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## Skeletal secretion of FGF-23 regulates phosphate and vitamin D metabolism

L. Darryl Quarles

Division of Nephrology, Department of Medicine, University of Tennessee HealthScience Center, 19 South Manassas Street, Memphis, TN 38163, USA

L. Darryl Quarles: dqarles@uthsc.edu

### Abstract

The discovery of fibroblast growth factor 23 (FGF-23) has expanded our understanding of phosphate and vitamin D homeostasis and provided new insights into the pathogenesis of hereditary hypophosphatemic and hyperphosphatemic disorders, as well as acquired disorders of phosphate metabolism, such as chronic kidney disease. FGF-23 is secreted by osteoblasts and osteocytes in bone and principally targets the kidney to regulate the reabsorption of phosphate, the production and catabolism of 1,25-dihydroxyvitamin D and the expression of  $\alpha$ -Klotho, an anti-ageing hormone. Secreted FGF-23 plays a central role in complex endocrine networks involving local bone-derived factors that regulate mineralization of extracellular matrix and systemic hormones involved in mineral metabolism. Inactivating mutations of *PHEX*, *DMP1* and *ENPP1*, which cause hereditary hypophosphatemic disorders and primary defects in bone mineralization, stimulate *FGF23* gene transcription in osteoblasts and osteocytes, at least in part, through canonical and intracrine FGF receptor pathways. These FGF-23 regulatory pathways may enable systemic phosphate and vitamin D homeostasis to be coordinated with bone mineralization. FGF-23 also functions as a counter-regulatory hormone for 1,25-dihydroxyvitamin D in a bone–kidney endocrine loop. FGF-23, through regulation of additional genes in the kidney and extrarenal tissues, probably has broader physiological functions beyond regulation of mineral metabolism that account for the association between FGF-23 and increased mortality and morbidity in chronic kidney disease.

### Introduction

In the past decade, bone has been recognized to be an endocrine organ that releases at least two hormones into the circulation, osteocalcin and fibroblast growth factor 23 (FGF-23). Osteocalcin is proposed to participate in an endocrine axis whereby bone regulates energy metabolism. This axis is initiated by the systemic release of undercarboxylated osteocalcin from bone resorption, which regulates insulin secretion, insulin sensitivity and energy expenditure by the activation of GPRC6A, a G-protein coupled receptor located in  $\beta$  cells and other target tissues.<sup>1–3</sup> A second endocrine axis, which involves the release of the

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hormone FGF-23 from osteoblasts and osteocytes, regulates systemic phosphate homeostasis and vitamin D metabolism.<sup>4</sup> This Review focuses on the rapidly growing knowledge of this regulation by FGF-23.<sup>5</sup>

FGF-23, along with FGF-19 and FGF-21, is a member of the FGF family, which are released into the circulation owing to their low binding affinity for heparin. These proteins act as endocrine factors by virtue of an evolutionarily engineered C-terminus that permits interaction with FGF receptor- $\alpha$ -Klotho co-receptor complexes in cell membranes of target tissues. These hormonal FGFs differ from canonical FGFs that have high heparin affinity and act as extracellular autocrine or paracrine ligands for nearby cell-surface membrane FGF receptors.<sup>5</sup> They also differ from intracellular, nuclear isoforms of FGFs, as exemplified by high-molecular weight (HMW)-FGF-2, which interact with intranuclear FGFR-1 to directly activate gene transcription.<sup>6</sup> FGF-19 regulates bile acid metabolism in the liver, whereas FGF-21 regulates lipid metabolism in white adipose tissue.

Our understanding of the role of FGF-23 is changing the traditional view of the regulation of several physiological processes, including bone mineralization, phosphate homeostasis and vitamin D metabolism. Endocrine networks that have become apparent from the unveiling of FGF-23 regulation and functions are also modifying our categorization of hereditary hypophosphatemic and hyperphosphatemic disorders. Furthermore, this new knowledge is challenging how several diseases are conceptualized, including the definition of vitamin D deficiency, the pathogenesis of secondary hyperparathyroidism in chronic kidney disease (CKD) and the relationship between disordered mineral metabolism and cardiovascular mortality.

## Overview of FGF-23

FGF-23 is a ~32 kDa protein with an N-terminal FGF homology domain and a novel 72-amino-acid C-terminus.<sup>7</sup> Both osteoblasts and osteocytes produce and secrete FGF-23.<sup>7-10</sup> Circulating FGF-23 binds to and activates receptor complexes consisting of FGFR-1, FGFR-3 or FGFR-4 and the transmembrane  $\beta$  glucuronidase  $\alpha$ -Klotho,<sup>11-14</sup> which is located in target tissues. Excess FGF-23 in both humans and mouse models causes hypophosphatemia, suppression of 1,25-dihydroxyvitamin D levels and rickets or osteomalacia.<sup>15-19</sup>

The major target for FGF-23 is the kidney, where increments in FGF-23 inhibit renal phosphate reabsorption by decreasing the expression and membrane insertion of Na<sup>+</sup>-dependent co-transporters. In the kidney, excess FGF-23 also suppresses circulating levels of 1,25-dihydroxyvitamin D by inhibiting the enzyme CYP27B1 (which converts 25-hydroxyvitamin D to 1,25-dihydroxyvitamin D) and possibly by stimulating the catabolism of 1,25-dihydroxyvitamin D by activating 24-hydroxylase (CYP24) in the proximal tubule.<sup>15-16,20-24</sup> Elevated FGF-23 is associated with reduced  $\alpha$ -Klotho expression and preliminary studies suggest that FGF-23 directly suppresses  $\alpha$ -Klotho message expression in the distal tubule.<sup>25,26</sup> Many unanswered questions exist regarding FGF-23 effects on specific tubular segments (proximal and/or distal), the specific FGFR receptors mediating its effects (that is, FGFR-1, FGFR-3 and/or FGFR-4) and the full complement of gene products

directly and/or indirectly regulated by FGF-23 in the kidney,<sup>14,27</sup> By contrast, reductions in FGF-23 in both humans and mouse genetic models are known to cause tumoral calcinosis, characterized by hyperphosphatemia, increased 1,25-dihydroxyvitamin D and soft tissue calcifications.<sup>9,20,28–32</sup>

