



# **Pathogenesis of FGF23-Related Hypophosphatemic Diseases Including X-linked Hypophosphatemia**

Tatsuro Nakanishi<sup>1,2</sup> and Toshimi Michigami<sup>1,\*</sup>

- <sup>1</sup> Department of Bone and Mineral Research, Research Institute, Osaka Women's and Children's Hospital, Izumi 594-1101, Japan; tnrebo@wch.opho.jp
- <sup>2</sup> Department of Pediatrics, Osaka University Graduate School of Medicine, Suita 565-0871, Japan
- Correspondence: michigami@wch.opho.jp; Tel.: +81-725-56-1220

Abstract: Since phosphate is indispensable for skeletal mineralization, chronic hypophosphatemia causes rickets and osteomalacia. Fibroblast growth factor 23 (FGF23), which is mainly produced by osteocytes in bone, functions as the central regulator of phosphate metabolism by increasing the renal excretion of phosphate and suppressing the production of 1,25-dihydroxyvitamin D. The excessive action of FGF23 results in hypophosphatemic diseases, which include a number of genetic disorders such as X-linked hypophosphatemic rickets (XLH) and tumor-induced osteomalacia (TIO). Phosphate-regulating gene homologous to endopeptidase on the X chromosome (PHEX), dentin matrix protein 1 (DMP1), ectonucleotide pyrophosphatase phosphodiesterase-1, and family with sequence similarity 20c, the inactivating variants of which are responsible for FGF23-related hereditary rickets/osteomalacia, are highly expressed in osteocytes, similar to FGF23, suggesting that they are local negative regulators of FGF23. Autosomal dominant hypophosphatemic rickets (ADHR) is caused by cleavage-resistant variants of FGF23, and iron deficiency increases serum levels of FGF23 and the manifestation of symptoms in ADHR. Enhanced FGF receptor (FGFR) signaling in osteocytes is suggested to be involved in the overproduction of FGF23 in XLH and autosomal recessive hypophosphatemic rickets type 1, which are caused by the inactivation of PHEX and DMP1, respectively. TIO is caused by the overproduction of FGF23 by phosphaturic tumors, which are often positive for FGFR. FGF23-related hypophosphatemia may also be associated with McCune-Albright syndrome, linear sebaceous nevus syndrome, and the intravenous administration of iron. This review summarizes current knowledge on the pathogenesis of FGF23-related hypophosphatemic diseases.

Keywords: phosphate; fibroblast growth factor 23; osteocytes; rickets; osteomalacia

# 1. Introduction

Phosphorus is an essential nutrient that mediates the majority of biological processes, including the integrity of cell membranes, the maintenance and inheritance of genetic information, energy metabolism, and the regulation of protein function by phosphorylation [1]. In vertebrates, phosphorus also contributes to skeletal mineralization as a constituent of hydroxyapatite (calcium-phosphate crystals). In the human adult body, approximately 90% of total phosphorus is stored in bone, while the remainder is present in the soft tissues and less than 1% in extracellular fluid [2]. In serum, the majority of phosphorus exists as free ions of inorganic phosphate (Pi), such as HPO<sub>4</sub><sup>2–</sup> and H<sub>2</sub>PO<sub>4</sub><sup>–</sup>, at a ratio of 4:1 at physiological pH [3]. Serum levels of Pi are influenced by age, dietary intake, serum pH, and so on.

Since phosphate is indispensable for the formation of hydroxyapatite, its chronic deficiency or wasting leads to impaired skeletal mineralization, namely, rickets in children and osteomalacia in adults [4]. At the beginning of this century, fibroblast growth factor 23 (FGF23) was identified as the molecule responsible for autosomal dominant hypophosphatemic rickets (ADHR) and tumor-induced hypophosphatemic osteomalacia (TIO) [5,6].



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**Copyright:** © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Since then, our understanding of the mechanisms underlying phosphate metabolism and the pathogenesis of various related diseases has increased. This review discusses current concepts on the role of FGF23 in phosphate homeostasis and the pathogenesis of FGF23-related hypophosphatemic diseases.

#### 2. Phosphorus Homeostasis

In mammals, the phosphate balance in postnatal life is maintained by its intestinal absorption, renal excretion, and accumulation in and release from bone and soft tissue [1]. The insufficient absorption, excess renal excretion, or excessive shift of Pi to bone or soft tissue may cause hypophosphatemia [7]. Since phosphate is indispensable for skeletal mineralization, chronic hypophosphatemia leads to rickets and osteomalacia.

