

Overview of the FGF23-Klotho axis

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Abstract Recent studies have identified a novel bone–kidney endocrine axis that maintains phosphate homeostasis. When phosphate is in excess, fibroblast growth factor-23 (FGF23) is secreted from bone and acts on the kidney to promote phosphate excretion into urine and suppress vitamin D synthesis, thereby inducing negative phosphate balance. One critical feature of FGF23 is that it requires Klotho, a single-pass transmembrane protein expressed in renal tubules, as an obligate coreceptor to bind and activate FGF receptors. Several hereditary disorders that exhibit inappropriately high serum FGF23 levels are associated with phosphate wasting and impaired bone mineralization. In contrast, defects in either FGF23 or Klotho are associated with phosphate retention and a premature-aging syndrome. The aging-like phenotypes in Klotho-deficient or FGF23-deficient mice can be rescued by resolving hyperphosphatemia with dietary or genetic manipulation, suggesting a novel concept that phosphate retention accelerates aging. Phosphate retention is universally observed in patients with chronic kidney disease (CKD) and identified as a potent risk of death in epidemiological studies. Thus, the bone–kidney endocrine axis mediated by FGF23 and Klotho has emerged as a novel target of therapeutic interventions in CKD.

Keywords Klotho · FGF23 · Phosphate · Vitamin D · CKD

Introduction

The blood phosphate level is determined by counterbalance between absorption of dietary phosphate from the intestine, mobilization from bone (the major reservoir of calcium and phosphate in the body), and excretion from the kidney into urine [1]. These processes are coordinately regulated by several endocrine factors. Vitamin D and parathyroid hormone (PTH), which have been extensively studied as hormones that regulate calcium metabolism [2], are also involved in phosphate metabolism. The active form of vitamin D (1,25-dihydroxyvitamin D₃) is synthesized in the kidney and acts on the intestine to increase absorption of dietary calcium and phosphate. It also acts on bone to stimulate osteoclastogenesis and promote mobilization of calcium and phosphate from the reservoir, thereby increasing blood levels of both calcium and phosphate. PTH acts on the kidney to promote both vitamin D synthesis and phosphaturia (phosphate excretion into urine). As a result, unlike vitamin D, PTH can selectively increase blood calcium levels without concomitant increase in blood phosphate levels [3].

Recent studies have identified fibroblast growth factor-23 (FGF23) as a novel hormone that lowers blood phosphate levels [4–6]. When phosphate is in excess, FGF23 is secreted from bone and acts on the kidney to induce phosphaturia and suppress vitamin D synthesis, thereby inducing a negative phosphate balance to maintain phosphate homeostasis [6–9]. FGF23 requires Klotho protein as a coreceptor for high affinity binding to cognate FGF receptors (FGFRs). The purpose of this review was to overview recent progress in our understanding of endocrine regulation of phosphate metabolism by FGF23 and Klotho and to discuss its potential role in the pathophysiology of chronic kidney disease (CKD).

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FGF23

FGF23 was originally identified as a gene mutated in patients with autosomal dominant hypophosphatemic rickets (ADHR) [5]. Patients with ADHR carry missense mutations in the *FGF23* gene that confer resistance to proteolytic degradation of the FGF23 protein [10]. As a result, ADHR patients exhibit high serum FGF23 levels. Because FGF23 has an activity that induces negative phosphate balance, ADHR patients exhibit phosphate-wasting phenotypes such as hypophosphatemia and rickets. Although the protease(s) that inactivates FGF23 remains to be identified, it cleaves FGF23 at the $^{176}\text{RXXR}^{179}$ motif and generates two inactive fragments [11]. Missense mutations in this critical motif (R176Q, R179Q/W) make the protein resistant to the protease and increase the half-life of FGF23 in the blood [12]. Several other phosphate-wasting syndromes associated with FGF23 excess have been identified, including tumor-induced osteomalacia (TIO), X-linked hypophosphatemia (XLH), and autosomal recessive hypophosphatemic rickets (ARHR). TIO is caused by FGF23-producing tumors [13]. XLH and ARHR are caused by mutations in the *PHEX* (a phosphate-regulating gene with homologies to endopeptidases on the X-chromosome) gene [14] and the *DMP-1* (dentin matrix protein-1) gene [15, 16], respectively. These two genes are expressed in osteocytes in the bone where FGF23 is primarily produced and secreted. Recent animal studies have demonstrated that *PHEX* and *DMP-1* may be involved in the regulation of FGF23 gene expression. *Hyp* mice, which have deletions in the *Phex* gene, show high FGF23 expression in the bone, high serum levels of FGF23, hypophosphatemia, and impaired bone mineralization, as observed in XLH patients [17, 18]. *DMP-1* knockout mice also exhibit high FGF23 expression and phosphate-wasting phenotypes, as observed in ARHR patients [15]. The precise mechanism by which *PHEX* and *DMP-1* proteins suppress expression of the *FGF23* gene remains to be determined.

