Effects of different training modalities on phosphate homeostasis and local vitamin D metabolism in rat bone

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Objectives: Mechanical loading may be an important factor in the regulation of bone derived hormones involved in phosphate homeostasis. This study investigated the effects of peak power and endurance training on expression levels of fibroblast growth factor 23 (FGF23) and 1α-hydroxylase (CYP27b1) in bone.

Methods: Thirty-eight rats were assigned to six weeks of training in four groups: peak power (PT), endurance (ET), PT followed by ET (PET) or no training (control). In cortical bone, FGF23 was quantified using immunohistochemistry. mRNA expression levels of proteins involved in phosphate and vitamin D homeostasis were quantified in cortical bone and kidney. C-terminal FGF23, 25-hydroxyvitamin D3, parathyroid hormone (PTH), calcium and phosphate concentrations were measured in plasma or serum.

Results: Neither FGF23 mRNA and protein expression levels in cortical bone nor FGF23 plasma concentrations differed between the groups. In cortical bone, mRNA expression levels of sclerostin (SOST), dental matrix protein 1 (DMP1), phosphate-regulating gene with homologies to endopeptidases on the X chromosome (PHEX) and matrix extracellular phosphoglycoprotein (MEPE) were lower after PT compared to ET and PET. Expression levels of CYP27b1 and vitamin D receptor (VDR) in tibial bone were decreased after PT compared to ET. In kidney, no differences between groups were observed for mRNA expression levels of CYP27b1, 24-hydroxylase (CYP24), VDR, NaPi-IIa cotransporter (NPT2a) and NaPi-IIc cotransporter (NPT2c). Serum PTH concentrations were higher after PT compared to controls.

Conclusion: After six weeks, none of the training modalities induced changes in FGF23 expression levels. However, PT might have caused changes in local phosphate regulation within bone compared to ET and PET. CYP27b1 and VDR expression in bone was reduced after PT compared to ET, suggesting high intensity peak power training in this rat model is associated with decreased vitamin D signalling in bone.
Full Title: **Effects of different training modalities on phosphate homeostasis and local vitamin D metabolism in rat bone.**

Short Title: **Effects of training on phosphate homeostasis and vitamin D in bone.**

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Abstract

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Introduction

Mechanical forces associated with physical exercise cause deformation of bone tissue.

These deformations activate an acute response in osteocytes, in particular production of cytokines and signalling molecules such as nitric oxide (NO), bone morphogenetic proteins (BMPs), Wnts, and prostaglandin E2 (PGE2) (1). These factors affect osteoblast and osteoclast recruitment, differentiation and activation locally at the sites that are subjected to high strains.

Since whole bone response to mechanical loading depends on the strain rate and magnitude (2,3), peak power training with high peak loads with a high frequency is generally considered to result in a higher bone response than endurance training. Moreover, it was reported that a combination of training types even better maintained bone mineral density in postmenopausal women (4).

Recently, it was proposed that mechanical loading affects gene expression and local protein activity of fibroblast growth factor 23 (FGF23) (5). FGF23 mRNA expression in mouse bone has been shown to increase after 6 days of endurance training (6). Moreover, it was suggested FGF23 might be a ‘molecular mediator of the whole-body effects of exercise originating from bone’ (7). Indeed, serum FGF23 increased after endurance exercise in rat and human (8,9), indicating a potential systemic effect of local mechano-response in bone; FGF23 is produced exclusively in bone but acts on the kidneys, where it decreases the reabsorption and increases excretion of phosphate and also suppress 1α -hydroxylase (CYP27b1), reducing its ability to activate vitamin D and subsequently impairing calcium absorption (10). To our knowledge, the direct effect of different training types on FGF23 expression within bone has not been reported yet.

A known stimulator of FGF23 production is systemic 1,25-dihydroxyvitamin D3 (1,25(OH)2D3) (11) or in vitro locally produced 1,25(OH)2D3 (12). Inhibitors of FGF23
production are mineralization-regulating proteins such as phosphate-regulating gene with homologies to endopeptidases on the X chromosome (PHEX) and dental matrix protein 1 (DMP1) (13), possibly through alterations in bone mineralization. Mechanical loading could either affect FGF23 gene expression directly, or indirectly by alterations in the expression of FGF23-regulating proteins known to be stimulated by mechanical loading, such as DMP1 (14,15) and matrix extracellular phosphoglycoprotein (MEPE) (16) as well as presumably locally converted 1,25(OH)2D3.

The active metabolite 1,25(OH)2D3 is hydroxylated from 25-hydroxyvitamin D3 (25(OH)D3) by 1α-hydroxylase (17), which is, among other tissues, expressed by human bone cells (18,19). The regulation of activity of 1α-hydroxylase within bone is yet to be determined. Van Driel et al. demonstrated that in cultured bone cells 25(OH)D3, PTH and calcium did not affect CYP27b1 mRNA expression and activity (18). In contrast, treatment of primary human osteoblasts with high doses of calcium did increase CYP27b1 mRNA expression (20). Pulsatile fluid flow, an in vitro model for mechanical loading, increased CYP27b1 expression in primary human osteoblasts (21), suggesting that 1α-hydroxylase within bone is at least partly regulated by mechanical loading. Among healthy, young women, serum 25(OH)D3 did not differ between different training groups (22). However, among rats an increase in serum 1,25(OH)2D3 after endurance training compared to controls was observed (23). Whether vitamin D signalling within bone is affected by mechanical loading in vivo remains to be elucidated.

