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The reciprocity of skeletal muscle and bone: an evolving view from mechanical coupling, secretory crosstalk to stem cell exchange

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Introduction: Muscle and bone constitute the two main parts of the musculoskeletal system and generate an intricately coordinated motion system. The crosstalk between muscle and bone has been under investigation, leading to revolutionary perspectives in recent years.

Method and results: In this review, the evolving concept of muscle-bone interaction from mechanical coupling, secretory crosstalk to stem cell exchange was explained in sequence. The theory of mechanical coupling stems from the observation that the development and maintenance of bone mass are largely dependent on muscle-derived mechanical loads, which was later proved by Wolff's law, Utah paradigm and Mechanostat hypothesis. Then bone and muscle are gradually recognized as endocrine organs, which can secrete various cytokines to modulate the tissue homeostasis and remodeling to each other. The latest view presented muscle-bone interaction in a more direct way: the resident mesenchymal stromal cell in the skeletal muscle, i.e., fibro-adipogenic progenitors (FAPs), could migrate to the bone injury site and contribute to bone regeneration. Emerging evidence even reveals the ectopic source of FAPs from tissue outside the musculoskeletal system, highlighting its dynamic property.

Conclusion: FAPs have been established as the critical cell connecting muscle and bone, which provides a new modality to study inter-tissue communication. A comprehensive and integrated perspective of muscle and bone will facilitate indepth research in the musculoskeletal system and promote novel therapeutic avenues in treating musculoskeletal disorders.

KEYWORDS

musculoskeletal system, inter-tissue communication, muscular diseases, bone regeneration, fibro-adipogenic progenitors

1 Introduction

The skeletal muscle and the bone constitute the two major components of the musculoskeletal system. They are in charge of the voluntary body movement, ranging from walking, jumping, facial expression, mastication to respiration. It has long been assumed that muscle and bone are a structural and mechanical integrity. The anatomical adjacency and mechano-transduction inside the tissues facilitate their growth and functional fulfillment. Furthermore, studies in the past 20 years highlighted the endocrine roles of muscle and bone, opening more avenues to the interaction between

the two tissues. It was not until recently that a direct connection had been established. The resident mesenchymal stromal cell in the skeletal muscle, i.e., fibro-adipogenic progenitors (FAPs), could migrate to the bone injury site and contribute to bone regeneration. This three-stage view of the muscle and bone relationship has greatly expanded the connotation of the musculoskeletal system and set up a new model for inter-organ communication.

2 Mechanical coupling

Because of the principal role in locomotion, the first noticed and studied aspect of muscle-bone interaction was mechanical. Anatomically, bones are "cornerstones" and muscles are more like "appendants": skeletal muscles connect bones and joints through tendons, while bones provide firm attachment sites in return (Frontera and Ochala, 2015). In contrast, muscles are "commanders" and bones are "executors" when completing physical activities. During physical activities, skeletal muscles are specified and actuated by the nervous system (Jayasinghe et al., 2022), followed by the asynchronous slide of actin and myosin filaments, which induces muscle contraction and thus generates a variety of motion-torque patterns at joints (Sweeney and Hammers, 2018; Ludvig et al., 2022), enabling the multidirectional movement of bones (Sylvester et al., 2021).