## Mutations that elevate FGF-23 levels

The study of hypophosphatemic disorders, which are caused by eight distinct gene mutations that result in a common phenotype,<sup>27</sup> has contributed to our understanding of the regulation and function of FGF-23 (Table 1). Study of autosomal dominant hypophosphatemic rickets, which is caused by mutations in an RXXR site in FGF-23 that prevents its cleavage, helped to define the functional role of FGF-23 as an essential hormone that regulates phosphate and vitamin D metabolism. Comparative analysis of the other seven mutations associated with the hypophosphatemic disorders X-linked hypophosphatemic rickets, autosomal dominant hypophosphatemic rickets, autosomal recessive hypophosphatemic rickets 1, autosomal recessive hypophosphatemic rickets 2, osteoglophonic dysplasia, McCune–Albright syndrome and Jansen metaphyseal chondrodysplasia provided insights into local and systemic regulation of *FGF23* gene transcription in osteoblasts and osteocytes (Table 1).<sup>4,27,33</sup> In addition to these disorders, epidermal nevus, which can result from somatic mosaicism for mutations in the *FGFR3* gene or the *PIK3CA* gene,<sup>34</sup> is associated with increased circulating FGF-23 levels.<sup>35</sup> Opsismodysplasia, a rare hereditary spondylo(epi) chondrodysplasia, is also characterized by hypophosphatemia, abnormal vitamin D metabolism and elevated FGF-23 levels, but the disease-causing gene mutation has not been identified.<sup>36</sup> Elevated FGF-23 levels also cause acquired hypophosphatemic disorders, such as tumor-induced osteomalacia, and are an important adaptive response in CKD.<sup>37</sup>

## Mutations causing impaired mineralization

Why is FGF-23 produced predominately by osteoblasts and osteocytes in bone? Studies of X-linked hypophosphatemia and autosomal recessive hypophosphatemic rickets in both humans and mouse models indicate that extracellular matrix mineralization in bone is linked to renal handling of phosphate through the release of FGF-23. X-linked hypophosphatemia in humans,<sup>27</sup> and the corresponding hypophosphatemic (Hyp) mouse model, are caused by mutations of the *PHEX* gene, which encodes the endopeptidase PHEX (also known as HYP).<sup>9,38</sup> The *Phex* gene in mice is highly expressed in differentiated osteoblasts and osteocytes, and conditional deletion of *Phex* in a mature osteoblast lineage *in vivo* in mice is sufficient to reproduce the Hyp phenotype.<sup>39</sup> Although an initial study suggested that PHEX metabolizes FGF-23,<sup>40</sup> subsequent studies failed to establish PHEX-dependent cleavage of FGF-23 *in vitro*.<sup>8,41,42</sup> Rather, *Phex* mutations lead to elevations of FGF-23 levels owing to increased *Fgf23* gene transcription in osteoblasts and osteocytes in mice.<sup>43</sup> In this regard, increased *Fgf23* gene transcription is present in isolated osteoblasts and bone marrow stromal cell cultures that differentiate into osteoblasts derived from Hyp mice and in Hyp bone explanted to wild-type mice.<sup>38</sup>

Autosomal recessive hypophosphatemic rickets 1, caused by inactivating mutations of *DMP1* in humans<sup>44</sup> and the corresponding *Dmp1*<sup>-/-</sup> mouse model,<sup>45</sup> results in intrinsic stimulation of FGF-23 expression in osteoblasts and osteocytes. DMP-1 is a SIBLING

(small integrin-binding ligand, N-linked glycoprotein) extracellular matrix protein that acts as a nucleator of mineralization and activates signaling pathways in osteoblasts and osteocytes via extracellular matrix–cell-surface interactions.<sup>46</sup> DMP-1 is cleaved *in vitro* into 37 kDa and 57 kDa fragments by BMP-1<sup>47</sup> and MMP-2.<sup>48</sup> The NH<sub>2</sub>-terminal fragment is a proteoglycan with a chondroitin sulfate chain attached at serine position 74 that binds to pro-MMP-9 and might sequester growth factors,<sup>49</sup> whereas the C-terminal fragment has an RGD recognition sequence for binding to integrins and an ASARM motif that may mediate binding to PHEX. *Phex* and *Dmp1* mutations in mice result in impaired mineralization that is independent of the concomitant hypophosphatemia.<sup>38,45</sup> This impairment may be related to PHEX binding to ASARM peptides that are located within SIBLING proteins, such as MEPE<sup>44</sup> and DMP-1, which act to inhibit mineralization.<sup>50</sup> Nonadditive effects on FGF-23 expression are observed in compound mutant *Hyp/Dmp1*<sup>-/-</sup> mice, which suggests that common intrinsic abnormalities of mineralization and/or alterations of the extracellular matrix milieu are coupled to FGF-23 expression.<sup>4,51–53</sup>

Two additional genetic abnormalities regulating pyrophosphate metabolism also support a coupling of mineralization with FGF-23 expression in mouse models. Inorganic pyrophosphate, an inhibitor of calcification, is generated by the enzyme E-NPP1 and transported by ANK-1 into the extracellular matrix environment, where it is converted to the inorganic phosphate required for matrix mineralization by the alkaline phosphatase enzyme TNAP, which is expressed in osteoblasts. Inactivating mutations of *ENPP1*, which cause hereditary generalized arterial calcification of infancy, also cause autosomal recessive hypophosphatemic rickets 2, which is characterized by FGF-23-mediated hypophosphatemia.<sup>54,55</sup> The mechanism whereby E-NPP1 increases FGF-23 expression is not known; however, the inactivation of E-NPP1 (which reduces pyrophosphate levels in soft tissues and causes ectopic calcification), also deprives TNAP of its substrate, which results in local phosphate depletion in bone that leads to osteomalacia and increased FGF-23 expression. In addition, inactivation of ANK-1, a pyrophosphate transporter located in osteoblasts, results in impaired mineralization of extracellular matrix and nearly a 10-fold increase in FGF-23 expression in bone in mice.<sup>56</sup>

In summary, mutations in the genes that encode DMP-1, PHEX, ANK-1 and E-NPP1, which block the mineralization of extracellular matrix, lead to increased FGF-23 production by osteoblasts and osteocytes.<sup>57,58</sup> These observations suggest a physiological need to coordinate bone mineralization and kidney handling of phosphate and vitamin D metabolism via the release of FGF-23.