Pi absorption in the intestines is mediated by two processes: a passive, paracellular diffusion process and an active, transcellular transport process. The latter is mediated by the type IIb Na<sup>+</sup>/Pi co-transporter (NaPi-IIb), which is encoded by the *SLC34A2* gene in humans [8]. The expression of NaPi-IIb in the intestines is up-regulated by the low dietary intake of Pi and 1,25-dihydroxyvitamin D (1,25(OH)<sub>2</sub>D), an active metabolite of vitamin D [9,10]. A dietary deficiency of phosphate is rare because the majority of foods contain large amounts of phosphate. However, the insufficient action of vitamin D decreases the absorption of Pi in the intestines.

In the kidneys, the majority of Pi filtered in the glomeruli is mainly reabsorbed by type IIa and IIc Na<sup>+</sup>/Pi co-transporters (NaPi-IIa and NaPi-IIc) localized at the brush border membrane (BBM) of the proximal tubules [11–13]. NaPi-IIa and NaPi-IIc are encoded by the *SLC34A1* and *SLC34A3* genes, respectively, in humans. Inactivating variants in the *SLC34A3* gene cause hereditary hypophosphatemic rickets with hypercalciuria (HHRH) [14]. HHRH is characterized by renal Pi wasting, hypophosphatemia, increased levels of serum 1,25(OH)<sub>2</sub>D, and secondary hypocalciuria. On the other hand, inactivating variants in the *SLC34A1* gene have been shown to cause Fanconi renotubular syndrome 2, infantile hypercalciuria 2, and nephrolithiasis/osteoporosis associated with hypophosphatemia [1].

Endocrine factors including 1,25(OH)<sub>2</sub>D, parathyroid hormone (PTH), and FGF23 play critical roles in phosphate homeostasis. As described above, 1,25(OH)<sub>2</sub>D increases the intestinal absorption of Pi by up-regulating NaPi-IIb [10]. PTH increases the renal excretion of Pi by decreasing the amounts of NaPi-IIa and NaPi-IIc that localize in the BBM [15–17]. FGF23 functions as the central regulator of phosphate metabolism, as described in the following section.

## 3. Roles of FGF23 in Mineral Metabolism

FGF23 is a 32-kDa protein that consists of 251 amino acids including a 24-amino acid signal peptide at its N terminus [6]. It is mainly produced by osteoblast lineage cells, particularly osteocytes, which terminally differentiate from osteoblasts and are embedded within the bone matrix [18]. In adult bone, osteocytes make up more than 90% to 95% of all bone cells. Although the location and inaccessibility of osteocytes in the mineralized bone matrix delayed our understanding of their cellular functions, studies conducted in the last few decades have revealed their roles in bone homeostasis. Osteocytes are essential for controlling the bone mass in postnatal life by sensing mechanical strain and regulating the formation and resorption of bone [19]. They also produce sclerostin, an inhibitor of Wnt/ $\beta$ -catenin signaling. Mechanical strain reduces the production of sclerostin by osteocytes, which in turn enhances Wnt/ $\beta$ -catenin signaling and increases bone formation [20]. In addition, mouse studies suggested that the receptor activator of nuclear factor kappa B ligand produced by osteocytes controls the postnatal bone mass [21]. The expression of FGF23 by osteocytes indicates that they also play critical roles in phosphate homeostasis.

FGF23 belongs to the FGF19 subfamily together with FGF19 and FGF21, based on their unique features among FGFs [22]. Members of the FGF19 family exert their effects on distant target organs and tissues in an endocrine manner, and their low binding affinity

to heparin/heparan sulfate has been suggested to confer endocrine functions due to less binding to heparan sulfate surrounding their producing cells and their entry into the circulation [22]. Members of the FGF19 subfamily require the single-pass transmembrane protein Klotho,  $\alpha$ Klotho for FGF23, and  $\beta$ Klotho for FGF19 and FGF21, for their signal transduction through an FGF receptor (FGFR) [22–24]. A previous study demonstrated that the N-terminal domain of FGF23 bound to FGFR while its C-terminal domain bound to  $\alpha$ Klotho [22].

Since FGF23 requires  $\alpha$ Klotho together with FGFR to exert its effects, organs and tissues expressing both  $\alpha$ Klotho and FGFR, such as the kidneys, parathyroid glands, placenta, and choroid plexus, may be the physiological targets of FGF23 [25,26]. In the kidneys, which are the main target, FGF23 increases urinary Pi excretion by suppressing the expression of NaPi-IIa and NaPi-IIc. In addition, FGF23 decreases the production of 1,25(OH)<sub>2</sub>D by suppressing the renal expression of *Cyp27b1* encoding 25-hydroxyvitamin D 1 $\alpha$ -hydroxylase and increasing that of *Cyp24a1* encoding 25-hydroxyvitamin D-24-hydroxylase, which leads to a reduction in the absorption of Pi in the intestines [6] (Figure 1).



