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# Recent advances in primary cilia in bone metabolism

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Primary cilia are microtubule-based organelles that are widespread on the cell surface and play a key role in tissue development and homeostasis by sensing and transducing various signaling pathways. The process of intraflagellar transport (IFT), which is propelled by kinesin and dynein motors, plays a crucial role in the formation and functionality of cilia. Abnormalities in the cilia or ciliary transport system often cause a range of clinical conditions collectively known as ciliopathies, which include polydactyly, short ribs, scoliosis, thoracic stenosis and many abnormalities in the bones and cartilage. In this review, we summarize recent findings on the role of primary cilia and ciliary transport systems in bone development, we describe the role of cilia in bone formation, cartilage development and bone resorption, and we summarize advances in the study of primary cilia in fracture healing. In addition, the recent discovery of crosstalk between integrins and primary cilia provides new insights into how primary cilia affect bone.

#### KEYWORDS

primary cilia, bone formation, chondrogenesis, bone resorption, fracture healing, integrins

### **1** Introduction

Cilia are highly conserved organelles distributed on the cell surface and present in most cells of the human body (1). The cilium consists of a basal body and an axoneme (2), and its formation is mainly carried out through a transport mechanism that is completely homologous to that of Chlamydomonas reinhardtii and IFT proteins (3) which is closely related to the cell cycle (4), and it is a complex process that involves the assembly of microtubule structures and protein transport (5), including the fusion of centromeres with the plasma membrane (3) ciliary secretion (6), and the transport of Golgi vesicles (7). Early studies suggested that there are two modes of ciliary axoneme formation, namely nine double microtubules surrounding a central pair of single microtubules (9 + 2); and a central pair of microtubules missing (9 + 0) mode, due to the lack of axonemal dynamo proteins responsible for ciliary motility in 9 + 0 cilia are considered immotile and are also referred to as primary cilia (8).

However, with the in-depth study of primary cilia, some studies have found that the three-dimensional structural model of primary cilia established by cryo-cryo-tomography reveals that the structure of primary cilia is different from that of the "9 + 0" pattern (9),

including a recent study that found that the "9 + 0" pattern arrangement exists only in the matrix of the primary cilium and that the structure of primary cilium changes to unstructured bundles of EB1-decorated microtubules and actin filaments just a few micrometers away from the matrix (10). Primary cilia contain hundreds and thousands of proteins, but interestingly primary cilia do not synthesize any proteins themselves. Ciliary proteins are mainly dependent on the intraflagellar transport system (IFT) for transport to the cilium. IFT is a bidirectional transport system operated by IFT protein complexes (IFT complex A, IFT complex B) and IFT motors, which play an important role in cilia assembly and maintenance (11, 12). IFT is regulated by the transition zone (TZ) at the base of the cilia (13), and during cilia assembly activates kinesin-II to transport large protein complexes from the cytosol to the tip of the cilium, the transported proteins are dissociated and released, and then kinesin-II is inactivated, kinesin-II is activated and binds to the IFT protein complex a, and retrograde movement towards the base of the cilium (4). Also, IFT is required to mediate many signaling functions (14) IFT mediates Hedgehog, Wnt, PDGFR, Notch, TGF-β, mTOR and other signaling pathways have been reported in detail in other reviews (4, 15, 16).

A large body of data shows that primary cilia play a crucial role in vertebrate development and human genetic diseases by regulating a variety of fundamental processes that contribute to the maintenance of cellular activity and tissue homeostasis, including cell differentiation and vertebrate development, cell cycle control, and signaling (17, 18). IFT is capable of initiating signaling pathways linking mechanical or chemical stimuli to intracellular transduction cascades that act at various sites in the body (19, 20). Many human diseases are associated with defects in ciliary structure or localized proteins in cilia (21). In recent years, it has been found that ciliopathy symptoms often involve skeletal development, highlighting the importance of primary cilia in skeletal development. This review summarizes recent findings on the role of primary cilia and the ciliary transport system in osteogenesis and development, We describe the role of cilia in bone formation, cartilage development, and bone resorption, and we also summarize advances in the study of primary cilia in fracture healing.