Osteocytes may respond to mechanical loading by altering the expression of FGF23 and CYP27b1. As the response of osteocytes to mechanical loading is determined by the type of loading, FGF23 production and 1α-hydroxylation may be differentially changed according to the type of loading. We hypothesize that mechanical loading induced by physical exercise influences
FGF23 production and 1α-hydroxylation in bone, with the greatest response after a combination of both peak power training and endurance training. Therefore, the aim of this study is to investigate how different kinds of training modalities affect the production of phosphate regulating proteins and 1α-hydroxylation by rat bone tissue \textit{in vivo}. 
Materials and Methods

Experimental design and training protocol

The animal experiment was approved by the Animal Experiment Committee of the VU University Amsterdam with permit number FBW 10-03, and described previously (24). Briefly, 38 female Wistar rats at the age of 13 weeks were assigned to four groups: control (n=8), peak power training (PT, n=10), endurance training (ET, n=10) and a combined training (PET, n=10). The rats were trained on a treadmill for 6 weeks, 5 days a week for 1 session a day (ET and PT group). For the peak power training, rats performed 10 sprints of 15 sec in gallop at a maximal attainable velocity on a progressively increasing slope starting at 10% reaching up to 40% by the end. Endurance training consisted of treadmill running for 10 min at a speed of 16 m/min without a slope which was gradually increased up to 45 min of length with a speed of 26 m/min (trotting) on a 10% slope. The rats following a combination of training types (PET) conducted 2 training sessions a day: peak power training in the morning and endurance training 8 h later. Previously, an 8h rest period was shown sufficient to restore full mechanosensitivity (25). All rats were sacrificed 22 h after the last training by cardiac injection with Euthasol 20% (AST Farma B.V., Oude water, The Netherlands). The animals were group housed, rodent diet (Teklad Global 16% Protein Rodent Diet, Madison, WI, USA) containing 1% calcium, 0.7% phosphorus and 1.5 IU/g vitamin D3 was provided ad libitum and the rats were kept on a reversed 12 h light/dark cycle to provide the training during their active period of the day.

Immunohistochemistry

Immunohistochemistry was performed on the right tibial shaft to analyse protein expression of FGF23. Non-decalcified tibiae were fixed in 4% phosphate buffered
paraformaldehyde and after dehydration in increasing alcohol series, embedded in 80% methylmethacrylate (MMA) (BDH Chemicals, Poole, England) with 20% dibuthylphtalate (Merck, Darmstadt, Germany). Longitudinal five-micrometer-thick tissue sections were cut using a Polycut 2500 S microtome (Reichert-Jung, Nussloch, Germany). To remove MMA, tissue sections were incubated in 50% xylene and 50% chloroform. Subsequently, sections were rehydrated in a series of decreasing alcohol concentrations. Decalcification was done with 1% acidic acid, followed by quenching of endogenous peroxidases with 3% H2O2 in 40% methanol/PBS. Tissue sections were incubated with 10% goat serum to prevent unspecific binding of the secondary antibody. Subsequently, sections were incubated overnight at 4°C with polyclonal 1:200 rabbit anti-FGF23 (AB_2104625, Santa Cruz Biotechnology Sc-50291, CA, USA). The next day, tissue sections were incubated with polyclonal 1:100 biotinylated goat-anti-rabbit (AB_2313609, Dako E0432, Heverlee, Belgium) for 1 h and with 1:200 horseradish peroxidase labelled streptavidin (Invitrogen, Life Technologies, Bleiswijk, The Netherlands) for 1 h. Signal enhancement was established by treatment with tyramide (Invitrogen, Life Technologies, Bleiswijk, The Netherlands) for 10 min followed by a second 1 h incubation with 1:200 horseradish peroxidase labelled streptavidin. Chromogenesis was performed by treatment of the sections with AEC reagent (Invitrogen, Life Technologies, Bleiswijk, The Netherlands) for 6 min and by counterstaining with haematoxylin. Finally, the sections were mounted with ClearMount Mounting Solution (Invitrogen, Life Technologies, Bleiswijk, The Netherlands) and covered with a coverslip.

For each rat, two or three longitudinal tissue sections (depending on the quality of the tissue section), separated by 150 µm, were analysed using an Olympus BX51 Microscope at 200x magnification. The entire length of the tibial shaft was divided in alternating longitudinal regions
of interest of 50 µm long, in which all osteocytes across the entire thickness of the cortex were counted manually. Per tissue section, up to 400 osteocytes were counted. The amount of positively stained osteocytes was compared to the total number of osteocytes and expressed as fraction positively stained osteocytes compared to the total number of osteocytes.