The muscle and the bone also interact through mechanical loads that culminate in remodeling (Goodman et al., 2015). The mechanical load exerted by bones on muscles originates from the elastic force accumulated by the deformation of bones under stress and has not been accurately measured. But mechanical loads imposed by muscles on bones are easier to comprehend and appreciate. During embryogenesis, muscle serves as a functional force generator early in development, exerting an increased mechanical load that can be translated into signals that combine with the genetic program of organogenesis on neighboring tissues as development proceeds (Felsenthal and Zelzer, 2017) and lasts throughout life. Most importantly, the development and maintenance of bone mass are largely dependent on musclederived mechanical loads (Goodman et al., 2015). As Wolff's law pointed out, bone size and geometry would change according to the strain applied. It was calculated that force produced in muscle contraction accounted for more than 70% of the bending moments imposed on the lower limb. Therefore, skeletal muscle was considered the primary source of mechanical loading of the bone, which became the central principle of the Utah paradigm. It was thus reasonable to infer that bone mass and mechanical properties would be allometric scaled to the peak muscle force. To depict a comprehensive prospect of the mechanical coupling, mechanostat hypothesis put forward a disuse-adapted-overload model where force over 3,000 microstrain would cause bone formation, while force under 5,00 microstrain would lead to bone resorption. In addition to generating growth stimulus on the bone, the skeletal muscle can also exacerbate the bone growth defect, as seen in the adolescent idiopathic scoliosis (Fadzan and Bettany-Saltikov, 2017). Strong paravertebral muscle on the convex side will reduce the mechanical loading on the spinal bone, thus promoting the bone growth. While weak paravertebral muscle on the concave side can increase the load and impede bone growth, resulting in aggravated curvature (Shao et al., 2023).

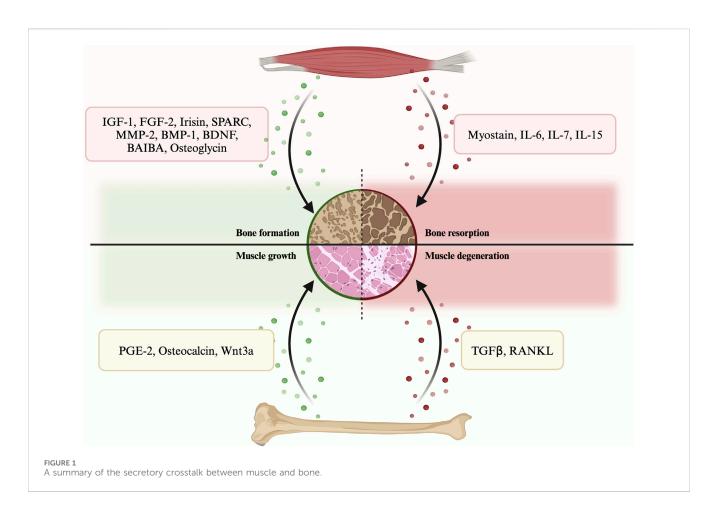
The Wolff's law, the Utah paradigm and the mechanostat hypothesis lay the foundation for the muscle-bone mechanical coupling theory. As the theory postulated, the growth in muscle strength should precede the growth in bone strength, which was proved by a longitudinal study examining the pubertal growth revealing that the peak of lean body mass preceded the peak of bone mineral content by an average of 0.51 years in girls and 0.36 years in boys (Rauch et al., 2004). Similarly, in a mice hindlimb suspension study, it was demonstrated that the cortical thickness of femur and tibia decreased 7 days later than the loss of gastrocnemius and quadricep muscle mass (Wang et al., 2020). These later observations verified the theoretic deduction.

3 Secretory crosstalk

Apart from generating and maintaining strength by mechanical coupling, the musculoskeletal system also influences the metabolism and function of other organs, including multi-organ insulin sensitivity (Bowden Davies et al., 2018), cardiac health (Wu et al., 2021) and risk of cancer (Jurdana, 2021). Since 1960s, researchers had hypothesized that skeletal muscle possess "humoral" factors, because electrical stimulation of dysfunctional muscles in patients with spinal cord injury induces many of the same physiological changes as in uninjured individuals (Kjaer et al., 1996; Mohr et al., 1997), implying that the musculoskeletal system can affect multiple organs in a non-neural conduction manner. The identification of myostatin confirms the existence of muscle secretory factor, which functions specifically as a negative regulator of skeletal muscle growth (McPherron et al., 1997). Since then, more than 650 myokines have joined the growing list of muscle secretory factors (Khan and Ghafoor, 2019). These muscle secretory factors that are produced, expressed and released by muscle fibers and exert either autocrine, paracrine or endocrine effects are classified as myokines (Severinsen and Pedersen, 2020), making skeletal muscle more than just a component in our locomotor system. When it comes to the other player in the musculoskeletal system, bone is generally thought as a torpid organ. But in recent years, studies have shown an active role of bone. It can function as an endocrine organ by producing many cytokines and proteins called osteokines to modulate glucose and energy metabolism as well as phosphate metabolism (Wang et al., 2021a).