# 2 The role of primary cilia in bone metabolism

### 2.1 The role of primary cilia in bone formation

The skeletal system is a dynamic and highly coordinated system that involves a balance between bone formation by osteoblasts and bone resorption by osteoclasts (22). Bone formation is predominantly executed by osteoblasts (23), and primary cilia have been shown to have an important role in osteogenesis (24– 26). It has been shown that primary cilia regulate the osteogenic differentiation of bone marrow-derived mesenchymal stem cells (27) and adipose-derived stem cells (28), and that they are involved in the process of osteoblast alignment and differentiation (29),

regulating skeletal development in the embryonic period and bone formation in adulthood (30), and that knockdown of primary cilia reduces load-induced bone formation (31). In recent years, the role of pulsed electromagnetic fields in promoting bone formation has been gaining attention, however, primary cilia play a key role in this. It has been shown that primary cilia act as sensory organs involved in pulsed electromagnetic field-mediated bone formation (32). He (33) found that PEMF stimulates osteogenic differentiation mainly by activating the NOS/NO/sGC/cGMP/PKG signaling pathway and thus stimulating osteogenic differentiation, and it is noteworthy that all components of this signaling pathway, including iNOS, eNOS, sGC, PKG-1, and PKG-2 are localized in the primary cilia, and that eNOS needs to be phosphorylated within the primary cilia in order to be phosphorylated. As well as Liu (25) also reported that the primary cilia of osteoblasts in rats exposed to microgravity rapidly became shorter or even disappeared, and this phenomenon was accompanied by a significant reduction in osteogenic differentiation and mineralization of osteoblasts. In addition, it has been shown that cellular function can barely be maintained under conditions without mechanical loading and the effect of inducing osteoblasts to become bone cells is greatly limited (34), whereas primary cilia, as an important part of mechanotransduction, are involved in mechanosensing and signaling that regulates the behavior of mesenchymal bone progenitors, osteoblasts and osteoclasts (35).

The IFT protein family also plays an important role in bone formation (36), and the knockdown of IFT proteins leads to a reduction in mechanically stimulated paracrine signaling that promotes osteogenesis and affects osteogenesis (37). Guleria (23) found that silencing of IFT52 disrupts primary ciliogenesis preventing normal osteogenic differentiation. IFT52 silencing resulted in a significant decrease in mRNA levels of runt-related transcription factor 2 (Runx2), secreted phosphoprotein 1 (Spp1) and bone gamma carboxyglutamate protein (Bglap). the decrease in Patched-1 (Ptch1) demonstrated that silencing of Ift52 attenuates the upregulation of this microsignaling pathway (Hh). IFT140 as a core protein of IFT complex A has been shown to be closely associated with bone formation marker expression during differentiation of BMMSCs (38). Tao (39) established the Osx-Cre; Ift140flox/flox mouse model and found that the mice exhibited dwarf phenotypes such as short bone length, low bone mass and reduced bone mineral deposition rate as well as reduced expression of osteoblast markers. These results suggest that IFT140 is indispensable in the process of bone formation. Interestingly, IFT140 is temporally and spatially specific during bone formation, with the current study indicating that IFT140 mainly acts during the early stages of osteoblast differentiation (40). In addition, the differentiation programs of adipogenesis and osteoclastogenesis are in competition with each other and balance each other, with mechanisms that promote adipogenesis inhibiting osteoclastogenesis (41). A recent study demonstrated that IFT20 controls MSCs allocation by regulating glucose metabolism during skeletal development. IFT20 deficiency in MSCs promotes adipocyte formation, which enhances RANKL expression and reduces bone formation. Mechanistically, IFT20 deletion in MSCs reduces glucose tolerance, thereby inhibiting glucose uptake, lactate