In contrast to patients with ADHR, ARHR, XLH, and TIO that exhibit phosphate-wasting phenotypes, patients with familial tumoral calcinosis (FTC) exhibit phosphate-retention phenotypes including hyperphosphatemia and ectopic calcification associated with low serum FGF23. FTC is caused by mutations in the *GALNT3* gene that encodes a glycosyl transferase called UDP-N-acetyl- α -D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-3 (ppGaNTase-3) [19]. This enzyme is required for O-glycosylation of FGF23 at Thr¹⁷⁸, which resides within the cleavage motif of FGF23 [20]. It is likely that loss of O-glycosylation at Thr¹⁷⁸ increases susceptibility of FGF23 to proteolytic degeneration, resulting in low serum levels of FGF23. Mice that completely lack FGF23 (*Fgf23*^{-/-} mice) develop severe

phosphate-retention phenotypes characterized by extensive soft tissue calcification and hyperphosphatemia [21]. These studies on phosphate-wasting and phosphate-retention syndromes in mice and humans have provided unequivocal evidence indicating that FGF23 is an essential hormone for maintaining phosphate homeostasis.

Although FGF23 belongs to the FGF ligand superfamily [22], phylogenetic and sequence analyses have segregated FGF23 and two additional FGFs (FGF19 and FGF21) from the other FGF family members [23]. These three atypical FGFs—namely, FGF19, FGF21, and FGF23—are collectively called endocrine FGFs because they function as endocrine factors, unlike the other classic FGFs that basically function as paracrine and/or autocrine factors [24]. The molecular basis behind the endocrine mode of action may lie in the fact that these endocrine FGFs have low affinity to heparan sulfate (HS). In general, FGF ligands have a conserved core region with 12 antiparallel β strands, where the HS-binding domain resides [25–27]. However, the HS-binding domain of endocrine FGFs deviates from that of the other paracrine-acting FGFs and prohibits formation of hydrogen bonding between HS and amino acid residues in the HS-binding domain, which is the basis of affinity to HS [28, 29]. This unique structural feature reduces affinity of endocrine FGFs to HS and allows them to escape from HS-rich extracellular matrices and enter into systemic circulation. Although the low affinity of endocrine FGFs to HS may be advantageous for the endocrine mode of action, it may be disadvantageous for signal transduction through FGFRs because HS is required for high-affinity binding of FGFs to FGFRs. FGFRs are single-pass transmembrane receptor tyrosin kinases that dimerize upon binding FGFs. HS participates in the FGF-FGFR interaction and promotes formation of a 2:2:2 FGF-FGFR-HS signaling complex, which is essential for efficient activation of FGFR tyrosine kinase [30]. Thus, endocrine FGFs may require a cofactor(s) other than and/or in addition to HS to secure efficient dimerization and activation of FGFR. In fact, FGF23 cannot activate FGF signaling in most cultured cells, even when they express FGFRs endogenously, whereas classic FGFs such as FGF2 can activate FGF signaling in these cells [31]. Identity of the putative cofactor(s) required for FGF23 to activate FGFRs was not clear until it was realized that the phenotypes of FGF23-deficient mice are identical with those of *Klotho*-deficient mice.