**Quantitative polymerase chain reaction (qPCR)**

**Tissue preparation and RNA-extraction**

Left tibiae and kidneys were snap frozen and stored at -80°C until further analysis. One week before tibial RNA-isolation and 40 h prior to kidney RNA-isolation, tibiae and kidneys were stored in RNAlater® - ICE (Ambion, Life Technologies, Bleiswijk, The Netherlands). The kidneys were homogenised using a Dounce Homogeniser (Sigma-Aldrich, Zwijndrecht, The Netherlands). Subsequent processing occurred according to protocol from the column-based ‘FavorPrep Tissue Total RNA Purification Maxi Kit’ (Favorgen Biotech corp., Huissen, The Netherlands). For RNA isolation of the tibiae, proximal and distal ends of the tibiae were cut off. Bone marrow was removed by flushing the diaphysis with ice cold RNAse free water. Diaphyseal cortical bone was pulverised with a freezer-mill (SPEX 6750, Glen Creston, Stanmore, England) and incubated in Trizol (Invitrogen, Life Technologies, Bleiswijk, The Netherlands) for 1 h at 37°C. After the first Trizol-extraction, a chloroform-isomyl alcohol extraction was performed, followed by a second Trizol-extraction. Both kidney and tibial RNA were treated with DNAse (Promega, Leiden, The Netherlands) to remove DNA-contamination.
**Reversed Transcription**

After RNA-isolation, 100 ng of total RNA was reverse-transcribed using 10 ng/μl random primers (Roche, Almere, The Netherlands) and 5 U/μl M-MLV Reverse Transcriptase in a mixture with 5 mM MgCl₂, 1x RT-buffer, 1 mM dNTPs each, 1M betaine and 0.40 U/μl RNAsin (Promega, Leiden, The Netherlands). A total volume of 20 μl was incubated for 10 min at 25°C, 1 h at 37°C and 5 min at 95°C.

**Quantitative PCR**

A total volume of 25 μl containing 3 μl of cDNA, 1000 nM primers and 12,5 μl SYBR Green Supermix (Bio-Rad, Veenendaal, The Netherlands) was amplified in the iCycler system (Bio-Rad, Veenendaal, The Netherlands) using the primers as described in Table 1. PCR consisted of an initial denaturation step for 3 min at 95°C, followed by 40 amplification cycles (15 sec at 95°C, 1 min at 60°C). Subsequently, a melting curve was run from 50°C to 95°C to check the specificity of the reactions.

The following mRNA levels were assessed in the tibia: 1α-hydroxylase (CYP27b1), 24-hydroxylase (CYP24), vitamin D receptor (VDR), fibroblast growth factor 23 (FGF23), dental matrix protein 1 (DMP1), phosphate-regulating gene with homologies to endopeptidases on the X chromosome (PHEX), matrix extracellular phosphoglycoprotein (MEPE) and sclerostin (SOST).

The following mRNA levels were assessed in the kidney: CYP24, CYP27b1, VDR, NaPi-IIa cotransporter (NPT2a or SLC34a1) and NaPi-IIc cotransporter (NPT2c or SLC34a3).

Levels of tibial and kidney mRNA were expressed relative to the average of reference genes hypoxanthine phosphoribosyltransferase (HPRT) and porphobilinogendeaminase (PBGD)
using the $2^{\Delta CT}$ method. All samples were assessed in duplicate or triplicate on a single 96 well-plate per gene.

Serum Biochemical Analysis

Blood was obtained by puncture of the vena cava during general anaesthesia prior to sacrifice. Rat C-terminal FGF23 Elisa Kit (Immutopics, San Clemente, CA, USA) was used to measure FGF23 in EDTA plasma. Serum 25(OH)D3 was detected using liquid chromatography – tandem mass spectrometry (LC-MS/MS) according to standardised methods (26). Serum PTH was measured using the Rat Intact PTH Elisa Kit (Immutopics, San Clemente, CA, USA). Serum calcium and phosphate concentrations were measured using an Elecsys platform (Roche Diagnostics, Mannheim, Germany).

Statistical analysis

GraphPad Prism 6.0 (GraphPad Software Inc., La Jolla, CA, USA) was used for data analysis. Differences between the four groups were tested with a Kruskal-Wallis Test, followed by a Dunn’s post-hoc test. A value of $p<0.05$ was considered to be significant. Results are reported as means per group. Within figures, each dot represents the mean value of a single rat, whereas the bars represent the means per group.
Results

Animal experiment

Of the thirty rats that performed training, three rats (two of the PT group and one of the ET group) were excluded for not fulfilling the required training protocol. This resulted in a total sample size for controls n=8, ET n=9, PT n=8 and PET n=10. The final body weight of the included rats (n=35) did not differ among the groups. Tibial RNA-isolation for one control rat failed. We could not collect plasma samples for one control, three ET-rats, two PT-rats and one PET-rat.