### Mutations linking mineralization and FGF-23

A major knowledge gap exists for the mechanism whereby ‘intrinsic factors’ disrupting the process of bone mineralization (namely, by inhibiting E-NPP1, ANK-1, PHEX and DMP-1 functions) lead to the stimulation of FGF-23 gene expression in osteoblasts and osteocytes. However, study of additional single gene mutations in humans and mouse genetic models that cause FGF-23-mediated hypophosphatemia have identified an unexpected role of canonical and integrative nuclear FGFR-1 signaling in the regulation of FGF-23 expression

in bone; both of these types of FGF receptor signaling have been linked to *Phex* and *Dmp1* mutations and increased *Fgf23* gene transcription in mice.

In addition, osteoglophonic dysplasia, an autosomal dominant bone dysplastic disorder caused by activating mutations in FGFR-1, is associated with hypophosphatemia and elevated FGF-23 levels.<sup>59</sup> Several pieces of data support a role of FGFR-1 in the regulation of FGF-23 in bone. First, FGFR-1 is expressed in osteoblasts and osteocytes.<sup>60</sup> Second, both DMP-1 and PHEX act through common pathways regulating FGFR-1 activity to stimulate *Fgf23* gene transcription in osteoblast and osteocytes in compound mutant mouse models.<sup>61</sup> Third, pharmacological inhibition of FGFR-1 also inhibits *Fgf23* transcription in bone in animal models.<sup>62</sup> Fourth, the importance of integrative nuclear FGFR-1 signaling in activation of FGF-23 in bone is further supported by the observations that overexpression of HMW-FGF-2, the ligand for nuclear FGFR-1, in transgenic mice stimulates FGF-23 expression in bone and that HMW-FGF-2 is increased in the bone of adult Hyp mice.<sup>63</sup> Integrative nuclear FGFR-1 signaling activates the transcription factor CREB, which is present in the proximal *Fgf23* promoter, which suggests that the promoter contains a possible binding site for FGFR regulation of *FGF23* gene transcription. Finally, FGF2 administration *in vivo* to mice also induces hypophosphatemia and impairs matrix mineralization in mice.<sup>64,65</sup>

Thus, both canonical and integrative nuclear FGFR-1 pathways appear to be involved in regulating FGF-23 expression in bone. These observations have led to an organizing hypothesis that a physiological function of FGF-23 is to couple bone mineralization with FGF-23 production in osteoblasts and osteocytes via activation of FGFR-1 via yet-to-be defined mechanisms intrinsic to the bone milieu.<sup>27</sup> Regardless, the regulation of FGF-23 by FGFR-1 suggests that the generation of a hormonal FGF-23 from the ancestral FGF gene is an evolutionary adaptation to provide a mechanism to link paracrine actions of canonical FGFs to systemic effects.

Whereas mutations in the genes that encode DMP-1, PHEX, ANK-1 and E-NPP1 block bone mineralization and lead to increased FGF-23 production by osteocytes,<sup>57,58</sup> nutritional osteomalacia is associated with decreased FGF-23 expression and, paradoxically, further suppression of FGF-23 after treatment with vitamin D and healing of rachitic bone disease.<sup>66</sup> These findings suggest that defective mineralization *per se* (that is caused by vitamin D deficiency) is not sufficient to stimulate FGF-23 production in bone. Although the mechanisms of these disparate findings are not clear, differences in bone phosphate flux, which is increased during healing of vitamin D deficiency and decreased in defective mineralization caused by *PHEX*, *DMP1*, *ANK1* and *ENPP1* mutations, might account for these different responses. Also, bone remodeling might regulate FGF-23 expression. In this regard, inhibition of bone turnover mediated by osteoprotegerin or alendronate increases FGF-23 expression.<sup>67</sup> The mechanism mediating the effects of bone remodeling on FGF-23 expression is also not clear, but alterations in phosphate influx into and out of bone could also be involved. Parathyroid hormone (PTH), leptin, estrogens and glucocorticoids also help to coordinate the regulation of FGF-23 and bone remodeling,<sup>68</sup> which suggests the possible presence of more-complex endocrine networks involving regulation of FGF-23 and energy metabolism (see below).

### FGF-23 regulation by PTH signaling mutations

Jansen metaphyseal chondrodysplasia, which is caused by a mutation in the *PTH1R* gene that renders the corresponding PTH/PTHr receptor constitutively active, is associated with high FGF-23 concentrations in the circulation and low serum phosphate and inappropriately normal 1,25-dihydroxyvitamin D<sub>3</sub> levels.<sup>69</sup> Transgenic mice with constitutive activation of PTH receptor signaling in osteocytes exhibit increased bone mass and remodeling and increased circulating FGF-23 levels.<sup>70</sup> McCune–Albright syndrome, which is caused by activating mutations of the *GNAS* gene (which encodes the Gα<sub>s</sub> subunit that is coupled to the PTH receptor), also results in increased FGF-23 expression in the fibrodysplastic lesions that are characteristic of the syndrome.<sup>71</sup> Furthermore, serum levels of FGF-23 correlate with disease burden bone turnover markers in this syndrome. However, activating mutations of *GNAS* are necessary but not sufficient to increase FGF-23 expression, as evidenced by increased *FGF23* transcripts in only a subset of the fibrodysplastic lesions showing *GNAS* mutations.<sup>72</sup> Controversies regarding PTH regulation of FGF-23 are discussed below.