**Figure 1.** Effects of FGF23 on kidneys. FGF23 produced in bone binds to FGFR and  $\alpha$ Klotho in the kidneys and suppresses the renal expression of NaPi-IIa and NaPi-IIc Na<sup>+</sup>/Pi co-transporters, which increases urinary Pi excretion and reduces serum Pi levels. In addition, FGF23 decreases the production of 1,25(OH)<sub>2</sub>D by the down-regulation of 25-hydroxyvitamin D-1 $\alpha$ -hydroxylase (1 $\alpha$ OHase) and up-regulation of 25-hydroxyvitamin D-24-hydroxylase (24OHase), which reduces the intestinal absorption of Pi and further lowers serum Pi levels.

The findings of animal studies have suggested that FGF23 suppresses the production and secretion of PTH in both  $\alpha$ Klotho-dependent and -independent manners [27,28]. Since 1,25(OH)<sub>2</sub>D and PTH have been shown to increase the expression of FGF23 [29–32], these factors counter-regulate with FGF23 to maintain phosphate homeostasis. The findings of mouse studies have indicated that maternal FGF23 exerts its effects on the placenta to increase the expression of *Cyp24a1*; however, it does not alter placental Pi transport [33].

FGF23 is inactivated by cleavage between  $\operatorname{Arg}^{179}$  and  $\operatorname{Ser}^{180}$  via a subtilisin-like proprotein convertase, which recognizes the motif  $\operatorname{R}^{176}XXR^{179}/\operatorname{S}^{180}$  [34]. FGF23 is *O*glycosylated at Thr<sup>178</sup> by the enzyme UDP-*N*-acetyl- $\alpha$ -D-galacosamine:polypeptide *N*acetylgalactosaminyltransferase 3 (GalNAc-T3), which is suggested to prevent cleavage by the subtilisin-like proprotein convertase [35]. Inactivating mutations in the *GALNT3* gene encoding GalNAc-T3 result in the cleavage of intact FGF23 before secretion, and the resultant deficiency of intact FGF23 causes hyperphosphatemic familial tumoral calcinosis (HFTC), a rare autosomal recessive disease characterized by hyperphosphatemia, normal or elevated levels of serum 1,25(OH)<sub>2</sub>D, and ectopic calcification [36,37]. Inactivating mutations in the *FGF23* gene itself and the *KLOTHO* gene encoding  $\alpha$ Klotho were also found to be responsible for HFTC [38–40].

Both Fgf23-knockout mice and Klotho-deficient mice exhibit hyperphosphatemia with increased renal Pi reabsorption and elevated serum  $1,25(OH)_2D$  levels, which resemble the features in patients with HFTC [25,41]. These mutant mice also show severe growth retardation with abnormal skeletal phenotype and short lifespan. Interestingly, Fgf23-knockout mice display a marked accumulation of osteoid despite hyperphosphatemia [41]. Although mechanisms for this counterintuitive observation remain unclear, it may suggest the effects of FGF23 to be independent of phosphate metabolism.

## 4. Pathogenesis of FGF23-Related Hypophosphatemic Diseases

### 4.1. FGF23-Related Hypophosphatemic Diseases

Since the FGF23- $\alpha$ Klotho axis plays a central role in phosphate homeostasis, its disruption causes hyperphosphatemic conditions, as described in the previous section. On the other hand, the excessive action of FGF23 underlies various hypophosphatemic diseases, which are characterized by urinary phosphate wasting, hypophosphatemia, and inappropriately low levels of serum 1,25(OH)<sub>2</sub>D [42,43]. Phosphate is indispensable for skeletal mineralization; therefore, chronic hypophosphatemia due to excessive FGF23 leads to rickets in children and osteomalacia in adults. The impaired production of 1,25(OH)<sub>2</sub>D contributes to the resistance of FGF23-related hypophosphatemic rickets/osteomalacia to native vitamin D. FGF23-related hypophosphatemic rickets/osteomalacia include various conditions such as genetic diseases (Tables 1 and 2). The following sections will describe the pathogenesis of each condition.