and ATP production. In addition, the deletion of IFT20 significantly reduced the activity of TGF- $\beta$ -Smad2/3 signaling and decreased the binding activity of Smad2/3 to the Glut1 promoter, thereby downregulating Glut1 expression. These results suggest that IFT20 plays an important role in the balance between osteogenesis and adipogenesis through the TGF- $\beta$ -Smad2/3-Glut1 axis (42). In addition, numerous reports of Ift80 and Ift88 being ablated in developing mouse limbs lead to defects in skeletal pattern formation (43–45), as well as a large number of skeletal ciliopathies in which mutations have been found on IFT43, IFT121, IFT122, and IFT80, IFT52, IFT172, and IFT56 (46).

In addition, it was found that polycystic kidney disease (PKD) gene product polycystin-1 (PC1), which is localized to primary cilia, can regulate bone development and osteoblast function through the regulation of the osteogenic transcription factor Runx2-II, and modulates intracellular calcium-dependent signaling in osteoblasts, and that depletion of intracellular calcium blocks PC1 activation of the Runx2-II P1, demonstrating the importance of the intracellular calcium pathway in PC1-mediated Runx2-II P1 promoter activity in osteoblasts. It is suggested that the polycystin complex in bone functions as an intracellular pathway that senses external developmental signals and translates them into the regulation of Runx2-II expression during bone formation (47). Moreover, Xiao (48) found that Pkd1 has a mechanotransduction role in bone, and conditional deletion of Pkd1 in osteoblasts revealed the reductions of bone mineral density, bone trabecular volume, and expression of Runx2, Osteocalcin, and Dmp1 in Dmp1-Cre; Pkd1flox/+, and Pkd1<sup>Dmp1-cKO</sup> mice have all correlated with the Pkd1 gene dosage. Furthermore, Pkd1-deficient osteoblasts showed a markedly diminished response to fluid shear stress in vitro. These results suggest that Pkd1 has a direct mechanosensing role in osteoblasts to regulate osteoclast function and skeletal homeostasis. As well, studies have reported that silencing Pkd2 in osteoblasts also leads to reduced Runx2 expression and impaired biomechanical properties in mouse bone, thereby affecting bone formation. Interestingly, deletion of Pkd2 also led to reduced PPARy expression, resulting in reduced bone marrow fat in vivo (49). These results suggest that Pkd1 and Pkd2 have synergistic roles in osteoblast differentiation and opposite roles in adipogenesis, and we therefore suggest that the Pkd1 and Pkd2 signaling pathways can independently influence the mesenchymal spectrum of bone (50, 51). Qiu (52) explored the mechanism and found that the expression of Gli2, a gene downstream of the Hh signaling pathway, and Axin2, a gene downstream of Wnt/β-catenin, were significantly reduced in osteoblasts of Kif3aOc-cKO mice compared with Kif3aflox/+ mice, and these results indicate that Kif3a deficiency affects the transduction of signaling pathways, such as Hh and Wnt/β-catenin, which in turn affects bone formation. However, some studies have found that the effect of Kif3a on osteoblasts may be mediated through polycystins (53), which needs to be further investigated.

We summarize recent relevant studies exploring the role of IFT proteins in bone formation (Table 1).

Skeletal tissue development in mammals is complex and involves multiple signaling pathways interacting with each other, Primary cilia on osteoblasts are the arena for the confluence of many signaling pathways (17), and the interactions between primary cilia-related signaling pathways during osteogenesis are provided in the figure (Figure 1).