3.1 Muscle secretory factors to bone

In terms of their effects on bone regeneration, myokines can be categorized into two types: bone formation factors and bone resorption factors (Figure 1; Supplementary Table S1). The former type includes various myokines such as insulin-like growth factor 1 (IGF-1), fibroblast growth factor 2 (FGF-2), irisin, secreted protein acidic and rich in cysteine (SPARC), matrix metalloproteinase 2 (MMP-2), bone morphogenetic protein 1 (BMP-1), brain-derived neurotrophic factor (BDNF), β -aminoisobutyric acid (BAIBA) etc. On the other hand,



myokines that promote bone resorption include myostatin (GDF-8), interleukins, etc. These myokines play a crucial role in bone remodeling and regeneration.

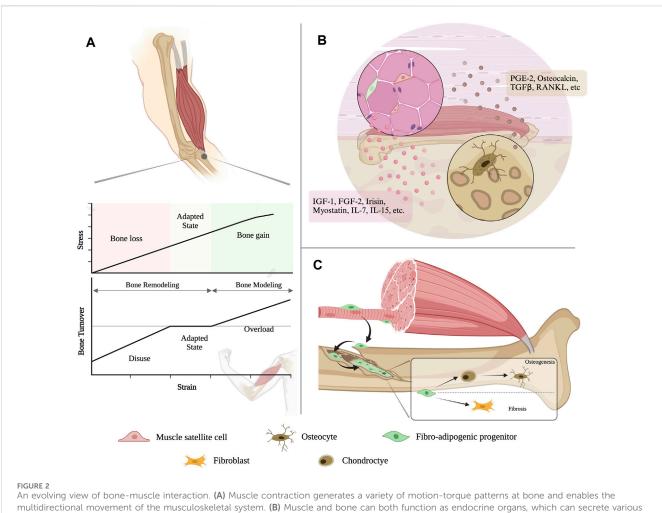
3.1.1 Bone formation myokines

The established role of IGF-1 and FGF-2 in bone physiology pertains to their ability to maintain proper bone matrix levels and bone mass by promoting osteoblast proliferation and differentiation (Novais et al., 2021; Chen et al., 2022; Mazziotti et al., 2022). *In vitro* studies demonstrated that myotubes secreted IGF-1 and FGF-2 (Clarke and Feeback, 1996; Henningsen et al., 2010), while *in vivo* experiments revealed high concentrations of IGF-1 in wound secretions from muscle flaps and increased local FGF-2 release following muscle injury (Vogt et al., 2005; D'Amore et al., 1994), in addition to elevated circulating FGF-2 levels (D'Amore et al., 1994). Furthermore, the localization of IGF-1 and FGF-2 at the muscle-periosteum interface suggested that muscle-derived IGF-1 and FGF-2 may play a role in bone regeneration through paracrine or endocrine actions (Hamrick et al., 2010).

Irisin, a hormone induced by exercise, is a fragmented product of fibronectin type III domain-containing protein 5(FNDC-5) and acts as a linkage between muscles and other tissues (Waseem et al., 2022). It has been demonstrated that Irisin expression is positively associated with bone mineral density and bone strength (Singhal et al., 2014; Palermo et al., 2015; Colaianni et al., 2019). More recently, irisin has also been shown to have anabolic effects on bone in animal models. Increased circulating irisin levels enhance osteogenesis in mice by inducing osteoblastogenesis and inhibiting osteoclastogenesis in bone cell lines (Zhang et al., 2017a). And the administration of irisin prevents both disuseinduced and peroxide-induced osteocyte apoptosis (Kim et al., 2018; Storlino et al., 2020). On the other hand, Irisin acts directly on osteoclast progenitors to increase differentiation and promote bone resorption, implying it may also serve as an important counterregulatory hormone that targets both osteoclasts and osteocytes (Estell et al., 2020).