FGF23 protein expression in tibiae

As shown in figure 1, immunohistochemical analysis of FGF23 protein expression by cortical osteocytes in the tibiae did not reveal any differences in the number of stained cells between the four groups. FGF23 staining in trabecular bone could unfortunately not be quantified, due to low contrast between positive osteocytes and aspecific staining of the marrow.

mRNA expression in tibiae

Figure 2A shows that FGF23 mRNA expression levels did not differ amongst the groups. mRNA levels of osteocyte maturation markers SOST, DMP1, PHEX and MEPE were significantly lower after PT compared to those after ET and PET, as illustrated by figures 2B-E, respectively. Furthermore, figures 2F and 2G show that CYP27b1 and VDR mRNA expression levels were lower after PT compared to those after ET. Expression of CYP24 mRNA was below detection limit.
mRNA expression in kidneys

Assuming that FGF23 production would be altered by physical exercise, we investigated mRNA expression levels of FGF23 responsive genes in the kidney to assess whether this would have systemic influences. Figure 3 reveals that there were no significant differences in CYP27b1, CYP24, VDR, NPT2a and NPT2c mRNA expression levels. Therefore, in kidney tissue physical exercise did not induce any changes in mRNA expression levels of FGF23-responsive genes and VDR.

Serum biochemical analysis

Figure 4 shows there were no significant differences in plasma c-term FGF23 concentration between groups. However, we did see an increase in serum PTH concentration in the PT-group compared to the controls. Systemic serum 25(OH)D3, calcium and phosphate levels were not significantly different between groups.
Discussion

This study aimed to investigate whether different kinds of training modalities affect phosphate homeostasis and 1α-hydroxylation in rat bone in vivo.

In contrast to our hypothesis, we observed no differences between groups in FGF23 expression in both bone and serum. In line with our results, FGF23 serum levels were not affected by either submaximal exercise or high intensity exercise with a bicycle ergometer among young men (27). Yet, Lombardi et al. described increased FGF23 serum concentrations among participants of the multiple-stage bicycle race Giro d’Italia (9). However, confounding factors in that study could be the high dietary intake as well as weight loss, suggesting the observed changes might be an effect of altered metabolic state rather than mechanical loading of bone cells. Moreover, these cyclists showed signs of induced bone resorption (28), possibly as a consequence of the heavy metabolic stress in the absence of mechanical loading, whereas we assume a situation of bone formation after mechanical loading in our model. Our findings also stand in contrast with those of Li et al., who reported increased serum FGF23 after acute exercise, exhaustive exercise, and chronic exercise among mice (8). However, these mice trained for only one week, whereas the rats in our model performed exercise for up to 6 weeks. As Li et al. also described that FGF23 promotes exercise endurance, it is possible the rats in our study at first responded to training with FGF23-upregulation to adapt to the exercise, but that after 6 weeks the rats were adapted and thus no longer showed increased FGF23 at the time of measurement.

Quantification of FGF23 substrates, such as genes involved in vitamin D metabolism and sodium-phosphate co-transporters in kidney tissue, ruled out that post-translational modification differed amongst groups. As FGF23 is considered the major phosphate regulating hormone (10),
it is not surprising that the lack of change in FGF23 or FGF23 substrates was paralleled with unchanged phosphate concentrations in serum after 6 weeks of training. Therefore, these results indicate that FGF23 production in bone and systemic phosphate homeostasis were not affected after 6 weeks of treadmill running.

Surprisingly, PT caused a decrease in both CYP27b1 and VDR mRNA levels in tibial bone compared to mRNA levels after ET, suggesting decreased vitamin D signalling in bone in the PT group. We had expected to observe an increase in local vitamin D signalling after PT compared to ET. Possibly, PT might have been too strenuous, causing fatigue damage of the bone leading to apoptosis of osteocytes(29). This process could also have accounted for the observed decreased vitamin D signalling after PT. In line with other studies (9,22), no differences between groups were observed for serum 25(OH)D concentrations, suggesting the observed decreased vitamin D signalling is rather a local process within bone than a systemic effect of exercise.

We hypothesized that alterations in FGF23 expression and 1α-hydroxylation in bone tissue in response to mechanical loading are related; both FGF23 expression and 1α-hydroxylation are likely to be altered by mechanical loading and both processes may be involved in regulation of each other. As we did not observe differences in FGF23 production but did observe decreased vitamin D signalling in bone tissue, we cannot confirm this hypothesis. Previous studies show contradicting results regarding this issue. In cultured rat osteoblasts FGF23 mRNA expression levels appeared to be enhanced by “autocrine/paracrine action of osteoblast-derived 1α,25(OH)2D” (12). In contrast, incubating cultured primary human osteoblasts with FGF23 did not cause changes in mRNA expression of CYP27b1 and VDR (20). Moreover, in human bone samples an association between FGF23 and CYP27b1, CYP24 or
VDR mRNA expression could not be shown (30). Possibly, in our rat model an acute change in FGF23 expression and 1α-hydroxylation in bone tissue may have occurred, but a new bone balance between the two was established at the time of investigation.

Furthermore, PTH levels were higher after PT compared to controls 22 hours after the last training. It is likely that the observed PTH concentrations after PT reflect an increased basal level of PTH in this group after six weeks of training. In general, a transient increase of PTH is observed during and directly after exercise, depending on the type of exercise (31), and most studies report normalization of PTH serum concentrations shortly after exercise (31–33), but some studies observed a prolonged increase in PTH 24 hours after a single exercise (34–36).