## 2.2 The role of primary cilia in bone resorption

Osteoclasts (OCs) are the cells responsible for physiological bone resorption and play an important role in maintaining bone

| TABLE 1 | Intraflagellar | transport i | n bone | formation. |
|---------|----------------|-------------|--------|------------|
|---------|----------------|-------------|--------|------------|

| Name         | Study   | Ref  |
|--------------|---|------|
| Ift52        | Silencing of Ift52 impaired primary ciliogenesis and hindered osteogenic differentiation. Up-regulation of the Hedgehog (Hh) pathway during osteogenesis is attenuated in Ift52-silenced cells  |      |
| IFT140       | Osx-Cre; If1140flox/flox mice exhibit pygmy phenotypes such as short bone length, low bone mass and reduced bone mineral deposition rate and reduced expression of osteoblast markers.  |      |
| Pkd1/<br>PC1 | Deletion of Pkd1 in Mature Osteoblasts, Reduced Expression of Runx2-II, Osteocalcin, Dmp1 and Phex in Bone of Pkd1flox/+ and Pkd1Dmp1-cKO Mice.   | (48) |
| Pkd2/<br>PC2 | Silencing of Pkd2, Oc-Cre;Pkd2flox/null (Pkd2Oc-cKO) mice exhibit reduced bone mineral density, trabecular volume, cortical thickness, mineral deposition rate, and impaired biomechanical properties of bone.Pkd2 deficiency leads to reduced Runx2 expression in bone and affects osteoblast differentiation <i>in vivo</i> . | (49) |
| Kif3a        | Colα1(I) 2.3-Cre;Kif3afl/fl mice have reduced new bone formation under mechanical ulnar loading.  | (51) |
|              | Kif3aOc-cKO mice developed osteoporosis at 6 weeks of age, as evidenced by reduced femoral mineral density, trabecular volume, and cortical thickness. Shh-mediated expression of Gli2 as well as Wnt3a-mediated expression of β-catenin and Axin2 were attenuated.   | (52) |
| Ift20        | Using Osterix-Cre and inducible type I collagen-CreERT, it was found that deletion of IFT20 in osteoblast cell lines resulted in impaired cell alignment and reduced bone mass. The results indicate that IFT20 regulates polarity and cell alignment during bone development through ceramide-pPKCζ-β-catenin signaling        | (54) |
| IFT46        | Knockdown of IFT46 results in the absence of cilia in different tissues.IFT46 morphants exhibit axial shortening, defective neural tube closure, and abnormal craniofacial development.   | (55) |
| SAC          | PEMF stimulation promoted intracellular sAC expression and increased cAMP concentration in hypoxia-exposed primary osteoblasts. Blockade of intracellular sAC inhibited the PEMF-induced decrease in HIF-1 $\alpha$ expression and increase in osteoblast differentiation.  | (56) |

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**FIGURE 1** Crosstalk in primary cilia-related signaling pathways during osteogenesis. BMP inhibits Wnt/β-catenin signaling by increasing the expression of the Wnt antagonists Dkk1 and Sost and blocking β-catenin nuclear translocation; on the other hand, BMP promotes Wnt/β-catenin signaling by forming a co-transcriptional complex with β-catenin/Runx2 and antagonizing the function of Dvl by increasing the expression of Wnt. TGF-β promotes Wnt activity by increasing Wnt ligand expression. Hh signaling promotes BMP signaling by enhancing BMP-2 expression through Gli. In addition, Notch intracellular structural domain (NICD1) enhances BMP-induced angiogenesis and thus promotes bone formation. BMP, Bone Morphogenetic Proteins; Runk2, Recombinant Runt Related Transcription Factor 2; TGF-β, transforming growth factor-β; ERK1/2, extracellular regulated protein kinases; Hh, Hedgehog; SMO, Smoothene.