Disorder	Causes	Incidences
ADHR	Activating variants of FGF23	Very rare
XLH	Inactivating variants of PHEX	1:20,000 live birth
ARHR1	Inactivating variants of DMP1	Very rare
ARHR2	Inactivating variants of ENPP1	Very rare
Raine syndrome	Inactivating variants of FAM20C	Very rare
Osteoglophonic dysplasia	Activating variants of FGFR1	Very rare
Jansen's metaphyseal chondrodysplasia	Activating variants of PTH1R	Very rare
TIO	Phosphaturic mesenchymal tumor, FN1-FGFR1, FN1-FGF1 fusion genes	Most frequent among the acquired disorders
McCune-Albright Syndrome	GNAS somatic activating variants in bone lesions	Very rare
Linear sebaceous nevus syndrome	KRAS/HRAS/NRAS somatic activating variants in skin/bone lesions	Very rare
I.V. administration of iron preparations	Saccharated ferric oxide, iron polymaltose	Very rare

Table 1. FGF23-related hypophosphatemic diseases and their causes.

#### 4.2. Autosomal Dominant Hypophosphatemic Rickets (ADHR)

ADHR (OMIM #193100) is caused by missense variants in the FGF23 gene. Since the responsible variants occur at Arg<sup>176</sup> or Arg<sup>179</sup> within the RXXR/S motif recognized by subtilisin-like proprotein convertase, the resultant mutant FGF23 protein is resistant to cleavage-mediated inactivation [5]. However, this disease shows incomplete penetrance. Patients with ADHR variants do not always have high serum levels of intact FGF23, and the disease may occur with an early or delayed onset and variable expressivity [44]. Late-onset ADHR primarily manifests in post-pubertal women who are prone to iron deficiency [45]. Serum iron levels were previously shown to negatively correlate with serum levels of both the C-terminal fragment of FGF23 and intact FGF23 in ADHR patients; however, serum iron levels also negatively correlated with C-terminal FGF23 levels in healthy control subjects, whereas no relationship was observed with intact FGF23 levels [45]. Farrow et al. generated FGF23-knock-in mice carrying the R176Q ADHR point mutation (ADHR mice)

and found that a low-iron diet increased bone FGF23 mRNA levels and the serum levels of both intact and C-terminal FGF23 with hypophosphatemia in ADHR mice. On the other hand, wild-type mice fed the low-iron diet showed normal serum levels of intact FGF23 and phosphate, but an elevated level of the C-terminal fragment of FGF23 [46]. Furthermore, the chelation of iron up-regulated the expression of *Fgf23* in a cultured osteoblastic cell line, which involved hypoxia-inducible factor 1 $\alpha$ . Collectively, these findings suggest that iron deficiency increases the expression of *Fgf23* in bone and also that the FGF23 protein is cleaved in iron deficiency to maintain normal serum levels of FGF23 and normophosphatemia in control subjects, whereas the cleavage resistance of mutant FGF23 leads to the accumulation of intact FGF23 and hypophosphatemia in ADHR subjects [46].

Disorder **Clinical Features** Renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, ADHR rickets/osteomalacia, incomplete penetrance, often delayed onset Renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, XLH rickets/osteomalacia, short stature, bone deformity, dental problems, enthesopathy Renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, ARHR1 rickets/osteomalacia, short stature, bone deformity, dental problems, enthesopathy Renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, ARHR2 rickets/osteomalacia, short stature, bone deformity, dental problems, enthesopathy Osteosclerosis of early onset, poor life prognosis in most cases, characteristic face, renal Pi Raine syndrome wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, rickets/osteomalacia, short stature, bone deformity, dental problems, enthesopathy Multiple radiolucent areas in metaphysis, rhizomelic short stature, characteristic face, renal Pi Osteoglophonic dysplasia wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels Jansen's metaphyseal Short-limbed short stature, usually hypercalcemia, skull sclerosis, renal Pi wasting, chondrodysplasia hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels Renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, TIO rickets/osteomalacia, improved by tumor resection Fibrous dysplasia, café-au-lait skin pigmentation, endocrinologic abnormalities, renal Pi McCune-Albright Syndrome wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels Linear sebaceous nevus, abnormalities in central nervous system, ocular anomalies, bone Linear sebaceous nevus syndrome defects, renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)2D levels I.V. administration of iron Renal Pi wasting, hypophosphatemia, low or normal serum 1,25(OH)<sub>2</sub>D levels, preparations rickets/osteomalacia, improved by discontinuation of the responsible drugs

Table 2. Clinical features of FGF23-related hypophosphatemic diseases.