Low basal PTH concentrations are associated with higher physical fitness (34). Another interesting observation is that PT compared to ET and PET showed lower expression of osteocyte maturation markers SOST, DMP1, PHEX and MEPE. Therefore, it seems that PT caused a suppression of mature osteocyte function, which was reversed by additional endurance training. In general, DMP1 and MEPE are known to increase after mechanical loading (14–16), whereas SOST is downregulated by mechanical loading (37,38).

DMP1, MEPE and PHEX are not only markers of osteocyte maturation, but are functionally involved in phosphate regulation and bone mineralization (39). DMP1 and PHEX are known to be involved in inhibition of FGF23 production (13). Therefore, it was expected that reduction of these factors would lead to an increase in FGF23 production in the PT group in our rat model. However, after 6 weeks of peak training FGF23 production and function in systemic phosphate homeostasis appeared unchanged, which does not exclude that local changes in phosphate regulation in bone tissue may have occurred after PT. Moreover, it has been suggested that DMP1 and MEPE cause local changes in stiffness and mineralization of canaliculi and lacunae,
thereby altering the osteocytic response to mechanical loading (40). Possibly a new balance has
been established between DMP1, MEPE and FGF23 after 6 weeks of peak training, in which
FGF23 carries out its systemic functions whereas DMP1 and MEPE have a local function in
lacunar remodelling.

Local mechano-response in bone has the potential to lead to systemic effects, because
bone is an endocrine organ (41), and changes in bone resorption and formation can lead to
alterations in systemic calcium and phosphate. However, in comparison to the many local
mechano-responsive changes in tibial bone in this rat model, the only observed systemic effect
was an alteration of PTH concentrations after PT. PTH concentrations respond to many other
stimuli than just mechanical loading, such as changes in the adrenergic system, pH and or lactic
acid (42). As neither systemic FGF23, 25(OH)D3, calcium and phosphate, nor any of the genes
measured in kidney tissue differed between groups, it seems that the effects of mechanical
loading in this model did not lead to major systemic differences between groups.

Remarkably, we did not observe any statistically significant effect of PET compared to
controls or ET. Therefore, it seems that the observed changes in gene and protein expression
after PT have been reversed by additional ET. These results should be interpreted with caution;
the rats performing PET had more training than the other groups. By performing both training
programs, they underwent a higher amount of strain, which might be a confounding factor.

In this study organs were harvested 22 hours after the last training session in order to
study contractile muscle force characteristics in situ (24). Therefore, acute effects of mechanical
loading on mRNA concentrations might have already disappeared at the time of investigation.
The observed differences after PT might indicate a possible new steady state.
Another limiting factor to our study is the lack of accurate FGF23-staining in trabecular bone. Trabecular bone is generally considered to be metabolic active and possible effects of exercise on FGF23 protein expression within bone marrow could unfortunately not be quantified.
Conclusion

In conclusion, six weeks of physical exercise did not cause significant differences in phosphate homeostasis on a systemic level. However, mRNA expression levels of genes involved in local phosphate regulation and vitamin D signalling within bone tissue differed among groups, suggesting a local mechano-response rather than a systemic response in this rat model.
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1. Klein-Nulend J, Bacabac RG, Bakker AD. Mechanical loading and how it affects bone cells: The role of the osteocyte cytoskeleton in maintaining our skeleton. Eur Cells Mater. 2012;24:278–91.

2. Rubin CT, Lanyon LE. Regulation of bone mass by mechanical strain magnitude. Calcif Tissue Int. 1985;37(4):411–7.

3. Turner CH, Owan I, Takano Y. Mechanotransduction in bone: role of strain rate. Am J Physiol. 1995 Sep;269(3 Pt 1):E438-42.

4. Martyn-St James M, Carroll S. A meta-analysis of impact exercise on postmenopausal bone loss: the case for mixed loading exercise programmes. Br J Sports Med. 2009;43:898–908.

5. Sapir-Koren R, Livshits G. Osteocyte control of bone remodeling: is sclerostin a key molecular coordinator of the balanced bone resorption-formation cycles? Osteoporos Int. 2014;25(12):2685–2700.

6. Gardinier JD, Al-Omaishi S, Morris MD, Kohn DH. PTH Signaling Mediates Perlacon Remodeling During Exercise. Matrix Biol. 2016;(52–54):162–75.

7. Qi Z, Liu W, Lu J. The mechanisms underlying the beneficial effects of exercise on bone remodeling: Roles of bone-derived cytokines and microRNAs. Prog Biophys Mol Biol. Elsevier Ltd; 2016;122(2):131–9.

8. Li DJ, Fu H, Zhao T, Ni M, Shen FM. Exercise-stimulated FGF23 promotes exercise performance via controlling the excess reactive oxygen species production and enhancing mitochondrial function in skeletal muscle. Metabolism. Elsevier Inc.; 2016;65(5):747–56.

9. Lombardi G, Corsetti R, Lanteri P, Grasso D, Vianello E, Marazzi M, et al. Reciprocal regulation of calcium-/phosphate-regulating hormones in cyclists during the Giro d’Italia 3-week stage race. Scand J Med Sci Sports. 2014;24:779–87.