mass homeostasis in the body (57). The current study indicates that osteoclasts derived from the hematopoietic lineage do not have primary cilia (58), but nevertheless, primary cilia are essential for their occurrence. PEMFs were found to inhibit osteoclast differentiation by scavenging reactive oxygen species (59) and modulating the Akt/mTOR signaling pathway (60). Wang (61) found that PEMFs affect bone resorption mainly by regulating RANKL/OPG to inhibit osteoclast formation, and this process is closely related to primary cilia, and its deletion of primary cilia was found to significantly enhance osteoclast formation and bone resorption. Parathyroid hormone (PTH) receptor 1 (PTH1R) is a B-group G protein-coupled receptor (GPCR), which plays an important role in bone metabolism, and activation of PTH1R was found to regulate osteoclasts. Tirado-Cabrera (62) reported the localization of PTH1R to primary cilia, which confirmed that primary cilia in osteoblasts are necessary for proper communication with osteoclasts.IL-6 secretion is increased when PTH1R is silenced. Previous studies have shown that IL-6 enhances osteoblast-mediated osteoclastogenesis by promoting the activities of JAK2 and RANKL (63). IFT80 can promote tumor necrosis factor (TNF) receptor-associated factor 6 (TRAF6) proteasomal degradation through binding to Casitas B-lineage lymphoma protoogene-b (Concbl-b), which regulates OC formation. Knockdown of IFT80 leads to Cbl-b ubiquitination and elevated TRAF6 levels, which overactivated the nuclear factor  $\kappa\beta$  (NF- $\kappa\beta$ ) receptor

activator (RANKL) signaling axis (RANK/RANKL), thereby increasing OC formation, while IFT80 overexpression significantly inhibits OC formation and the downstream signaling pathway of RANKL/RANK (64).

# 2.3 The role of primary cilia in cartilage development and repair

Chondrogenesis denotes an intricately intricate process (65). While the importance of primary cilia in the progression of cartilage has been previously examined (66), recent investigations have furnished novel evidence regarding certain factors localized to primary cilia, including THM2, PTHrP, GRK2, and Gpr161, which participate in the development of cartilage (Table 2).

In addition, due to the unique composition of cartilage, its restoration following an injury proves highly challenging. Recently, Tao (81) found that activation of ciliated Hh signaling by Smoothened agonists (SAG) when the growth plate is damaged significantly accelerates growth plate repair after injury, and his further study found that chondrocytes in the resting and proliferating zones are dynamically ciliated during growth plate repair. Furthermore, conditional deletion of the cilia core gene Ift140 in cartilage disrupts cilia-mediated Hh signaling in the growth plate. These results show that ciliary Hh signaling by

| Name          | Study related to cartilage development   | Ref               |
|---------------|--|-------------------|
| THM2          | The small phenotype of Thm2-/-, Thm1aln/+ mouse mutants is accompanied by structural defects in the small mandible and long bones (tibia) and decreased bone mineral density.  | (46)              |
| PTHrP         | PTHrP chondrocytes are maintained in a Wnt-inhibited environment in the resting zone of the growth plate and regulate the formation, expansion and differentiation of chondrocytes in the resting zone.  | (67)              |
| GRK2          | GRK2 deficiency leads to ATD through defects in ciliary signaling; GRK2 is a positive regulator of Hh signaling, and its absence leads to inhibition of GL13 activity. inactivation of GRK2 leads to defects in cilia accumulation in SMO of chondrocytes as well as impaired Wnt signaling typical of GRK2-/- cells.                        | (68)              |
| crocc2        | crocc2 affects chondrocyte ability to sense and respond to the mechanical environment, with downstream effects on cartilage morphogenesis.   | ( <del>69</del> ) |
| FGF2/<br>FGF8 | FGF2 and FGF8 are important growth factors for the growth of mandibular condylar cartilage in young mice and have a limited role in cartilage of aged mice.  | (70)              |
| Dicam         | Dicam expressed in quiescent and proliferating chondrocytes of the growth plate enhances growth plate chondrocyte proliferation and maturation with enhanced Ihh and PTHrP signaling.  | (71)              |
| NG2           | NG2/CSPG4 affects chondrocyte differentiation by ERK1/2 signaling  | (72)              |
| TRPV4         | TRPV4 mutations alter BMP signaling in chondrocytes and prevent proper chondrocyte hypertrophy   | (73)              |
| PC2           | Polycystin is required for mechanical transduction of chondrocytes   | (74)              |
| HDAC6         | Overexpression of HDAC6 causes mitochondrial dysfunction and promotes reactive oxygen species production, leading to ECM degradation.  | (75)              |
| Arl13b        | Associated with altered hedgehog signaling in chondrocytes.  | (76,<br>77)       |
| HIF-2α        | Up-regulation of HIF-2 α expression increases the expression of multiple catabolic genes and promotes ECM degradation in articular cartilage.  | (78)              |
| Gpr161        | Knockdown of Gpr161 affected endochondral bone formation and prevented periarticular chondrocytes from differentiating into columnar chondrocytes while causing a lack of Ihh signaling.   | (79)              |
| Piezo1        | Overexpression of Piezo1 in osteoarthritic cartilage and cultured chondrocytes under shear stress and Piezo1 silencing inhibits nuclear translocation<br>of YAP and down-regulates MMP13 and ADAMTS5 expression Activation of Piezo1 promotes mechanically-induced cartilage degradation through<br>the YAP-MMP13/ADAMTS5 signaling pathway. | (80)              |