Klotho

The *klotho* gene, named after a Greek goddess who spins the thread of life, was originally identified as a gene mutated in a mouse strain that inherits a premature-aging

syndrome in an autosomal recessive manner [32]. Mice defective in *klotho* gene expression develop multiple aging-like phenotypes around 3–4 weeks after birth, including growth retardation, hypogonadotropic hypogonadism, rapid thymus atrophy [33], skin atrophy, sarcopenia, vascular calcification, osteopenia [34], pulmonary emphysema [35–37], cognition impairment [38], hearing disturbance [39], and motor neuron degeneration [40], and die around 2 months of age. In contrast, transgenic mice that overexpress *Klotho* live longer than wild-type mice [41]. Thus, the *Klotho* gene may be an aging suppressor gene that extends life span when overexpressed and accelerates aging when disrupted [42]. Furthermore, polymorphisms in the human *KLOTHO* gene are associated with life span [43] as well as bone mineral density [44–46], high-density lipoprotein (HDL) cholesterol level, blood pressure, stroke [47], coronary artery disease [48], and cognitive function [49], suggesting that *Klotho* may be involved in the regulation of aging processes in humans. The *Klotho* gene encodes a single-pass transmembrane protein that belongs to a family 1 glycosidase [50] and is expressed primarily in renal tubules in the kidney and choroid plexus in the brain [32]. Although recombinant *Klotho* protein was reported to have weak β -glucuronidase activity in vitro [51], physiological relevance of the β -glucuronidase activity in vivo was not clear.

The clue to understanding *Klotho* protein function was the fact that FGF23-deficient mice and *Klotho*-deficient mice develop identical phenotypes. FGF23-deficient mice not only exhibit phosphate retention but also develop multiple aging-like phenotypes [21], which is reminiscent of *Klotho*-deficient mice. Conversely, *Klotho*-deficient mice not only develop a premature-aging syndrome but also exhibit hyperphosphatemia [52, 53], which is reminiscent of FGF23-deficient mice. These observations suggested that *Klotho* and FGF23 might function in a common signal transduction pathway. In fact, *Klotho* protein forms a constitutive binary complex with several FGF receptor isoforms (FGFR1c, 3c, 4) and significantly increases the affinity of these FGFRs specifically to FGF23 [31]. Thus, *Klotho* protein functions as an obligate coreceptor for FGF23. This finding was later confirmed in an independent study [54]. The fact that FGF23 requires *Klotho* protein as a coreceptor explains why *Klotho*-deficient mice, FGF23-deficient mice, and mice lacking both *Klotho* and FGF23 [55] develop identical phenotypes. It also explains why extremely high serum FGF23 levels of *Klotho*-deficient mice [54] do not cause any adverse effects in *Klotho*-deficient mice [55]. In addition, kidney-specific expression of *Klotho* explains why FGF23 can identify the kidney as its target organ among many other tissues that express multiple FGFR isoforms. *Klotho* protein function is to compensate for the low affinity of FGF23 to heparan sulfate

and specifically support FGFR activation with FGF23, which represents a novel mechanism for confining target organs in redundant ligand-receptor interactions.

Endocrine regulation of phosphate metabolism

The bone–kidney endocrine axis mediated by FGF23 and *Klotho* has emerged as the major regulator of phosphate homeostasis [4, 9, 24, 56–58]. FGF23 has an activity that reduces the number of sodium-phosphate cotransporter type-2a (NaPi-2a) on the brush border membrane of proximal tubules, thereby promoting renal phosphate excretion [59–62]. Thus, FGF23 functions as a phosphaturic hormone. In addition, FGF23 suppresses synthesis and promotes degradation of 1,25-dihydroxyvitamin D₃ in proximal tubules [63]. FGF23 down-regulates expression of the *Cyp27b1* gene, which encodes 1 α -hydroxylase, the enzyme that synthesizes the active form of vitamin D (1,25-dihydroxyvitamin D₃) from its inactive precursor (25-hydroxyvitamin D₃). Furthermore, FGF23 up-regulates expression of the *Cyp24* gene that encodes 24-hydroxylase, the enzyme that hydrolyzes and inactivates 1,25-dihydroxyvitamin D₃. Thus, FGF23 functions as a counterregulatory hormone for vitamin D [8]. The ability of FGF23 to reduce serum 1,25-dihydroxyvitamin D₃ levels also contributes to induction of negative phosphate balance through reducing phosphate absorption from the intestine. Importantly, 1,25-dihydroxyvitamin D₃ up-regulates expression of the FGF23 gene [63] and closes a negative feedback loop (Fig. 1). Disruption of this negative feedback loop results in high serum 1,25-dihydroxyvitamin D₃ levels, as observed in *Klotho*-deficient mice, FGF23-deficient mice, and FTC patients. Recently, a patient carrying a homozygous missense mutation in the *KLOTHO* gene (H193R) was reported [64]. The patient exhibited phosphate-retention phenotypes similar to FTC patients, indicating that the H193R mutation is a loss-of-function mutation. This is the first case in humans exhibiting phosphate retention due to a defect in *Klotho* protein.