10. Bonewald LF, Wacker MJ. FGF23 Production by Osteocytes. Pediatr Nephrol. 2013;28(4):563–8.

11. Saji F, Shigematsu T, Sakaguchi T, Ohya M, Orita H, Maeda Y, et al. Fibroblast growth factor 23 production in bone is directly regulated by 1{alpha},25-dihydroxyvitamin D, but not PTH. Am J Physiol Renal Physiol. 2010 Nov;299:F1212–7.

12. Tang WJ, Wang LF, Xu XY, Zhou Y, Jin WF, Wang HF, et al. Autocrine/paracrine action of vitamin D on FGF23 expression in cultured rat osteoblasts. Calcif Tissue Int. 2010;86(5):404–10.

13. Martin A, Liu S, David V, Li H, Karydis A, Feng JQ, et al. Bone proteins PHEX and DMP1 regulate fibroblastic growth factor Fgf23 expression in osteocytes through a common pathway involving FGF receptor (FGFR) signaling. FASEB J. 2011;25(8):2551–62.

14. Gluhak-Heinrich J, Ye L, Bonewald LF, Feng JQ, MacDougall M, Harris SE, et al. Mechanical loading stimulates dentin matrix protein 1 (DMP1) expression in osteocytes in vivo. J bone Miner Res. 2003 May;18(5):807–17.
15. Yang W, Lu Y, Kalajzic I, Guo D, Harris MA, Gluhak-Heinrich J, et al. Dentin matrix protein 1 gene cis-regulation: use in osteocytes to characterize local responses to mechanical loading in vitro and in vivo. J Biol Chem. 2005 May 27;280(21):20680–90.

16. Reijnders CMA, Van Essen HW, Van Rens BTTM, Van Beek JHGM, Ylstra B, Blankenstein MA, et al. Increased expression of matrix extracellular phosphoglycoprotein (MEPE) in cortical bone of the rat tibia after mechanical loading: Identification by oligonucleotide microarray. PLoS One. 2013;8(11):1–12.

17. Lips P. Vitamin D physiology. Prog Biophys Mol Biol. 2006 Sep;92:4–8.

18. van Driel M, Koedam M, Buurman CJ, Hewison M, Chiba H, Uitterlinden AG, et al. Evidence for auto/paracrine actions of vitamin D in bone: 1alpha-hydroxylation expression and activity in human bone cells. FASEB J. 2006 Nov;20:2417–9.

19. van der Meijden K, Lips P, van Driel M, Heijboer AC, Schulten EAJM, den Heijer M, et al. Primary Human Osteoblasts in Response to 25-Hydroxyvitamin D3, 1,25-Dihydroxyvitamin D3 and 24R,25-Dihydroxyvitamin D3. PLoS One. 2014;9(10):1–11.

20. van der Meijden K, Essen HW, Bloemers FW, Schulten EAJM, Lips P, Bravenboer N. Regulation of CYP27B1 mRNA Expression in Primary Human Osteoblasts. Calcif Tissue Int. Springer US; 2016;99(2):164–73.

21. van der Meijden K, Bakker AD, van Essen HW, Heijboer AC, Schulten EAJM, Lips P, et al. Mechanical loading and the synthesis of 1,25(OH)2D in primary human osteoblasts. J Steroid Biochem Mol Biol. Elsevier Ltd; 2016;156:32–9.

22. Lester ME, Urso ML, Evans RK, Pierce JR, Spiering BA, Maresh CM, et al. Influence of exercise mode and osteogenic index on bone biomarker responses during short-term physical training. Bone. 2009;45:768–76.

23. Iwamoto J, Shimamura C, Takeda T, Abe H, Ichimura S, Sato Y, et al. Effects of treadmill exercise on bone mass, bone metabolism, and calciotropic hormones in young growing rats. J Bone Miner Metab. 2004;22:26–31.

24. Furrer R, Jaspers RT, Baggerman HL, Bravenboer N, Lips P, de Haan A. Attenuated increase in maximal force of rat medial gastrocnemius muscle after concurrent peak power and endurance training. Biomed Res Int. 2013 Jan;2013:935671.

25. Robling AG, Burr DB, Turner CH. Recovery periods restore mechanosensitivity to dynamically loaded bone. J Exp Biol. 2001;204:3389–99.

26. van der Meijden K, Bravenboer N, Dirks NF, Heijboer AC, den Heijer M, de Wit GMJ, et al. Effects of 1,25(OH)2D3 and 25(OH)D3 on C2C12 Myoblast Proliferation, Differentiation, and Myotube Hypertrophy. J Cell Physiol. 2016;231(11):2517–28.

27. Emrich IE, Baier M, Zawada AM, Meyer T, Fliser D, Scharhag J, et al. Plasma FGF23 does not rise during physical exercise as a physiological model of sympathetic activation. Clin Res Cardiol. 2018;

28. Lombardi G, Lanteri P, Graziani R, Colombini A, Banfi G, Corsetti R. Bone and Energy Metabolism Parameters in Professional Cyclists during the Giro d’Italia 3-weeks Stage Race. PLoS One. 2012;7(7):e42077.