TABLE 2 The role of proteins or signalling molecules on primary cilia in cartilage development.

THM2, modifier of Hedgehog signaling; PTHrP, Recombinant Parathyroid Hormone Related Protein; GRK2, G protein-coupled receptor kinase 2; Crocc2, encodes a key gene of the ciliary rootlets; FGF, Fibroblast Growth Factor; Dicam, Proteins localized to primary cilia; TRPV4, nonselective cation channel; PC2, tubulin deacetylase; HDAC6, a tubulin deacetylase; Arl13b, key regulator of the ciliary trafficking; HIF-2α, DNA-binding transcription factor; Gpr161, orphan G-proteincoupled receptor.

chondrocytes in the growth plate coordinates the cartilage regeneration response to GP injury. Additionally, primary cilia have also been reported to be involved in mechanical stress-mediated cartilage repair via the ERK/mTOR axis interacting with autophagy (82).

# 2.4 The role of primary cilia in fracture healing

Fracture healing is a significant clinical concern, and research has indicated that primary cilia and the ciliary transport system play a crucial role in the process of fracture healing. Through the targeted disruption of IFT88 in Prx1-Cre mice, Moore (65) observed that mice lacking primary cilia exhibited abnormally enlarged calluses, a substantial decrease in bone formation, and the presence of persistent cartilage nodules. Analysis of mRNA expression in early cartilage injury tissue revealed downregulation of osteogenesis, Hh signaling, and Wnt signaling, along with an upregulation of chondrogenesis and angiogenesis. Functionally, the deficiency of primary cilia resulted in delayed bone healing. Liu (83) conducted an investigation on the involvement of IFT80 protein in fracture healing, utilizing tamoxifen-induced Col2 $\alpha$ 1CreER mice. The study revealed that Col2 $\alpha$ 1-creIFT80f/f mice displayed smaller fracture calluses compared to IFT80f/f mice. Furthermore, the Col2 $\alpha$ 1-creIFT80f/f mice exhibited low-density/ porous woven bone tissue with significantly reduced bone volume, number of trabeculae, trabeculae thickness, and trabeculae spacing compared to the control mice. These findings indicate that the deletion of IFT80 led to a notable downregulation in the expression of angiogenic markers such as VEGF, PDGF, and angiopoietin, resulting in impaired vascularization of the fracture callus. From a mechanistic perspective, IFT80 deficiency reduces chondrocyte proliferation, cilia assembly and cartilage gene expression and differentiation by inhibiting TGF- $\beta$ I and TGF- $\beta$ R expression and Smad2/3 phosphorylation in fracture calluses, thereby downregulating the TGF- $\beta$  signaling pathway.