SPARC is a glycoprotein that plays a crucial role in various physiological processes such as extracellular matrix remodeling, tissue repair, and collagen mineralization promotion (Zhou et al., 2018). During muscle damage and regeneration, SPARC is notably upregulated (Jørgensen et al., 2009). The SPARC null mice was characterized by delayed fibrocartilage mineralization that leads to the loss of bone mass (Wang et al., 2021b), suggesting that SPARC positively regulates bone regeneration and facilitates fibrocartilage mineralization, thereby aiding bone regeneration.

Matrix metalloproteinase 2 (MMP-2), a member of the extracellular matrix metalloproteinase family, is actively secreted by myotubes *in vitro*. The protein is essential for maintaining bone mineral density and strength, as well as skeletal development (Hardy and Fernandez-Patron, 2020). In MMP-2 null mice, bone loss and reduced bone mineral density occurred, meanwhile the lack of MMP-2 affected the later stages of fracture healing tissue remodeling (Lieu et al., 2011). However, dysregulated MMP-2 expression also led to various diseases, including developmental



multidirectional movement of the musculoskeletal system. (B) Muscle and bone can both function as endocrine organs, which can secrete various cytokines to modulate the tissue homeostasis and remodeling to each other. (C) The resident mesenchymal stromal cell in the skeletal muscle, i.e., fibro-adipogenic progenitors (FAPs), could migrate to the bone injury site and contribute to bone regeneration.

osteolysis and metastatic osteolysis (Li et al., 2021). Although MMP-2 may have a potential role in promoting bone formation as a myokine, further exploration is necessary to understand its specific function. Bone morphogenetic protein 1 (BMP-1), a zinc-dependent metalloproteinase, promoted bone mineralization (Kessler et al., 1996; Kudo, 2019). Expression of BMP-1 was detected in human myotubes cultured *in vitro* and in muscle tissues subjected to blast trauma (Hittel et al., 2009; Jackson et al., 2011), which could partially explain the ectopic ossification after blast trauma (Jackson et al., 2011). However, the role of BMP-1 as a myokine in normal bone regeneration requires further investigation.

Brain-derived neurotrophic factor (BDNF) is a member of the neurotrophin family of growth factors (Colucci-D'Amato et al., 2020). In addition to its role in neuron growth, neuronal development, and synaptic plasticity and function, BDNF is also expressed and released by skeletal muscle as a myokine that is capable of enhancing lipid oxidation in muscles via activation of AMPK (Pedersen et al., 2009; Antony and Li, 2020). Physical activities has a positive effect on the release of BDNF (Kim and Kim, 2018; Behrendt et al., 2021), whose specific receptor tropomyosin-related kinase B (TrkB) was shown to be expressed at high levels in the osteoblasts (Asaumi et al., 2000). During fracture healing, BDNF was found to positively modulate the expression and secretion of VEGF from osteoblasts via TrkB/ERK1/2 signaling pathway (Zhang et al., 2017b). And AKT signaling was found to be activated downstream, followed by the upregulation of integrin β 1, therefore stimulating osteoblasts migration (Zhang et al., 2020). A recent study reported BDNF was related to the maintenance of mitochondrial quality (Ahuja et al., 2022). Taking it into consideration that mitochondrial dysfunction is involved in several degenerative bone and joint diseases, especially osteoporosis and osteoarthritis (Lee et al., 2021; Sun et al., 2021), BDNF may play a more important role in bone remodeling than it does in our current knowledge.