29. Jilka RL, Noble B, Weinstein RS. Osteocyte apoptosis. Bone. 2013;54(2):264–71.
30. Ormsby RT, Findlay DM, Kogawa M, Anderson PH, Morris HA, Atkins GJ. Analysis of vitamin D metabolism gene expression in human bone: Evidence for autocrine control of bone remodelling. J Steroid Biochem Mol Biol. Elsevier Ltd; 2013;144:110–3.

31. Maimoun L, Sultan C. Effect of Physical Activity on Calcium Homeostasis and Calciotropic Hormones: A Review. Calcif Tissue Int. 2009;85:277–86.

32. Scott JPR, Sale C, Greeves JP, Casey A, Dutton J, Fraser WD. The role of exercise intensity in the bone metabolic response to an acute bout of weight-bearing exercise. J Appl Physiol. 2011;110:423–32.

33. Gardinier JD, Mohamed F, Kohn DH. PTH Signaling During Exercise Contributes to Bone Adaptation. J Bone Miner Metab. 2015;30(6):1053–63.

34. Brahm H, Piehl-Aulin K, Ljunghall S. Bone Metabolism During Exercise and Recovery: The Influence of Plasma Volume and Physical Fitness. Calcif Tissue Int. 1997;61:192–8.

35. Bouassida A, Zellag D, Zaouali Ajina M, Gharbi N, Duclos M, Richelet JP, et al. Parathyroid hormone concentrations during and after two periods of high intensity exercise with and without an intervening recovery period. Eur J Appl Physiol. 2003;88:339–44.

36. Thorsen K, Kristoffersson A, Hultdin J, Lorentzon R. Effects of Moderate Endurance Exercise on Calcium, Parathyroid Hormone, and Markers of Bone Metabolism in Young Women. Calcif Tissue Int. 1997;60:16–20.

37. Robling AG, Niziolek PJ, Baldrige LA, Condon KW, Allen MR, Alam I, et al. Mechanical stimulation of bone in vivo reduces osteocyte expression of Sost/sclerostin. J Biol Chem. 2008 Feb 29;283(9):5866–75.

38. Gardinier JD, Al-omaishi S, Morris MD, Kohn DH. PTH signaling mediates perilacunar remodeling during exercise. Matrix Biol. International Society of Matrix Biology; 2016;52–54:162–75.

39. Rowe PSN. Regulation of Bone-Renal Mineral and Energy Metabolism: The PHEX, FGF23, DMP1, MEPE ASARM Pathway. Crit Rev Eukaryot Gene Expres. 2012;22(1):61–86.

40. Harris SE, Gluhak-Heinrich J, Harris MA, Yang W, Bonewald LF, Riha D, et al. DMP1 and MEPE expression are elevated in osteocytes after mechanical loading in vivo: Theoretical role in controlling mineral quality in the perilacunar matrix. J Musculoskelet Neuronal Interact. 2007;7(4):313–5.

41. Han Y, You X, Xing W, Zhang Z, Zou W. Paracrine and endocrine actions of bone - The functions of secretory proteins from osteoblasts, osteocytes, and osteoclasts. Bone Res. Springer US; 2018;6(16):1–11.

42. Bouassida A, Latiri I, Bouassida S, Zalleg D, Zaouali M, Feki Y, et al. Parathyroid hormone and physical exercise: a brief review. J Sports Sci Med. 2006;5:367–74.
Supporting information captions

Table 1. Details of primers used for quantitative PCR analysis. CYP24: 24-hydroxylase; CYP27b1: 1α-hydroxylase; DMP1: dental matrix protein 1; FGF23: fibroblast growth factor 23; HPRT: hypoxanthine phosphoribosyltransferase; MEPE: matrix extracellular phosphoglycoprotein; NPT2a: NaPi-IIa cotransporter; NPT2c: NaPi-IIc cotransporter; PBGD: porphobilinogendeaminase; PHEX: phosphate-regulating gene with homologies to endopeptidases on the X chromosome; SOST: sclerostin; VDR: vitamin D receptor.

Figure 1. Effects of different training modalities (ET: endurance training; PT: peak training; PET: combined peak- and endurance training) on the expression of FGF23 in rat tibiae as analysed by immunohistochemistry. (A) Fraction positively stained cortical osteocytes compared to total amount of osteocytes. (B) Representative image of FGF23 positively stained cortical osteocytes after PET. Arrows show positively stained osteocytes, whereas arrowheads point out FGF23 negative osteocytes. Bar represents 25 µm.

Figure 2. Effects of different training modalities (ET: endurance training; PT: peak training; PET: combined peak- and endurance training) on mRNA expression in rat tibiae as analysed by qPCR. Gene expression levels of (A) FGF23, (B) SOST, (C) DMP1, (D) PHEX, (E) MEPE, (F) CYP27b1 and (G) VDR were measured. Results were normalized for reference genes HPRT and PBGD. Significant differences are indicated as *p<0.05 and **p<0.01.

Figure 3. Effects of different training modalities (ET: endurance training; PT: peak training; PET: combined peak- and endurance training) on mRNA expression in rat kidneys as analysed...
by qPCR. Gene expression levels of (A) CYP27b1, (B) CYP24, (C) VDR, (D) NPT2a and (E) NPT2c were measured. Results were normalized for reference genes HPRT and PBGD.

Significant difference is indicated as *p<0.05.

