control and TβRII^{ocy-/-} lactating female mice. Representative μCT reconstructions show trabecular (top) and cortical (bottom) bones (A). Decline in trabecular bone parameters: bone volume fraction (BV/TV), trabecular thickness (Tb.Th.), trabecular spacing (Tb.Sp.), and trabecular number (Tb.N) (B-E), and cortical bone parameters: cortical bone volume (Ct.BA/TA) and cortical thickness (Ct. Th), (F-G) in virgin and lactating mice of both genotypes is shown below. Scale bar, 100 μm (n = 6–10 mice/group). Error bars indicate mean ± SEM, *p<0.05 compared to control virgins, \$p<0.05 compared to TβRII^{ocy-/-} virgin group and #p<0.05 compared to control lactation group, as calculated from one-way ANOVA with Tukey post-hoc test.

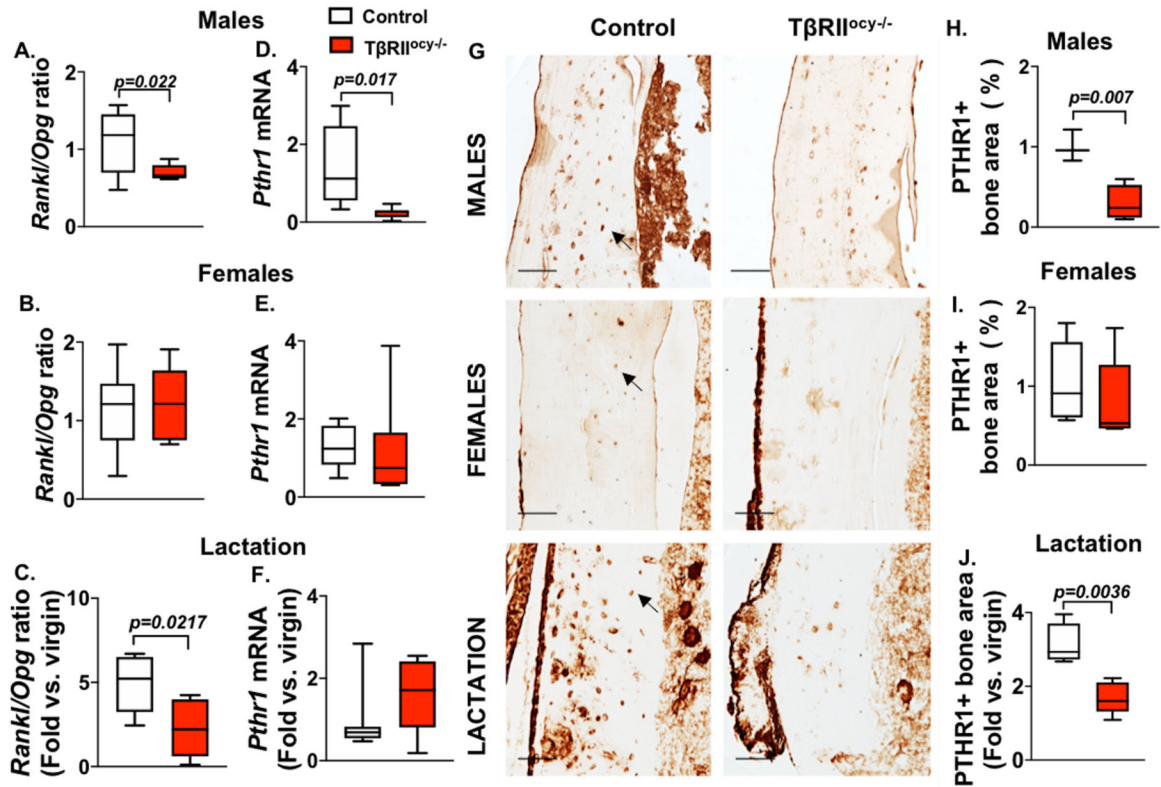


Figure 6. PTHR1 dependent signaling is reduced in $T\beta RII^{oc/-}$ males and lactating females. qPCR analysis for Rankl and Opg in control and $T\beta RII^{oc/-}$ males (A) and virgin (B) and lactating (C) females is presented as a ratio of Rankl/Opg mRNA, with lactating groups shown relative to corresponding virgin groups. *Pthr1* mRNA was also quantified by qPCR (D-F) and by immunohistochemistry (G-J). Percentage of osteocytes that stained positive for PTHR1 in bones of control and $T\beta RII^{oc/-}$ male and virgin and lactating female mice were quantified and normalized against total bone area as shown (H-J) (n=3–4 mice/group, 4 ROI/mouse). Error bars indicate mean \pm SEM, statistically significant differences were calculated from Student’s t-test.

Table 1:

μ CT of male and female (virgin and lactating) control and T β RII^{ocy-/-} mouse bones.