 β -aminoisobutyric acid (BAIBA) origins from mitochondrial valine catabolism and is increased by physical activities (Kamei et al., 2020). BAIBA served as a bone-protective factor that prevents osteocyte cell death induced by reactive oxygen species (Kitase et al., 2018). Osteoglycin is a secreted protein found in skeletal muscle production and is recognized as a crucial anabolic factor produced by muscle-derived cells. Its secretion into the bloodstream promoted bone formation (Tanaka et al., 2012).

3.1.2 Bone resorption myokines

Myostatin (GDF-8), known as a retro-myokine, is secreted by muscle and acts as an inhibitor of muscle hypertrophy (McPherron et al., 1997; Kim et al., 2021). It is expressed in fracture healing tissue and suppresses the initial recruitment and proliferation of osteogenic progenitor cells in the healing tissue (Kellum et al., 2009). Myostatin could impede osteoblast differentiation and activate osteoclast maturation, resulting in compromised bone structure, bone density, and contractile properties (Bialek et al., 2014; Suh et al., 2020; Zhi et al., 2020; Liu et al., 2021; Omosule et al., 2022; Tang et al., 2022).

Interleukin (IL) families are pro-inflammatory mediators secreted by various cell types across the body. Several ILs including IL-6, IL-7, IL-15 were identified as myokines. During physical activities, circulating IL-6 level surged, and the majority of which was proven to originate from muscle (Chowdhury et al., 2020). IL-6 drived osteoclastogenesis and led to a bone-resorbing outcome (Udagawa et al., 1995; Chowdhury et al., 2020). IL-7 and IL-15 are also secreted by skeletal muscle and could induce bone resorption either directly by inducing osteoclastogenesis (Ogata et al., 1999; Kim et al., 2017), or indirectly by acting on effector cells consisting macrophages and NK cells (Feng et al., 2015; Kim et al., 2020). However, there is also an opposite perspective for IL-15, suggesting it is crucial for osteoblastic matrix formation and bone mineralization (Loro et al., 2017).

All myokines mentioned above are involved in regulating bone metabolism through endocrine pathways, which ultimately resulted in promotion or inhibition of bone remodeling.

3.2 Bone secretory factors to muscle

According to their impacts on muscle, osteokines can be categorized into muscle growth osteokines and muscle degeneration osteokines. Examples of the former include prostaglandin E2 (PGE-2), osteocalcin and Wnt3a, while transforming growth factor β (TGF β) and receptor activator of nuclear factor kappa-B ligand (RANKL) belong to the latter. Since the concept of bone as an endocrine organ is relatively new, studies on osteokines are less in-depth than those on myokines.

3.2.1 Muscle growth oseokines

The osteokine PGE-2 is secreted at high levels by osteocytes, which has four types of receptors executing different functional roles. Stimulating the receptor EP4 significantly enhanced myoblast proliferation (Mo et al., 2015). While down-regulating the receptor EP2 promoted the fusion of human muscle progenitors in vitro and improved their transplantation capability (Sakai-Takemura et al., 2020). Osteocalcin is mainly produced by mature osteoblasts, but also by osteocytes (Bonewald, 2019). And pervious study has shown osteocalcin favored uptake and catabolism of nutrients in muscle and was necessary for adaptation to exercise (Mera et al., 2016). Wnt signaling is involved in the control of the myogenic program and the differentiation of satellite cells, the muscle resident stem cells (Fujimaki et al., 2014). A couple of Wnt signaling pathwayrelated factors were reported to support myogenesis and muscle function. Wnt1 is highly expressed in osteocytes and Wnt3a is produced by osteocytes in response to shear stress. They could both induce myogenesis but Wnt3a seems to be more potent than Wnt1 (Huang et al., 2017).

3.2.2 Muscle degeneration osteokines

TGF β is mainly produced by bone-forming osteoblasts (Bonewald, 2019). In osteolytic bone metastases cancer models, TGF β were released from the bone surface, and resulting in elevated oxidization of skeletal muscle proteins that contributed to muscle weakness (Waning et al., 2015). RANKL was first identified as a product of immune cells, but has since been shown to be produced by osteocytes to activate osteoclasts (Xiong et al., 2011). Injections of RANKL inhibitor to mice significantly increased the force of dystrophic EDL muscle (Dufresne et al., 2018), suggesting RANKL is a promising target to control muscle remodeling.