It should be noted that *Klotho* protein is expressed much more abundantly in distal convoluted tubules than in proximal tubules, whereas both phosphate reabsorption and vitamin D synthesis take place in proximal tubules. This discrepancy has raised two possibilities that are not mutually exclusive. One possibility is that, although *Klotho* expression levels in proximal tubules are not as high as distal convoluted tubules, FGF23 may signal through the FGFR-*Klotho* complex on proximal tubules and directly regulate NaPi-2a expression and vitamin D synthesis. In this case, the function of *Klotho* protein abundantly expressed in distal convoluted tubules must be addressed. The other possibility is that FGF23 may act first on distal

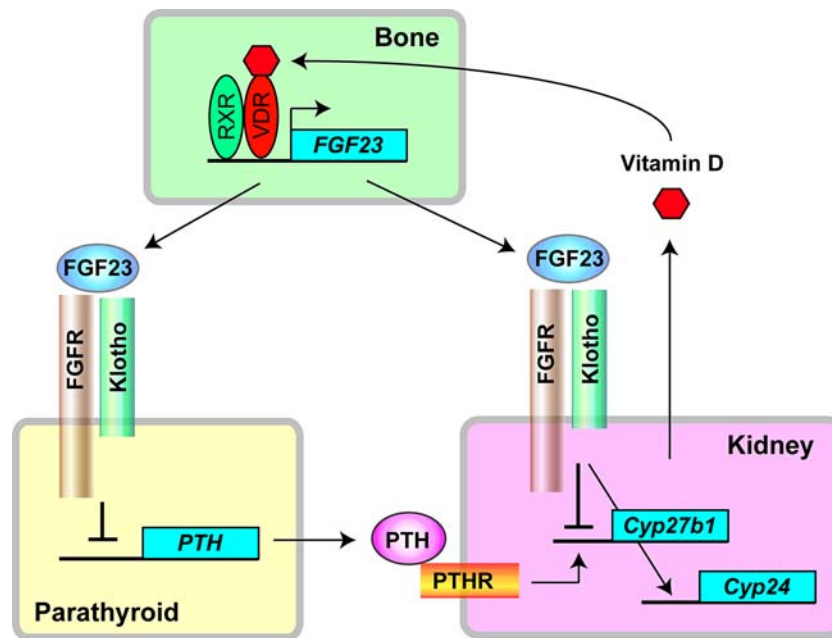


Fig. 1 The bone–kidney–parathyroid endocrine axes mediated by fibroblast growth factor-23 (FGF23) and Klotho. Active form of vitamin D (1,25-dihydroxyvitamin D₃) binds to vitamin D receptor (VDR) in the bone (osteocytes). The ligand-bound VDR forms a heterodimer with a nuclear receptor RXR and transactivates expression of the FGF23 gene. FGF23 secreted from bone acts on the Klotho-FGF receptor (FGFR) complex expressed in the kidney (the bone–kidney axis) and parathyroid gland (the bone–parathyroid axis). In the kidney, FGF23 suppresses synthesis of active vitamin D by down-regulating expression of the *Cyp27b1* gene and promotes its

inactivation by up-regulating expression of the *Cyp24* gene, thereby closing a negative feedback loop for vitamin D homeostasis. In the parathyroid gland, FGF23 suppresses production and secretion of parathyroid hormone (PTH). PTH binds to the PTH receptor (PTHR) expressed on renal tubular cells, leading to up-regulation of the *Cyp27b1* gene expression. Thus, suppression of PTH by FGF23 reduces expression of the *Cyp27b1* gene and serum levels of 1,25-dihydroxyvitamin D₃. This closes another long negative feedback loop for vitamin D homeostasis

convoluted tubules and then generate a secondary signal that instructs proximal tubules to reduce phosphate reabsorption and vitamin D synthesis. Recent animal studies may support the latter possibility. Despite the fact that proximal tubules primarily express FGFR3, knockout of the *Fgfr3* gene in *Hyp* mice, which have elevated serum FGF23 levels, failed to rescue their phosphate-wasting phenotypes [65]. Furthermore, it was reported that activation of the FGF signaling pathway was detectable only in distal convoluted tubules after injection of FGF23 into mice [66]. These findings suggest that activity of FGF23 may be independent of FGF signaling activation in the proximal tubule.