### Integrative physiology of FGF-23

Knowledge of how the FGF-23 bone–kidney axis is integrated with other hormonal networks is increasing, but the information is incomplete, and many gaps in knowledge remain. At present, FGF-23 is believed to be involved in at least three endocrine axes and preliminary data is emerging for crosstalk between the FGF-23 endocrine axes and other endocrine networks that regulate energy metabolism and aging.

#### FGF-23–vitamin D axis

FGF-23 is involved in a FGF-23–vitamin D endocrine loop (Figure 1). In this loop, 1,25-dihydroxyvitamin D stimulates FGF-23 production by bone through VDR-dependent mechanisms and elevated FGF-23 levels exert a negative feedback effect on the kidney to suppress 1,25-dihydroxyvitamin D expression.<sup>73,74</sup> Thus, FGF-23 acts as a counter-regulatory factor for 1,25-dihydroxyvitamin D.<sup>73</sup> Although FGF-23 is a phosphate-regulating hormone, like PTH, tight coupling between changes in circulating phosphate and FGF-23 concentrations are not present.<sup>73,75</sup> Some studies have failed to observe changes in FGF-23 levels in healthy individuals in response to either low or high phosphate diets.<sup>76,77</sup> By contrast, other studies have shown that alterations in dietary phosphate lead to changes in FGF-23 levels, albeit after a lag time of up to 1 week.<sup>78–83</sup> The effects of phosphate on FGF-23 levels are also modulated by vitamin D status, as increasing dietary phosphorus does not increase FGF-23 levels in the absence of the VDR in mouse models.<sup>15</sup> The related observation that FGF-23 levels are elevated in primary disorders of bone pyrophosphate or phosphate metabolism in *Enpp1* and *Ank1* mutant mice, raises the possibility that phosphate may indirectly regulate FGF-23 by affecting bone mineralization.<sup>73,75</sup>

#### PTH–vitamin D axis

Prior to the discovery of FGF-23, regulation of phosphate and vitamin D metabolism was understood in the context of the PTH–vitamin D endocrine loop, wherein PTH stimulates 1,25-dihydroxyvitamin D production that feeds back to the parathyroid gland to suppress PTH secretion through direct and indirect mechanisms (Figure 1). The primary function of



PTH is to maintain serum calcium levels in a narrow range. PTH secretion is stimulated by changes in serum calcium through activation of the calcium-sensing receptor in parathyroid chief cells. PTH, in turn, targets the kidney to decrease distal tubular calcium secretion and increase 1,25-dihydroxyvitamin D production by stimulating CYP27B1 activity and targets bone to increase calcium and phosphate efflux. PTH has phosphaturic actions that permit increments in serum calcium levels without elevations of serum phosphate levels, owing to 1,25-dihydroxyvitamin-D-mediated increases in calcium and phosphate absorption by the gastrointestinal tract.

PTH can indirectly stimulate FGF-23 serum levels by increasing 1,25-dihydroxyvitamin D synthesis, which in turn would directly stimulate FGF-23 production. The absence of parathyroid glands in *Gcm2*<sup>-/-</sup> mice results in low 1,25-dihydroxyvitamin D levels and decreased FGF-23 concentrations that are restored to normal by administration of 1,25-dihydroxyvitamin D supplements.<sup>73</sup> The essential role of 1,25-dihydroxyvitamin D in the regulation of FGF-23 expression is supported by the study of *Vdr*<sup>-/-</sup> mice, which have markedly elevated PTH levels but low circulating levels of FGF-23 in the absence of VDR-mediated gene transcription.<sup>15</sup>

### A PTH–FGF-23 axis?

Compelling but not incontrovertible evidence exists for a PTH–bone feedback loop, wherein PTH stimulates FGF-23 expression in bone and FGF-23 feeds back to the parathyroid gland to directly suppress PTH production.<sup>84</sup> In support of the efferent limb of the PTH–FGF-23 endocrine loop, activating mutations of the *PTH1R* and *GNAS* genes result in increased FGF-23 expression (Table 1).<sup>69</sup> FGF-23 concentration is also elevated in mice that overexpress a constitutively active PTH receptor in osteocytes and in some, but not all, animal models of excess PTH.<sup>85–87</sup> A strong association also exists between elevated FGF-23 levels and the severity of hyperparathyroidism in CKD and other disorders.<sup>12,88,89</sup> In addition, PTH increases FGF-23 expression and mediates the high FGF-23 levels observed in a CKD rat model fed a diet consisting of 0.75% adenine and 1.5% phosphate.<sup>86</sup> In these studies, early parathyroidectomy to reduce circulating PTH levels prevented the increase in FGF-23 levels observed in rats with adenine-induced renal failure. These investigators also found that continuous PTH administration at high doses (50 µg/kg per day) stimulated FGF-23 production in mice and also stimulated FGF-23 expression in cultured UMR-106 osteoblasts, an osteosarcoma-derived cell line. Finally, PTH has been shown to directly stimulate FGF-23 in UMR-106 osteoblasts<sup>86</sup> and primary osteoblasts, although the response in the later was transient.<sup>70,90</sup> In osteoblast culture models, PTH-dependent stimulation of FGF-23 was blocked by sclerostin overexpression, which suggests the involvement of Wnt signaling and/or alterations in bone remodeling.

Several findings, however, do not support the PTH–FGF-23 endocrine loop. In this regard, FGF-23 effects on the PTH secretion by the parathyroid gland are controversial.<sup>88,89</sup> FGFR-1 and α-Klotho are expressed in parathyroid chief cells, and FGF-23 directly suppresses *PTH* mRNA expression *in vitro* and decreases serum PTH *in vivo* in animal models,<sup>84</sup> consistent with a negative feedback afferent limb of a PTH–FGF-23 loop. However, elevated FGF-23 concentrations do not prevent the development of

hyperparathyroidism in various clinical disorders. Indeed, primary elevations in circulating FGF-23 levels in hereditary hypophosphatemic disorders as well as secondary increments in FGF-23 levels in CKD are both associated with increased, not decreased, PTH levels, suggesting that FGF-23 does not suppress PTH secretion.<sup>12</sup> Rather, the positive association between FGF-23 and PTH suggests that FGF-23 might promote the development of hyperparathyroidism.<sup>12,88,89</sup> The failure of FGF-23 to suppress PTH in CKD might also be explained by the resistance to FGF-23 in uremic parathyroid glands due to downregulation of  $\alpha$ -Klotho and FGFR expression.<sup>91,92</sup>