In addition, primary cilia serve as central hubs for signaling and can directly facilitate signaling pathways such as Hh, Wnt, and TGF $\beta$ . Serowoky (84) provided evidence demonstrating the essential role of the Hh signaling pathway in fracture healing. Research studies have indicated that Hh signaling is initially downregulated during the early phase of fracture repair, but becomes upregulated in the periosteum during the later stages of the healing process. Research studies have indicated that Hh signaling is initially downregulated during the early phase of fracture repair, but becomes upregulated during the early phase of fracture repair, but becomes upregulated during the early phase of fracture repair, but becomes upregulated in the periosteum

during the later stages of the healing process. However, Moore's findings, using lineage tracing of prx+ cells, showed that the increased Hh signaling did not align with Prx1-expressing cells and their descendants in the corpus callosum and new bone tissue (65). These results suggest that the regulation of Hh signaling in the absence of cilia is non-autonomous. Interestingly, in contrast to previous studies that reported an important role for IHH in the healing of mandibular fractures in zebrafish, Serowoky found a large role for SHH in rib fractures in mice, His findings indicated an increased expression of SHH in the formation of intrachondral and intramembranous bone at the fracture site (84). Notably, the conditional knockdown of Shh during the late stage of fracture severely hindered the formation of cartilage callus and impeded fracture healing. Moreover, the upregulation of Shh expression and the requirement for Shh and Smo seemed to be specific to massive fracture injuries, as they were not observed in small fracture healing scenarios. Intriguingly, it has been observed that Shh cKO mice do not depend on the Hh mechanism for repair near the cut end, but instead rely on Shh for fracture healing in the central region of the affected site (84). Additionally, the classical Wnt/ $\beta$ -catenin signaling pathway is upregulated during fracture healing and contributes to the promotion of repair (85). Moore's research demonstrated that the inhibition of Wnt led to an enlarged corpus callosum in wild-type mice, which aligns with the phenotype observed in mice lacking cilia (65). Furthermore, it has been discovered that TGF- $\beta$  signaling is upregulated during the initial phases of fracture healing (86). This signaling pathway promotes fracture repair by binding to cell surface receptors and triggering the phosphorylation of Smad proteins, particularly Smad2/3. The activation of Smad2/3 has various effects on fracture healing. It enhances the proliferation and differentiation of mesenchymal stem cells, stimulates chondrogenesis, increases the production of matrix proteins, and promotes neointima formation in fracture calluses (87). Additionally, bone morphogenetic protein-2 (BMP-2), a member of the TGF- $\beta$  superfamily, plays a significant role in activating osteoblasts and promoting bone formation. Activation of BMP-2 phosphorylates Smad and facilitates its translocation to the nucleus, activating p38 signaling. This, in turn, upregulates the expression of Runt-associated transcription factor 2 (RUNX2), which further promotes the process of fracture healing (88).

In recent years, there has been growing interest in understanding the association between diabetes and fracture risk. Numerous studies have highlighted that type 1 diabetes mellitus (T1DM) can impede fracture healing by impacting osteoblasts and their precursor cells (89, 90). Recently, Liu conducted a study to examine the impact of T1DM on primary cilia using a mouse model of streptozotocin-induced diabetes. The study also investigated the influence of primary cilia on fracture healing by utilizing osteoblast cell lines with IFT80 deletion through the use of OSX;IFT80f/f mice. The findings of the study demonstrated that diabetes hindered the expression of cilia genes and impeded the formation of primary cilia to a similar degree as observed in normoglycemic mice with IFT80 deficiency. Furthermore, both diabetic mice and normoglycemic mice with osteoblast cilia deletion exhibit delayed fracture healing significantly reduced bone density and mechanical strength, and decreased vascular formation. These results confirm the importance of primary cilia in bone regeneration during the

fracture healing process (83). Indeed, a recent report by Chinipardaz highlighted that diabetes leads to the loss of primary cilia in chondrocytes and osteoblasts. Furthermore, the study revealed that in the diabetic environment, FOXO1 plays a significant role in reducing cilia formation by inhibiting the expression of cilia genes. As a result, this disruption in primary cilia contributes to the delay in the fracture healing process (24). This finding sheds new light on the critical role of primary cilia in the context of fracture healing and offers valuable insights into the mechanisms involved.