PTH plays an important role not only in calcium metabolism but also in phosphate homeostasis. Like FGF23, PTH has an activity that induces phosphaturia [3]. However, in contrast to FGF23, PTH up-regulates expression of the *Cyp27b1* gene and increases serum 1,25-dihydroxyvitamin D₃ levels [2]. Recent studies showed that the parathyroid gland is one of the few organs that express a decent amount of Klotho protein endogenously, indicating that the parathyroid may be a target organ of FGF23. In fact, FGF23 down-regulates PTH expression and suppresses PTH secretion in vivo and in vitro [67, 68]. The ability of FGF23 to reduce serum PTH levels may further

enhance the activity of FGF23 as a counterregulatory hormone for vitamin D and contribute to a long negative feedback loop involving bone, kidney, and parathyroid gland (Fig. 1). However, it should be noted that patients with CKD typically exhibit secondary hyperparathyroidism associated with high serum FGF23 levels, which seemingly contradicts the ability of FGF23 to suppress PTH secretion and production. Considering that Klotho expression is positively regulated by vitamin D [53], one possible explanation is that low serum vitamin D levels in CKD patients may reduce Klotho expression not only in the kidney (discussed below) but also in the parathyroid glands and make these organs resistant to FGF23.

Phosphate toxicity

Defects in either Klotho or FGF23 disrupt the negative feedback loops that maintain phosphate and vitamin D homeostasis, resulting in high serum phosphate and vitamin D levels. High serum vitamin D promotes intestinal absorption of calcium and induces hypercalcemia as well. Importantly, this metabolic state characterized by high serum phosphate, calcium, and vitamin D levels is

associated with a premature aging syndrome, as observed in Klotho-deficient mice and FGF23-deficient mice. These observations imply that phosphate, calcium, and/or vitamin D may be toxic when retained and thus accelerate aging. Several animal studies have supported this notion. First, vitamin-D-deficient diet not only restored serum phosphate and calcium levels but also rescued several aging-like phenotypes in Klotho-deficient mice and FGF23-deficient mice [53, 69]. Second, ablation of vitamin D activity in Klotho-deficient mice and FGF23-deficient mice by disrupting the *Cyp27b1* gene [70, 71] or vitamin D receptor gene [72] also rescued hyperphosphatemia, hypercalcemia, and the premature aging syndrome. Lastly, low phosphate diet rescued shortened life span and vascular calcification in FGF23-deficient mice and Klotho-deficient mice [69, 73]. These studies provide evidence that the premature aging syndrome caused by defects in the bone–kidney endocrine axis is due to retention of phosphate, calcium, and/or vitamin D. It should be noted that low phosphate diet rescued FGF23-deficient mice despite the fact that it further increased already high serum calcium and vitamin D levels [69], suggesting that phosphate, but not calcium or vitamin D, is primarily responsible for the aging-like phenotypes. It is likely that low vitamin D diet and ablation of vitamin D activity rescued accelerated aging through reducing serum phosphate levels, although it remains to be determined whether high serum vitamin D and/or calcium levels are a prerequisite for phosphate to accelerate aging.

Chronic kidney disease

Phosphate retention is universally observed in patients with CKD. Hyperphosphatemia has been identified as a potent, independent risk of death [74]. Why does hyperphosphatemia increase mortality? One likely explanation is that high blood phosphate levels trigger vascular calcification and accelerate life-threatening complications such as cardiovascular events [75]. Vascular calcification is a very common complication in CKD and has been shown to contribute to the high morbidity and mortality in terms of cardiovascular events. The National Kidney Foundation task force indicated that the cardiovascular mortality of a 35-year-old patient on dialysis is equivalent to that of an 80-year-old healthy individual, rendering CKD to be one of the most potent accelerators of aging [76]. In addition, the American Heart Association announced that CKD should be included in the highest-risk group for cardiovascular disease and that patients with CKD should receive aggressive therapeutic measures to reduce morbidity and mortality [77]. Thus, lowering blood phosphate levels is expected to reduce vascular calcification and cardiovascular events, thereby improving prognosis of CKD patients. In fact, CKD patients with hyperphosphatemia

(≥ 6.5 mg/dl) were reported to have higher risk for death resulting from cardiovascular disease than those with the lower serum phosphate levels (< 6.5 mg/dl) [78]. Based on these observations, control of blood phosphate levels < 6.5 mg/dl has been proposed as one of the most important therapeutic goals for improving life expectancy of CKD patients.