Several additional observations also fail to support direct effects of PTH on FGF-23 production by bone. First, important limitations exist to the mouse model of 'CKD', which involves high dietary phosphate and adenine intake, which can confound establishing cause-and-effect relationships between PTH and FGF-23 expression. Second, many studies fail to show that PTH directly stimulates FGF-23 production or FGF-23 promoter activity in osteoblasts *in vitro*,<sup>73</sup> or in calvarial cultures *ex vivo*.<sup>93</sup> Third, PTH administration has also been shown to either suppress or have no effect on FGF-23 expression in wild-type mice.<sup>67,93</sup> FGF-23 is also not elevated in patients with primary hyperparathyroidism.<sup>94</sup> Fourth, in studies showing a relationship between PTH and FGF-23 levels, the increase in FGF-23 could be indirect, owing to the stimulatory effects of PTH on 1,25-dihydroxyvitamin D levels.<sup>85,95,96</sup> Conversely, in settings of primary increments in FGF-23, such as occur in Hyp mice, elevations in PTH are likely to be mediated by FGF-23-mediated suppression of 1,25-dihydroxyvitamin D.<sup>96,97</sup> The mechanism underlying the variable effects of PTH on FGF-23 expression is not known, but insights into potential crosstalk between pathways known to regulate *Fgf23* gene transcription *in vivo* and integration of their downstream signaling pathways at the levels of the *Fgf23* promoter may explain the context-dependent effects of PTH as well as the molecular mechanism whereby bone mineralization regulates FGF-23 levels (Figure 1).

### Other possible FGF-23–hormonal axes

Given the importance of systemic phosphate homeostasis in intermediary metabolism, it is surprising that FGF-23 has not been linked to energy metabolism. Future elucidation of the novel bone–pancreas endocrine loop, wherein insulin receptors in osteoblasts stimulate bone turnover and release of undercarboxylated osteocalcin that regulates glucose homeostasis and energy balance, might identify new molecular pathways for FGF-23 regulation.<sup>3,97</sup> The speculation of a role of FGF-23 in energy metabolism is supported by evidence that the insulin-responsive PI3K–Akt–Sgk3 pathway regulates FGF-23 expression in bone in mice, with *Sgk3* knockout mice showing reductions in bone density, and 1,25-dihydroxyvitamin D and FGF-23 levels.<sup>98</sup>

Although many aspects of this bone–pancreas endocrine network remain controversial, peripheral actions of insulin to stimulate cellular phosphate uptake and bone actions to suppress FGF-23 might be of physiological importance. Indeed, insulin regulation of FGF-23 might lead to kidney retention of phosphate proportionate to the increased utilization of phosphate in peripheral tissues during enhanced energy metabolism. FGF-23 might also be linked to energy metabolism via its regulation of  $\alpha$ -Klotho. Indeed, excess



FGF-23 is associated with decreased  $\alpha$ -Klotho expression in the kidney, either directly or indirectly through alterations of 1,25-dihydroxyvitamin D levels, which can also regulate  $\alpha$ -Klotho gene transcription in distal tubular cells.<sup>14,99</sup> In addition to its membrane functions,  $\alpha$ -Klotho is also a circulating factor released by ecto-domain shedding and/or by transcription of an alternatively spliced isoform.<sup>100</sup> Circulating  $\alpha$ -Klotho has multiple functions, including the regulation of the activity of ion channels and growth factor receptors, such as insulin receptors and insulin-like growth factor-1 receptors.<sup>101</sup> A complex network could be envisioned whereby insulin regulates FGF-23 through activation of insulin receptors in osteoblasts and FGF-23, in turn, regulates circulating levels of  $\alpha$ -Klotho, which modulate insulin receptor function in multiple tissues. Other hormonal pathways might also be involved in FGF-23 regulation, including communication between adipose tissue and bone. For instance, leptin secreted by adipocytes has been shown to directly stimulate FGF-23 synthesis in bone cells.<sup>68</sup>

### Phosphate regulation of FGF-23 conundrum

Given that FGF-23 regulates serum phosphate levels, it is logical that serum phosphate would feed back to regulate FGF-23 secretion. Phosphate regulation of FGF-23 is controversial. Serum phosphate levels positively correlate with elevations in FGF-23 levels in end-stage renal disease,<sup>102</sup> but phosphate restriction failed to lower elevated FGF-23 levels in patients with CKD.<sup>103</sup> A study in patients with CKD and elevated FGF-23 levels who were randomly allocated to either sevelamer (a binder known to provide an acidic load that may alter bone phosphate flux) or calcium carbonate found that urinary phosphorus excretion and PTH decreased in both treatment groups, but FGF-23 levels only decreased in the sevelamer arm.<sup>104</sup>

A small reduction in FGF-23 levels (84 pg/ml versus 61 pg/ml) has been reported in patients with CKD fed a low-phosphate, vegetarian diet.<sup>78</sup> In an adenine-induced CKD rat model, sevelamer prevented the increase in PTH and FGF-23 levels, but the effects were delayed, requiring 2 weeks to achieve reductions in FGF-23 levels, whereas correction in hyperphosphatemia occurred rapidly.<sup>105</sup> A high-phosphate diet was shown to enhance, and a low-phosphate diet to inhibit, the elevation of serum FGF-23 levels in five of six nephrectomized rats, but this result was obtained after 4 weeks of dietary treatment.<sup>106</sup> At present, direct evidence that phosphate regulates *Fgf23* gene transcription is lacking.<sup>73</sup> Phosphate effects on FGF-23 might be indirectly mediated by bone mineralization, which could account for the delayed effects of phosphate on FGF-23 expression, and involve pathways similar to those affected by *PHEX*, *DMPI*, *ENPP1* mutations (*vide supra*).