#### 4.3. X-Linked Hypophosphatemic Rickets (XLH)

XLH (OMIM #307800) is the most common form of hereditary hypophosphatemic rickets. Patients with XLH have elevated serum levels of FGF23, which result in urinary Pi wasting, hypophosphatemia, and inappropriately low levels of  $1,25(OH)_2D$  [47]. XLH was initially called vitamin D-resistant rickets because of a poor response to treatment with native vitamin D at dosages that cure vitamin D-deficient rickets. XLH is caused by inactivating variants in the *phosphate-regulating gene homologous to endopeptidase on the X chromosome* (*PHEX*) located at Xp22.1, showing X-linked dominant inheritance [48]. Similar to FGF23, PHEX is expressed in osteoblast lineage cells and is more highly expressed in osteocytes [18,49]. Although its structure suggests that the product of *PHEX* functions as a cell surface-bound, zinc-dependent protease, its physiological substrates remain elusive. In hypophosphatemic *Hyp* mice, which harbor a large deletion in the *Phex* gene and are widely used as a model for XLH, the expression of *Fgf23* in osteocytes was found to be

increased [18,50]; however, FGF23 did not serve as a substrate for PHEX [51]. Therefore, the regulation of FGF23 by PHEX may be indirect and involve other molecule(s).

Since PHEX is highly expressed in osteocytes, in an attempt to clarify abnormalities in *Phex*-deficient osteocytes, we previously compared the gene expression profiles of osteoblasts and osteocytes in Hyp mice and wild-type littermates [18]. As osteoblasts mature into osteocytes, the expression of *dentin matrix protein* 1 (Dmp1) and *family with sequence* similarity 20c (Fam20c), which are responsible for autosomal recessive hypophosphatemic type 1 (ARHR1) and Raine syndrome (RNS), respectively, increased in both Hyp and wild-type cells, and these genes were up-regulated in Hyp cells, similar to Fgf23 [18]. These findings indicated the critical roles of osteocytes in phosphate homeostasis and also suggested complex abnormalities in *Phex*-deficient osteocytes. We found that the expression of the genes encoding canonical FGF ligands (Fgf1 and Fgf2), their receptors (Fgfr1-3), and early growth response 1, which is a target for FGFR activation, was also upregulated in *Hyp* osteocytes, indicating enhanced FGFR signaling [18]. Furthermore, Martin et al. suggested enhanced FGFR signaling in Hyp bone [52], and Xiao et al. demonstrated that the conditional deletion of *Fgfr1* in osteocytes and mature osteoblasts partially restored the overproduction of FGF23 and ameliorated hypophosphatemia and rickets [53]. The regulation of FGF23 production by FGFR signaling is also supported by osteoglophonic dysplasia, which is a rare skeletal dysplasia caused by activating mutations in *FGFR1* that is frequently associated with elevated serum FGF23 levels and hypophosphatemia [54].

Enhanced FGFR signaling in *Phex*-deficient osteocytes is of interest based on previous findings suggesting that FGFR plays a critical role in the transduction of signaling evoked by increased extracellular Pi [55–57]. In various cell types, treatment with high extracellular Pi activated FGFR for the regulation of gene expression. In an osteoblastic cell line, treatment with an FGFR inhibitor abolished the up-regulation of Dmp1 by increased extracellular Pi [56]. In HEK293 cells, the knockdown of FGFR1 diminished the Pi-induced phosphorylation of ERK1/2 [55]. More recently, the activation of FGFR1 by extracellular Pi was shown to increase the expression of *Galnt3* in bone, leading to an elevated serum level of FGF23 in mice [57]. Collectively, these findings suggest that FGFR1 is involved in the sensing of Pi availability. In consideration of this role of FGFR and enhanced FGFR signaling in the osteocytes of *Hyp* mice, abnormal Pi sensing may be involved in the pathogenesis of XLH.

Matrix extracellular phosphoglycoprotein (MEPE) is a member of the SIBLING (small integrin-binding ligand, N-linked glycoproteins) family, and was initially cloned from the tumor of a patient with TIO [58]. A genome-wide association study proposed MEPE as a factor influencing bone mineral density in humans [59]. MEPE contains an acidic serine-aspartate rich MEPE-associated motif (ASARM) consisting of 23 residues at the C terminus. ASARM peptides released from MEPE by cathepsin-mediated cleavage have been shown to inhibit mineralization. Rowe et al. demonstrated that PHEX bound to the ASARM motif in MEPE and the released ASARM peptide and its serum levels were elevated in *Hyp* mice [60,61]. The ASARM motif is also present in other SIBLING proteins, such as DMP1 and osteopontin, and PHEX may bind to these proteins at these motifs [58]. A previous study implicated ASARM peptides released from these SIBLING proteins in defective mineralization in XLH [62].