	Male		Female		Lactation	
	Control	T β RII ^{ocy-/-}	Control	T β RII ^{ocy-/-}	Control	T β RII ^{ocy-/-}
Distal Femur						
<i>Tb.BV/TV</i> (%)	0.18 ± 0.04	0.30 ± 0.07 [*]	0.08 ± 0.04	0.1 ± 0.05	0.04 ± 0.01 ^a	0.08 ± 0.02 ^b
<i>Tb.N</i> (1/mm)	5.30 ± 1.08	6.11 ± 0.87	2.44 ± 0.72	2.68 ± 1.13	1.76 ± 0.45	2.89 ± 0.84
<i>Tb.Th</i> (mm)	0.03 ± 0.003	0.05 ± 0.01 [*]	0.033 ± 0.01	0.035 ± 0.01	0.022 ± 0.002 ^a	0.027 ± 0.00 ^{a,b,c}
<i>Tb.Sp</i> (mm)	0.16 ± 0.05	0.12 ± 0.03 [*]	0.40 ± 0.12	0.42 ± 0.22	0.59 ± 0.2 ^a	0.35 ± 0.15 ^{b,c}
<i>SMI</i>	1.71 ± 0.36	0.52 ± 0.62 [*]	2.83 ± 0.43	2.74 ± 0.66	3.01 ± 0.27	2.48 ± 0.38
<i>Tb.Min</i> (mg HA/cm ³)	1065.5 ± 18.9	1098.3 ± 16.3 [*]	1103.85 ± 30.2	1096.5 ± 23.6	1010.5 ± 27.0 ^a	1035.3 ± 40.9 ^c
Midshaft Femur						
<i>Ct.BA/TA</i> (%)	0.28 ± 0.03	0.30 ± 0.03	0.28 ± 0.02	0.29 ± 0.02	0.21 ± 0.03 ^a	0.24 ± 0.02 ^{a,b,c}
<i>Ct.Th</i> (mm)	0.20 ± 0.01	0.22 ± 0.02 [*]	0.19 ± 0.01	0.20 ± 0.01	0.14 ± 0.006 ^a	0.16 ± 0.007 ^c
<i>Ct.SMI</i>	0.16 ± 0.41	2.82 ± 1.42 [*]	0.28 ± 0.14	0.12 ± 0.23	0.22 ± 0.03	0.10 ± 0.18
<i>Ct.Min</i> (mg HA/cm ³)	1385.8 ± 14.4	1337.8 ± 18.7 [*]	1362.60 ± 46.6	1332.5 ± 66.4	1330.3 ± 29.1	1319.2 ± 36.5

* p<0.05 significantly different from the male control virgin (Student's t-test),

^a p<0.05 significantly different from the control virgin,

^b p<0.05 significantly different from the control lactation, and

^c p<0.05 significantly different from the virgin T β RII^{ocy-/-} mice.

Table 2:

Flexural strength test derived macro-mechanical properties of 15-week-old male and virgin and lactating female control and T β RII^{ocy-/-} mice.

Flexural Strength parameters	Male		Female		Lactation	
	Control	T β RII ^{ocy-/-}	Control	T β RII ^{ocy-/-}	Control	T β RII ^{ocy-/-}
<i>Stiffness (N-mm²)</i>	189.8 ± 24.0	169.7 ± 28.9	173.9 ± 20.3	177.1 ± 24.4	129.6 ± 22.5 ^a	161.2 ± 14.7
<i>Yield load (N)</i>	14.4 ± 1.9	12.5 ± 2.3	14.0 ± 1.1	13.7 ± 1.1	12.0 ± 1.7 ^a	13.8 ± 1.2
<i>Post-yield displacement (%)</i>	0.87 ± 0.71	0.55 ± 0.25	0.54 ± 0.19	0.72 ± 0.19	0.59 ± 0.18	1.27 ± 0.51
<i>Modulus (GN/m²)</i>	9.06 ± 1.99	6.2 ± 1.51 [*]	9.0 ± 2.45	8.8 ± 2.63	7.88 ± 1.38	9.41 ± 1.40
<i>Yield Stress (Mpa)</i>	115.6 ± 13.9	80.4 ± 23.98 [*]	118.4 ± 21.8	108.7 ± 22.9	135.3 ± 29.1	135.9 ± 15.2
<i>Ultimate Stress (Mpa)</i>	189.5 ± 22.5	189.30 ± 37.5	191.05 ± 30.2	227.1 ± 43.5	167.3 ± 20.0	169.7 ± 4.45
<i>MOI (mm⁴)</i>	0.117 ± 0.02	0.158 ± 0.05	0.113 ± 0.03	0.118 ± 0.03	0.077 ± 0.02 ^a	0.084 ± 0.01

* p<0.05 significantly different from the male control virgin group (Student's t-test),

^a p<0.05 significantly different from the female control virgin group.