### Hypothesis for FGF-23 regulation

Current knowledge about FGFR, PTH and 1,25-dihydroxyvitamin D signaling, as well as the function of genes in which mutations result in increased FGF-23 expression, can be used to formulate a speculative model of the regulation of FGF-23 gene expression (Figure 1). This model organizes existing data into a testable model for regulation of *FGF23* gene transcription by local bone-derived and systemic factors under various pathological and physiological conditions.

This hypothetical model predicts that activation of FGFR-1, the predominant FGFR in osteocytes, is the common signaling pathway linking DMP-1, PHEX and mineralization to FGF-23 gene expression in osteocytes and osteoblasts. The role of FGFR-1 in the regulation of FGF-23 has been confirmed for *Phex* and *Dmp1* mutations in mouse and cell culture modes.<sup>61</sup> At this point, whether E-NPP1 and ANK-1 regulation of FGF-23 is through an FGFR-1-dependent pathway or other mechanisms is not known. The schema also shows that FGFR-1 activation can occur through both non-canonical and canonical pathways. The mechanisms linking PHEX and DMP-1 to FGFR-1 activation are not known. DMP-1 and PHEX are expressed in osteocytes, and DMP-1 potentially binds to PHEX via its ASARM motif and to integrins via its RGD sequence located in the 57 kDa C-terminal fragment, thereby providing a possible mechanism for FGFR-1 activation.<sup>107–110</sup> Alternatively, a block in mineralization *per se*, resulting from *PHEX*, *DMP1* or other mutations (such as *ENPP1* or *ANK1*) might lead to canonical activation of FGFR-1 via increased FGF ligand expression and/or alterations of FGF bioactivity.<sup>110</sup> The model also shows an intracrine effect of HMW-FGF-2, which increases FGF-23 expression in Hyp mice.<sup>63</sup> In addition, FGFRs are shown coupled to additional signaling pathways (Figure 1), including those involving PI3K–AKT, Ras–Raf–MAPK, and PLC $\gamma$ , which have not been investigated in regulation of *FGF23* gene expression.

The model also shows the effect of 1,25-dihydroxyvitamin D to stimulate *FGF23* gene transcription by vitamin-D-receptor-mediated mechanisms (Figure 1). In addition, PTH could directly or indirectly regulate *FGF23* gene transcription through possible involvement of sclerostin and the GSK-3 $\beta$ – $\beta$ -catenin pathway. Also included is the potential crosstalk between FGFR-1 and PTH signaling pathways via the common role of CREB in mediating transcriptional regulation. Sclerostin, an osteocyte-derived inhibitor of Wnt signaling, is decreased in bone of Hyp mutant mice, and osteoblasts derived from Hyp mutant mice have increased expression of  $\beta$ -catenin.<sup>57</sup> Moreover, regulation of *FGF23* gene expression by PI3K, Akt, Foxo1 and Sgk3 is supported by data from global Sgk3-deficient mice<sup>98</sup> and Foxo1-deficient<sup>111</sup> mice, which have phenotypes that are consistent with alterations in FGF-23 expression, but in which alterations in FGF-23 expression remain to be confirmed.

Not shown in this model is a possible post-transcriptional mechanism to regulate FGF-23 independent of *FGF23* gene transcription. As noted above, *FGF23* has conserved sites for proprotein convertase cleavage into inactive fragments. The relevance of FGF-23 in metabolism is demonstrated by the fact that mutations lead to familial tumoral calcinosis, which has the opposite phenotype of disorders of FGF-23 excess, and is characterized by hyperphosphatemia, elevated 1,25-dihydroxyvitamin D levels and soft-tissue calcifications. Familial tumoral calcinosis is caused by inactivating mutations of *FGF23*<sup>28</sup> or *GALNT3*,<sup>112</sup> which protects FGF-23 from proteolytic processing by O-glycosylating the protein. Familial tumoral calcinosis can also be caused by inactivating mutations of  $\alpha$ -Klotho, which result in loss of end-organ actions of FGF-23.<sup>113</sup> The physiological role of FGF-23 processing is not clear, but under several circumstances discordance between *FGF23* transcript levels and circulating FGF-23 concentrations exists that suggests an important role of post-translational regulation of FGF-23.

## Clinical implications

### FGF-23 in chronic kidney disease

The pathogenesis of CKD is traditionally viewed from the perspective of the PTH–vitamin D axis, and current treatments focus on suppressing PTH levels with active vitamin D analogues,<sup>114</sup> which can raise serum calcium and phosphate concentrations<sup>115</sup> and further stimulate FGF-23 production.<sup>116–119</sup> Cross-sectional studies in patients with CKD show early FGF-23 elevations in relation to reductions in glomerular filtration rates,<sup>120</sup> which are associated with reductions of levels of 1,25-dihydroxyvitamin D and increments in CYP24.<sup>121</sup> FGF-23 is markedly elevated in end-stage renal disease,<sup>102,120,122</sup> in which levels correlate with the degree of hyperphosphatemia,<sup>102,122</sup> predict refractory hyperparathyroidism in some studies<sup>92</sup> and are associated with increased mortality.<sup>123</sup>

Compelling data exists to support the scenario that FGF-23 is the initial event in CKD that leads to reductions in 1,25-dihydroxyvitamin D levels and secondary increments in PTH. Analysis of the expression of enzymes that regulate vitamin D metabolism, CYP27B1 and CYP24, suggest a pattern consistent with an effect of FGF-23 in CKD.<sup>96</sup> In a rat model of anti-glomerular basement membrane (antiGBM) nephritis, treatment with a neutralizing anti-FGF-23 antibody<sup>121</sup> increased serum levels of 1,25-dihydroxyvitamin D and Cyp27b1, reduced Cyp24 levels and suppressed PTH levels. Treatment with paricalcitol further elevates FGF-23 and suppresses PTH levels in end-stage renal disease.<sup>124</sup>