Growth retardation is often observed in patients with XLH: however, the underlying mechanisms have not yet been elucidated in detail. A study published by Fuente et al. demonstrated marked alterations in the structure, dynamics, and maturation of growth plate cartilage in growth-retarded young *Hyp* mice [63]. In the growth plates of *Hyp* mice, both proliferation and apoptosis rates of chondrocytes were reduced, and the hypertrophy and maturation of chondrocytes were severely disturbed. The spatial organization of the chondro-osseous junction and the primary spongiosa trabeculae were markedly deformed. These alterations in the growth plates might be the mechanisms for the growth retardation in *Hyp* mice. The authors also found an enhanced activation of the extracellular signal-regulated kinase (ERK)1/2 signaling pathway in the *Hyp* growth plates, implying an

involvement of FGF23 in these abnormalities [63]. Reduction in caspase-mediated apoptosis of hypertrophic chondrocytes was also reported in rachitic mice with low-phosphate dietinduced hypophosphatemia as well as in *Hyp* mice, which suggests that hypophosphatemia impairs apoptosis of hypertrophic chondrocytes, leading to rickets [64].

Although chondrocytes do not express  $\alpha$ Klotho, which is required for FGF23 to activate its downstream signaling pathways at physiological concentrations, soluble forms of  $\alpha$ Klotho are present in serum and cerebrospinal fluid [65] and have been implicated in the regulation of FGF23 signaling in cells without the transmembrane form of  $\alpha$ Klotho [66]. We previously demonstrated that FGF23 suppressed the linear growth of mouse metatarsal cartilage in cultures in the presence of soluble  $\alpha$ Klotho by decreasing the proliferation of chondrocytes, which suggests that suppressed chondrocyte proliferation by FGF23 plays a causative role in the growth retardation associated with XLH [67].

Since the placenta expresses FGFR1 and  $\alpha$ Klotho, high levels of FGF23 in pregnant women with XLH may affect their fetuses. We previously investigated this issue using pregnant *Hyp* female mice [33]. *Hyp* and wild-type female mice were mated with wild-type male mice, and the pregnant mothers and their male fetuses were subjected to analyses. FGF23 levels were higher in *Hyp* mothers than in wild-type mothers. *Hyp* fetuses and wild-type fetuses were obtained from mating between *Hyp* females and wild-type males. FGF23 levels in *Hyp* fetuses were approximately 20-fold higher than in their mothers, while wild-type fetuses from *Hyp* mothers had low levels of FGF23, as did fetuses from wild-type mothers, suggesting that FGF23 does not cross the placenta [33]. The expression of *Cyp24a1* was higher in the placentas of fetuses from *Hyp* mothers than in those of fetuses from wild-type mothers, which resulted in decreased levels of plasma 25-hydroxyvitamin D in fetuses from *Hyp* mothers. Therefore, increased levels of circulating FGF23 in *Hyp* mothers may exert direct effects on the placenta during pregnancy and alter fetal vitamin D metabolism via the regulation of *Cyp24a1* expression [33]. Further studies are needed to clarify whether similar phenomena occur with pregnancy in human patients with XLH.

The enthesis is a tissue that forms at the site of insertion of a tendon to bone and consists of a bony eminence, mineralized fibrocartilage, unmineralized fibrocartilage, and a tendon. It optimizes the transfer of mechanical force from muscle to bone, which is required for efficient movements [68]. Enthesopathy is a pathological change at the insertion of tendons and ligaments. Mineralizing enthesopathy is one of the complications of XLH and other types of FGF23-related hypophosphatemia and accounts for a high morbidity rate in adult patients [69]. Karaplis et al. previously reported that a transgenic mouse model overexpressing a secreted form of the human FGF23[p.R176Q] variant, which is resistant to cleavage, displayed mineralizing enthesopathy of the Achilles and planar facial insertions, suggesting the involvement of FGF23 in the development of mineralizing enthesopathy [70]. More recently, Liu et al. investigated the cellular and molecular mechanisms involved in the development of mineralizing enthesopathy in *Hyp* mice and reported that Achilles tendon entheses of *Hyp* mice showed the expansion of hypertrophic-appearing chondrogenic cells. In comparison with the entheses of wild-type mice, *Hyp* entheses exhibited the expansion of cells expressing the chondrogenic marker gene Sox9 and enhanced bone morphogenetic protein and Indian hedgehog signaling pathways, both of which play critical roles in chondrocyte differentiation [71]. Although oral phosphate salts and active vitamin D metabolites are administered as conventional medical treatments for XLH to correct their deficiencies, it does not prevent or ameliorate enthesopathies [70]. Burosumab, a humanized monoclonal neutralizing antibody to FGF23, has recently become available as a new treatment for XLH [47]. In Japan, burosumab has been approved for the treatment of all types of FGF23-related hypophosphatemic rickets/osteomalacia. In pediatric patients with XLH, improvements in the severity of rickets and biochemical parameters were greater in patients treated with burosumab than in those who continued conventional therapy [72]. Further studies are needed to clarify the effects of burosumab on the prevention and treatment of enthesopathies.