**Figure 4.** Effects of different training modalities on serum concentrations of factors involved in phosphate homeostasis. Serum concentrations of (A) c-terminal FGF23, (B) 25(OH)D3, (C) parathyroid hormone (PTH), (D) calcium and (E) phosphate were assessed. Significant difference is indicated as *p<0.05.
Figure 1 (on next page)

Effects of different training modalities on the expression of FGF23 in rat tibiae as analysed by immunohistochemistry.

(A) Fraction positively stained cortical osteocytes compared to total amount of osteocytes. (B) Representative image of FGF23 positively stained cortical osteocytes after PET. Arrows show positively stained osteocytes, whereas arrowheads point out FGF23 negative osteocytes. Bar represents 25 µm.

(ET: endurance training; PT: peak training; PET: combined peak- and endurance training)
Figure 2 (on next page)

Effects of different training modalities on mRNA expression in rat tibiae as analysed by qPCR.

Gene expression levels of (A) FGF23, (B) SOST, (C) DMP1, (D) PHEX, (E) MEPE, (F) CYP27b1 and (G) VDR were measured. Results were normalized for reference genes HPRT and PBGD. Significant differences are indicated as *p<0.05 and **p<0.01.

(ET: endurance training; PT: peak training; PET: combined peak- and endurance training)
Manuscript to be reviewed
**Figure 3** (on next page)

Effects of different training modalities on mRNA expression in rat kidneys as analysed by qPCR.

Gene expression levels of (A) CYP27b1, (B) CYP24, (C) VDR, (D) NPT2a and (E) NPT2c were measured. Results were normalized for reference genes HPRT and PBGD. Significant difference is indicated as *p<0.05.

(ET: endurance training; PT: peak training; PET: combined peak- and endurance training)
Figure 1: Gene expression levels of different genes under different treatments.

A. CYP27b1, comparing Control, ET, PT, and PET conditions.

B. CYP24, comparing Control, ET, PT, and PET conditions.

C. VDR, comparing Control, ET, PT, and PET conditions.

D. NPT2a, comparing Control, ET, PT, and PET conditions.

E. NPT2c, comparing Control, ET, PT, and PET conditions.
Figure 4 (on next page)

Effects of different training modalities on serum concentrations of factors involved in phosphate homeostasis.

Serum concentrations of (A) c-terminal FGF23, (B) 25(OH)D3, (C) parathyroid hormone (PTH), (D) calcium and (E) phosphate were assessed. Significant difference is indicated as *p<0.05.
Table 1 (on next page)

Details of primers used for quantitative PCR analysis.

CYP24: 24-hydroxylase; CYP27b1: 1α-hydroxylase; DMP1: dental matrix protein 1; FGF23: fibroblast growth factor 23; HPRT: hypoxanthine phosphoribosyltransferase; MEPE: matrix extracellular phosphoglycoprotein; NPT2a: NaPi-IIa cotransporter; NPT2c: NaPi-IIc cotransporter; PBGD: porphobilinogen deaminase; PHEX: phosphate-regulating gene with homologies to endopeptidases on the X chromosome; SOST: sclerostin; VDR: vitamin D receptor.
### Table 1.

**PCR primer sequences 5' --> 3'**

| Target gene | Forward | Reverse |
|-------------|---------|---------|
| CYP24       | GCT-GAT-GAC-AGA-CGG-TGA-GA | TGT-CGT-GCT-GTT-TCT-TCA-GG |
| CYP27b1     | CCC-GAC-ACA-GAA-ACC-TTC-AT | GGC-AAA-CAT-CTG-ATC-CCA-GT |
| DMP1        | GCG-ACT-CCA-CAG-AGG-ATT-TC | GTC-CCT-CTG-GGC-TAT-CTT-CC |
| FGF23       | GAT-GCT-GGC-TCC-GTA-GTG-AT | CGT-CGT-AGC-CTG-TCT-CTA-GC |
| HPRT        | GTG-TCA-TCA-GCG-AAA-GTG-GA | TAC-TGG-CCA-CAT-CAA-CAG-GA |
| MEPE        | AAG-ACA-AGC-CAC-CCT-ACA-CG | CCC-ACT-GGA-TGA-TGA-CTC-ACT |
| NPT2a       | AGT-GGC-CAA-TGT-CAT-CCA-GA | AGT-GAT-GGC-TGA-GGT-GAA-CA |
| NPT2c       | GGT-CAC-CGT-CCT-TGT-ACA-GA | GAC-GCC-CAT-GAT-GAT-AGG-GA |
| PBGD        | ATG-TCC-GGT-AAC-GGC-GGC | CAA-GGT-TTT-CAG-CAT-CGC-TAC-CA |
| PHEX        | CAG-GCA-TCA-CAT-TCA-CCA-AC | GGA-GGA-CTG-TGA-GCA-CCA-AT |
| SOST        | CAG-CTC-TCA-CTA-GCC-CCT-TG | GGG-ATG-ATT-TCT-GTG-GCA-TC |
| VDR         | ACA-GTC-TGA-GGC-CCA-AGC-TA | TCC-CTG-AAG-TCA-GCG-TAG-GT |