In this conceptual framework that integrates knowledge of FGF-23 with the PTH and vitamin D endocrine networks, early CKD does not represent a reduction in 1,25-dihydroxyvitamin D levels due to impaired conversion of 25-hydroxyvitamin D to 1,25-dihydroxyvitamin D caused by kidney disease *per se*; rather, FGF-23-mediated suppression of circulating 1,25-dihydroxyvitamin D levels is an active adaptive response, which protects against hyperphosphatemia through FGF-23-mediated reductions of CYP27B1 and increments in CYP24. Changes in the levels of these two enzymes lead to reduction in the levels of both 25-hydroxyvitamin D and 1,25-dihydroxyvitamin D and subsequent effects on gastrointestinal phosphate absorption. Moreover, these changes also increase levels of PTH, which acts in concert with FGF-23 to stimulate phosphaturia. On the other hand, in the late stages of CKD and end-stage renal disease, the dominant stimulus for FGF-23 elevations may be bone remodeling, due to elevated PTH or other factors stimulating FGF-23 expression in the setting of CKD. This hypothesis has been reviewed in great detail elsewhere.<sup>125</sup>

If this hypothesis is correct, FGF-23 might be an early biomarker for earlier interventions in CKD. In addition, treatment approaches to prevent elevated FGF-23 levels could become the initial therapeutic focus but may differ depending on the stage of CKD. As calcitriol increases FGF-23 levels, calcitriol-sparing therapies may be warranted, such as combined low-dose paricalcitol and calcimimetics, which can lower FGF-23 levels in patients with end-stage renal disease.<sup>114,126</sup> Further studies that elucidate the interrelationships between FGF-23, PTH, vitamin D and bone metabolism are of fundamental importance in understanding the pathogenesis and treatment of CKD-mineral bone disorder.

### Implications for low vitamin D levels

CKD is more prevalent in elderly adults than is detected by serum creatinine levels. FGF-23 elevations and increased CYP24 catabolism may contribute to the low levels of 25-hydroxyvitamin D in elderly patients. If so, future translational studies might assess FGF-23 and/or 24,25-dihydroxyvitamin D levels to interpret low 25-hydroxyvitamin D levels (for example, true deficiency or increased catabolism secondary to increased FGF-23 levels in subclinical CKD). This research would add a new perspective to the existing uncertainties regarding the definition of vitamin D insufficiency and effects of vitamin D supplementation.<sup>127</sup>

### FGF-23 and mortality and morbidity

Elevated circulating FGF-23 concentrations are associated with increased mortality in end-stage renal disease<sup>119</sup> and coronary artery disease,<sup>128</sup> progression of renal disease<sup>129</sup> and left ventricular hypertrophy, and increased fat mass and dyslipidemia in elderly patients.<sup>130</sup> Additional knowledge of direct and indirect effects of FGF-23 are needed to understand if these associations are causal or an epiphenomena related to co-variance of circulating FGF-23 levels with the factors causing these associations.

### Conclusions

A key component of the capability of bone to participate in phosphate homeostasis resides in its endocrine function to produce the hormone FGF-23,<sup>4,27</sup> a circulating factor produced predominately by osteoblasts and osteocytes that principally targets the kidney to regulate renal handling of phosphate and vitamin D metabolism. The current lack of understanding of the molecular mechanisms regulating FGF-23 release from osteoblasts and osteocytes is a critical barrier to fully understanding the physiological functions of FGF-23 and the pathogenesis of hereditary and acquired hypophosphatemic disorders. By the study of distinct, single gene mutations that lead to a common increase in FGF-23 production, several pathways for regulating *FGF23* gene transcription have been discovered that link mineralization of bone, vitamin D receptor function and PTH-dependent signaling pathways in the control of circulating FGF-23 levels. The integrative physiology of FGF-23 is being uncovered, and this hormone appears to be involved in several endocrine loops.

Understanding the physiological regulation and function of FGF-23 is also providing new schemas for elucidating the pathogenesis of various disorders of bone and mineral metabolism. In addition to providing a molecular explanation for hereditary hypophosphatemic and hyperphosphatemic disorders, FGF-23 regulation and function is challenging our view of how to interpret circulating 25-hydroxyvitamin D levels and the pathogenesis of secondary hyperparathyroidism. Moreover, the yet-to-be determined functions of FGF-23 in the regulation of  $\alpha$ -Klotho and other endocrine factors, as well as potential effects of FGF-23 on FGFR- $\alpha$ -Klotho complexes in tissues other than bone and kidneys may provide an explanation for the strong associations between elevated serum FGF-23 levels and increased mortality observed in patients with renal failure and in the general population.

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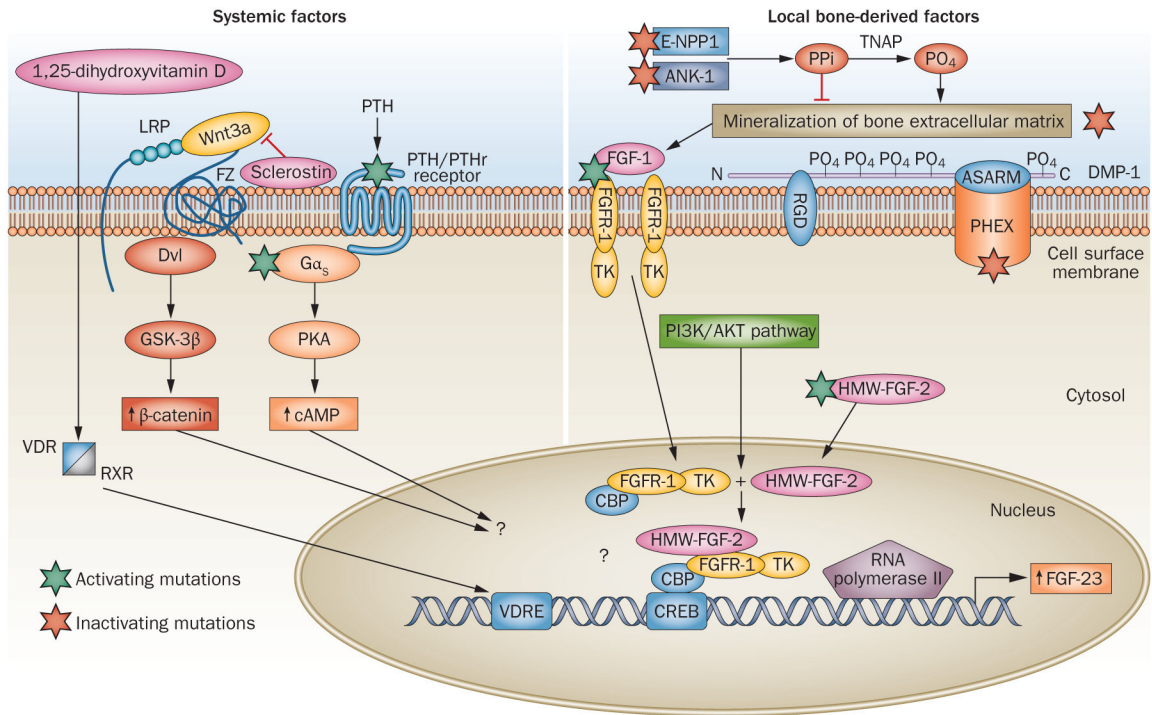
### Key points