## 4.4. Autosomal Recessive Hypophosphatemic Rickets Type 1 (ARHR1)

ARHR1 (OMIM #241520) is caused by inactivating variants of the *DMP1* gene [73,74]. DMP1 is an extracellular matrix protein belonging to the SIBLING family and is highly expressed in osteocytes as well as in dentin. Patients with ARHR1 manifest elevated FGF23 levels, hypophosphatemia, inappropriately low 1,25(OH)<sub>2</sub>D levels, and skeletal hypomineralization, similar to patients with XLH. *Dmp1*-null mice reproduced the phenotype of ARHR1 and exhibited defective osteocyte maturation and the up-regulated expression of *Fgf23* in osteocytes [73,74]. Although the pathogenesis of ARHR1 remains largely unknown, the findings of studies using *Phex*-deficient *Hyp* mice and *Dmp1*-null mice suggest that the overproduction of FGF23 is attributable to enhanced FGFR signaling in bone in both mouse models [52].

#### 4.5. Autosomal Recessive Hypophosphatemic Rickets Type 2 (ARHR2)

ARHR2 (OMIM #613312) also belongs to FGF23-related hypophosphatemic rickets and is caused by inactivating variants in the *ectonucleotide pyrophosphatase phosphodiesterase-1* (*ENPP1*) gene [75,76]. *ENPP1* encodes an enzyme that produces pyrophosphate (PPi), a potent inhibitor of mineralization, and inactivating variants in *ENPP1* are also responsible for hypermineralization disorders, such as generalized arterial calcification in infancy [77]. The ectoenzyme tissue non-specific alkaline phosphatase (TNSALP) facilitates skeletal mineralization by degrading PPi to produce Pi. Although PPi may regulate the production of FGF23, patients with hypophosphatasia, which is caused by inactivating variants in TNSALP, had normal levels of FGF23 despite elevated extracellular levels of PPi [78]. Therefore, the mechanisms by which ENPP1 deficiency results in the overproduction of FGF23 remain unclear. Since inactivating variants in ENPP1 cause conditions characterized by ectopic calcification and FGF23-related hypophosphatemia, a close relationship may exist between ectopic calcification and the overproduction of FGF23.

### 4.6. Raine Syndrome (RNS)

*FAM20C*, also known as *DMP4*, encodes a kinase that phosphorylates various secreted proteins. The proteins phosphorylated by FAM20C include FGF23 and members of the SIBLING family, such as DMP1, osteopontin, and MEPE [34,79]. Inactivating variants in the *FAM20C* gene are responsible for RNS (OMIM #259775). RNS is an autosomal recessive disease that is characterized by craniofacial malformation, osteosclerotic bone dysplasia, and a poor prognosis [80]. Surviving patients with mild RNS manifest hypophosphatemia due to elevated levels of FGF23 and dental anomalies [81,82]. *Fam20c*-null mice exhibited elevated levels of serum FGF23, hypophosphatemia, and dental anomalies [83]. These mice also showed low expression levels of *Dmp1* in osteocytes, which suggested that the down-regulated expression of DMP1 plays a causal role in the overproduction of FGF23 in RNS [83]. However, the overexpression of *Dmp1* failed to rescue the defects in *Fam20c*-null mice [84]. A previous study reported that FAM20C phosphorylated FGF23 on Ser<sup>180</sup>, which inhibited the *O*-glycosylation of FGF23 on Thr<sup>178</sup> by GalNAc-T3 and accelerated cleavage [34]. Therefore, inactivating variants in FAM20C may increase FGF23 levels by inhibiting its cleavage.