It is likely that dysregulation of the FGF23-Klotho endocrine axis may be involved in the mechanism by which CKD patients fail to maintain phosphate homeostasis. In fact, serum FGF23 levels are increased with advancing stages of CKD [79], whereas Klotho expression in the kidney is significantly decreased in CKD patients [80] and in various animal models of chronic and acute renal damage [81, 82]. Thus, CKD may be viewed as a state of FGF23 resistance caused by Klotho deficiency. This viewpoint explains several observations on phosphate metabolism in CKD that lack mechanistic insights. For example, epidemiological studies have indicated that serum FGF23 levels increase long before serum phosphate levels increase during the progression of CKD [79]. In other words, patients with early stages of CKD require higher serum FGF23 levels than normal people to maintain normal serum phosphate levels. This may represent compensation for end-organ resistance to FGF23 due to decreased Klotho expression in the kidney. It has been also known that serum vitamin D levels decrease long before serum phosphate levels increase during CKD progression [79]. This may be a result of the secondary hyper-FGF23-emia caused by decreased renal Klotho expression, because FGF23 has an activity that lowers serum vitamin D levels. In addition, epidemiological studies have indicated that high serum FGF23 levels are associated with poor prognosis in patients undergoing dialysis [83]. This may be explained by assuming that high serum FGF23 indicates low renal Klotho expression associated with severe renal damage. It remains to be determined whether decrease in Klotho expression is one of the earliest changes in the progression of CKD.

Of note, recent animal studies have shown that Klotho functions as a renoprotective factor. Although the mechanism is yet to be determined, overexpression of Klotho ameliorated progressive renal injury in mouse models of glomerulonephritis [81] and acute kidney injury [82]. Thus, it may be of therapeutic value for CKD to preserve Klotho expression in the kidney. Klotho expression is down-regulated by angiotensin II [84, 85] and up-regulated by peroxisome proliferator-activated receptor-gamma (PPAR γ) agonists such as thiazolidinediones [86]. These observations suggest that renoprotective effects of angiotensin-converting enzyme inhibitors and thiazolidinediones may be partly attributed to their potential for increasing or preserving Klotho expression in the kidney. Klotho expression is also up-regulated by 1,25-dihydroxyvitamin D₃ [53].

Thus, low serum vitamin D caused by secondary hyper-FGF23-emia further reduces Klotho expression, potentially leading to deterioration spiral of Klotho expression. The benefit of vitamin D replacement therapy may be partly attributed to interruption of this vicious cycle.

In addition to functioning as an obligate coreceptor for FGF23, Klotho protein functions as a humoral factor that regulates activity of several ion channels and growth-factor receptors [41], which represents a novel function of Klotho protein. The entire extracellular domain of Klotho protein is clipped by a membrane-anchored protease ADAM10/17 on the cell surface and released into the extracellular space [87]. In fact, Klotho ectodomain (secreted Klotho protein) is detectable in the blood, urine, and cerebrospinal fluid [41, 88]. The secreted Klotho protein in turn functions as a putative sialidase that removes terminal sialic acids in the glycans of several ion channels, including a calcium channel, transient receptor potential vanilloid type isoform 5 (TRPV5) [89, 90], and a potassium channel, renal outer medullary potassium channel-1 (ROMK1) [91]. Removal of sialic acids by secreted Klotho protein on the cell surface prevents internalization of these ion channels, resulting in increase in transepithelial calcium (Ca^{2+}) absorption and potassium (K^+) secretion in distal nephrons, respectively. Thus, Klotho protein not only regulates phosphate metabolism by functioning as a coreceptor for FGF23 but also regulates calcium and potassium metabolism by functioning as a humoral factor that modifies trafficking of TRPV5 and ROMK1. Significance of the secreted Klotho protein in the regulation of calcium and potassium homeostasis and in pathophysiology in CKD remains to be determined.

It has become increasingly clear that phosphate metabolism plays a critical role in the pathophysiology in CKD and that hyperphosphatemia should be aggressively treated to improve life expectancy of CKD patients. In this context, the Klotho and FGF23 axis is expected to be a novel target of therapeutic interventions in CKD.

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