- FGF-23 is a hormone produced by osteoblasts and osteocytes in bone that causes phosphaturia and inhibits 1,25-dihydroxyvitamin D production through its binding to FGFR- $\alpha$ -Klotho complexes in kidney tubules
- Primary elevations of circulating FGF-23 concentrations cause hereditary hypophosphatemic disorders and acquired hypophosphatemic disorders, and reductions in circulating FGF-23 concentrations cause familial tumoral calcinosis
- FGF-23 is regulated by local bone-derived factors through activation of FGFR-1 pathways and by systemic factors, including 1,25-dihydroxyvitamin D and parathyroid hormone (in a context-dependent fashion) in osteoblasts
- FGF-23 participates in several physiologically relevant endocrine axes involving 1,25-dihydroxyvitamin D or parathyroid hormone, and possibly additional hormonal networks involving secreted  $\alpha$ -Klotho and other kidney-derived factors
- Elevations in serum FGF-23 is an early adaptive response in chronic kidney disease, leading to maintenance of phosphate balance and development of secondary hyperparathyroidism through reductions in 1,25-dihydroxyvitamin D levels
- Elevations of circulating FGF-23 are an unexpectedly strong predictor of mortality in patients with renal failure, as a result of mechanisms that remain to be elucidated

### Review criteria

The articles upon which this Review was based were selected from a PubMed search using various combinations of keywords, including “hypophosphatemia”, “FGF-23”, “PTH”, “vitamin D”, “Cyp24”, “Klotho”, “bone”, “osteocalcin”, “hereditary hypophosphatemic disorders” for years ranging from 1953 to 2011. In addition, the Online Mendelian Inheritance in Man (OMIM) database was searched using the keyword “hypophosphatemia”. All papers included were full-text papers that were written in English, and data from primary sources in the reference lists of review papers were also confirmed.





**Figure 1.** A speculative model of *FGF23* gene transcriptional regulation. Four activating mutations or pathways (involving FGFR-1,  $G\alpha_s$  encoded by *GNAS*, PTH/PTHr receptor and HMW-FGF-2) and four inactivating mutations (involving PHEX, DMP-1, E-NPP1 and ANK-1) are associated with increased FGF-23 expression in bone. Local bone-derived factors that are linked to mineralization are shown on the right-hand side. ANK-1 and E-NPP1 regulate the transport and biosynthesis of pyrophosphate, and TNAP regulates the conversion of pyrophosphate to phosphate in the extracellular matrix mineralization process, whereas both PHEX and DMP-1 regulate bone mineralization through mechanisms that remain to be fully elucidated. Evidence exists in osteoblasts derived from the Hyp mouse model that defective mineralization is linked to the activation of FGFR-1 as well as HMW-FGF-2 integrative nuclear signaling pathways. The left-hand side of the figure shows systemic factors involved in FGF-23 regulation. 1,25-dihydroxyvitamin D is an important regulator of FGF-23 expression, acting through the VDR and VDRE. PTH can also stimulate FGF-23 through a sclerostin-dependent mechanism involving the Wnt- $\beta$ -catenin pathway, or through stimulation of *GNAS* and cAMP-dependent signaling pathways, as well as indirectly through stimulation of 1,25-dihydroxyvitamin D. Intrinsic and systemic factors are integrated at the levels of *cis*-acting elements in the proximal *FGF23* promoter that remain to be elucidated. A question mark (?) indicates areas of uncertainty. Abbreviations: Hyp, mouse model of X-linked hypophosphatemic rickets;  $PO_4$ , phosphate; PPI, pyrophosphate; TK, tyrosine kinase.

Genetic abnormalities that increase FGF-23 expression in bone

**Table 1**

Genetic abnormalities	Type of mutation									
	Inactivating					Activating				
Hypophosphatemic disorder	ARHR1	ARHR2	None	XLH	OGD	None	MAS	JMC		
OMIM#	241520	613312	None	307800	166250	None	174800	168468		
Mutated gene	<i>DMP1</i>	<i>ENPP1</i>	<i>ANK1</i>	<i>PHEX</i>	<i>FGFR1</i>	<i>FGF2-HMW</i>	<i>GNAS</i>	<i>PTH/PTHr receptor</i>		
Mouse model	<i>Dmp1</i> <sup>-/-</sup>	<i>Enpp1</i> <sup>-/-</sup>	<i>Ank1</i> <sup>K1/K1</sup>	Hyp	None	<i>Tg-FGF2-HMW</i>	<i>Tg-PTH/PTHr receptor</i>	<i>Tg-PTH/PTHr receptor</i>		

Abbreviations: ARHR, autosomal recessive hypophosphatemic rickets; FGF2-HMW, high-molecular-weight FGF2; Hyp, mouse model of XLH; JMC, Jansen metaphyseal chondrodysplasia; MAS, McCune-Albright syndrome; OGD, osteoglophonic dysplasia; Tg, transgene; XLH, X-linked hypophosphatemic rickets.