## 4.7. Tumor-Induced Osteomalacia (TIO)

TIO is a rare paraneoplastic syndrome characterized by urinary phosphate wasting, hypophosphatemia, and osteomalacia. Responsible tumors are generally benign, slow-growing phosphaturic mesenchymal tumors (PMT) [85]. The overproduction of FGF23 by tumors was previously shown to enhance the renal excretion of Pi and induce hypophosphatemia, low 1,25(OH)<sub>2</sub>D levels, and osteomalacia, which were cured by the surgical removal of the responsible tumor [6,86]. Lee et al. identified the fusion genes *Fibronectin 1* (*FN1*)-*FGFR1* and *FN1-FGF1* in subgroups of PMT and showed that immunoreactivity for FGFR1 was positive in 82% of PMT [87,88]. These findings suggest the involvement of the FGF/FGFR signaling pathway in the development of PMT.

## 4.8. Other Causes of FGF23-Related Hypophosphatemia

McCune-Albright syndrome (MAS, OMIM #174800) is characterized by polyostotic fibrous dysplasia, café-au-lait skin pigmentation, and precocious puberty, and is caused by a somatic activating variant in *GNAS1* encoding the subunit of the stimulatory G protein. MAS is clinically heterogeneous and may manifest various endocrinological abnormalities. Some patients with MAS exhibit hypophosphatemia, which results from the overproduction of FGF23 by abnormal skeletal progenitor cells in the bone lesions of fibrous dysplasia [89]. Serum levels of FGF23 in MAS patients correlate with disease activity [89], and significant hypophosphatemia only occurs in patients with a severe disease burden. A previous study suggested that the ratio of the C-terminal fragment of FGF23 to intact FGF23 was elevated by accelerated cleavage in the bone lesions of fibrous dysplasia [90].

Linear sebaceous nevus syndrome, also called Schimmelpenning-Feuerstein-Mims (SFM) syndrome (OMIM #163200), is characterized by congenital linear nevus sebaceous and abnormalities in neuroectodermal organs and is caused by somatic variants in *RAS* genes, including *KRAS*, *HRAS*, and *NRAS*, which are detectable in skin lesions [91,92]. Hypophosphatemia due to elevated levels of FGF23 is rarely associated with SFM syndrome. Lim et al. suggested that the source of FGF23 in SFM syndrome was bone lesions carrying *RAS* variants rather than skin lesions [93].

Osteoglophonic dysplasia (OMIM #166250) is a rare autosomal dominant disease characterized by rhizomelic dwarfism, non-ossifying bone lesions, craniosynostosis, and face abnormalities, and is caused by activating variants in the *FGFR1* gene. As discussed earlier, this disease may be associated with FGF23-related hypophosphatemia, indicating the involvement of FGFR1 in the regulation of FGF23 production [54].

Jansen's metaphyseal chondrodysplasia (OMIM #156400) is an autosomal dominant disease caused by an activating variant in the *PTH type 1 receptor* (*PTH1R*) gene [94]. Previous studies reported that FGF23-related hypophosphatemia may be associated with Jansen's metaphyseal chondrodysplasia [95]. This finding suggests that PTH signaling stimulates FGF23 production, which is also supported by the findings of several in vivo and in vitro studies [31,32,96].

FGF23-related hypophosphatemic rickets/osteomalacia may also be associated with the intravenous administration of saccharated ferric oxide or iron polymaltose [97,98]. The mechanisms by which these drugs cause the overproduction of FGF23 remain unclear; how-ever, their discontinuance rapidly restores elevated FGF23 levels and hypophosphatemia.

## 5. Conclusions

FGF23-related hypophosphatemia is characterized by urinary Pi wasting, hypophosphatemia, and inappropriately low levels of 1,25(OH)<sub>2</sub>D, and includes various types of hereditary rickets/osteomalacia, such as XLH, and acquired diseases, including TIO. The molecules responsible for hereditary rickets/osteomalacia are highly expressed by osteocytes, indicating that these cells play a central role in phosphate homeostasis. Since inactivating variants of PHEX, DMP1, ENPP1, and FAM20C lead to the overproduction of FGF23, these molecules appear to function as negative regulators of FGF23. Although the mechanisms underlying the overproduction of FGF23 remain unclear in most FGF23related hypophosphatemic diseases, enhanced FGFR signaling may be involved in the overproduction of FGF23 in XLH and ARHR1 as well as in TIO. Since FGFR1 is suggested to be involved in Pi sensing, abnormalities in Pi sensing may play a role in the pathogenesis of these diseases.

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