#### 2.5 On the crosstalk between primary cilia and integrins in bone

Integrins are heterodimeric transmembrane receptors composed of  $\alpha$  and  $\beta$  subunits that link the extracellular matrix to the intracellular skeleton at different focal adhesion (FA) sites (91). Integrins are linked to primary cilia via the actin cytoskeleton, recent research has revealed their direct involvement in linking the basal body of primary cilia to the actin cytoskeleton in a specific structure of cilia adhesion (92). In fact, cilia have long been reported to coordinate integrin signaling (93), however, recent studies have found that integrins can also influence cilia formation. A gene-wide screen reported a key role for integrin  $\beta 1$  in primary ciliogenesis, Failler (94) silenced TLN1, LAMB4, FERMT2, VTN, ITGB1 and Rac1 in Hs578T cells and indicated that ablation with shRNAs significantly increased the number of ciliated cells, with ablation of TLN1, FERMT2 and VTN having the most profound effect, resulting in a population of over 30% of cells assembling cilia. Interestingly, the knockdown of ITGB1 or Rac1 produced similar results, confirming that integrins are associated with ciliogenesis. Previous studies reported that integrins and can activate the mitogen-activated protein kinase (MAPK) pathway and that MAPK stimulates F-actin rearrangement and cilia lengthening by phosphorylating IFT motor proteins (95). In addition, recent studies have found that integrins and primary cilia interact in bone metabolism. Geoghegan (92) found that estrogen withdrawal associated with postmenopausal osteoporosis leads to disruption of  $\alpha v\beta 3$  integrins and reduced actin contractility, which in turn leads to cilia prolongation resulting in Hh pathway activation and paracrine signaling in osteoclasts. Stam (96) investigated the role of integrin  $\alpha 1\beta 1$  in primary cilia and the Factin cytoskeleton in response to osteoarthritic mediators, measuring primary cilia length and F-actin peak numbers in isolated wild-type and itga1-null chondrocytes, and showed that integrin  $\alpha 1\beta 1$  is required to mediate cilia lengthening and the formation of F-actin peaks in response to hypotonic stress and IL-1. In addition, integrins and primary cilia, together with other mechanosensing systems such as polycystins and TAZ, may also generate multifaceted mechanosensing networks in the skeleton. This suggests that mechanosensing and mechanotransduction pathways in bone are fundamental to understanding bone physiology and therefore the specific molecular mechanisms of mechanosensors in the bone microenvironment still need to be further explored in subsequent work.

### 3 Conclusions and perspectives

Recent advances in understanding the role of primary cilia in bone metabolism are reviewed in this paper. Current research on primary cilia has focused on the basic function of primary cilia, ciliopathy and primary ciliary signaling pathways, and the crosstalk between primary cilia and other receptors and ciliary proteins is largely unexplored. Furthermore, it has been observed that primary cilia undergo elongation and curvature in response to mechanical forces. However, no studies have yet investigated the impact of cilia elongation, bending amplitude, and frequency on signaling pathways and the development of bone metabolic disorders.

In conclusion, this paper reaffirms the importance of primary cilia in bone development and emphasizes the necessity for continued research on primary cilia-associated proteins and signaling pathways. By targeting these aspects, more advanced treatments for ciliopathies can be developed, offering enhanced efficacy in the future.

### Author contributions

FFL: Conceptualization, Investigation, Writing – original draft. HL: Investigation, Writing – review & editing. YWM: Writing – review & editing. RZ: Writing – review & editing. WW: Conceptualization, Project administration, Funding acquisition.

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The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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