4 Stem cell exchange

As the researches delve deeper, it came out that muscle-bone interaction went beyond mechanical and paracrine: stem cells from the skeletal muscle could actually migrate to the underlying bone and directly contribute to bone regeneration. On the other hand, there is currently no evidence of stem cell contribution from the bone to the skeletal muscle, which prompts us to view muscle as a more vigorous part in this pair.

4.1 Stem cell contribution from muscle to bone: traditional view and new perspective of skeletal stem cells

Bone maintains its structural integrity and functionality through a process that relies heavily on the activation of skeletal stem/ progenitor cells (SSPCs) (Jeffery et al., 2022). These specialized cells are capable of both self-renewal and multilineage differentiation into bone, cartilage, and stroma (Chan et al., 2018). Multiple sources of skeletal stem/progenitor cells (SSPCs) have been identified for bone repair, including bone marrow, growth plate, periosteum. Each population displays unique lineage capabilities and is involved in bone repair in varying degrees (Serowoky et al., 2020). SSPCs derived from bone marrow are primarily identified by the Leptin receptor (LepR) and are responsible for generating new osteoblasts in adult bone marrow, which can form ossicles supporting hematopoiesis in vivo (Zhou et al., 2014; Matsushita et al., 2020). Growth plate SSPCs express parathyroid hormone-related protein (PTHrP) and maintain the resting zone while providing a source of chondrocytes during bone repair (Mizuhashi et al., 2018). In the periosteum, Gil1+ SSPCs contribute to and are required for the growth and repair of skull bones (Zhao et al., 2015), whereas Prrx1+ cells broadly mark periosteal SSPCs in the limbs and elsewhere (Duchamp de Lageneste et al., 2018), re-establishing the pool of bone progenitor cells after injury. Additionally, Ctsk is a conservative marker of periosteal SSPCs, with Ctsk + cells contributing primarily to osteoblasts in cortical bone (Debnath et al., 2018).

In reconstructive surgeries, fractured bone with a muscular flap always heals better. Consequently, it has long been assumed that the

skeletal muscle tissue plays a protective role in bone fracture repair. But little was known about the cellular and molecular process. The Colnot team endeavored to explore the involvement of muscleoriginated stem cells in bone fracture healing (Abou-Khalil et al., 2015). At first, they investigated the contribution of muscle satellite cells (MuSCs), with lineage-tracing transgenic mice and transplantation experiments. However, they found rare MuSCs could form chondrocyte and the MuSCs contribution in the bone fracture callus was not substantial. In the end, they identified the critical role of BMP-2, which was secreted by MuSCs and promoted bone fracture healing. This was the first evidence that cells derived from muscle contributed to bone regeneration, though in a paracrine way. For further study, the Colnot team investigated another stem cell resident in the skeletal muscle-fibroadipogenic progenitors (FAPs), which are the mesenchymal stem cells closely coordinated with MuSCs in the orchestration of skeletal muscle regeneration. The authors transplanted EDL muscle grafts from Prrx1Cre; RosamTmG mice donors into wild-type hosts. They observed that progenitors originated from skeletal muscle gave rise to chondrocytes and osteoblasts for bone repair. Further scRNAseq analyses identified these progenitors as FAPs within skeletal muscle but expressing common markers with SSPCs. They also built a polytrauma mouse model to investigate the role of FAPs in musculoskeletal trauma and found that FAPs failed to undergo chondrogenesis under polytrauma, resulting in the non-union phenotype of fracture. Their sophisticated work verified the direct contribution of FAPs to bone repair and highlight the role of injury in mediating FAPs' behavior. FAPs has thus been confirmed as a new source of SSPCs.

4.2 Stem cell contribution from bone to muscle: an unanswered question

Although the myogenic capacity of SSPCs has not been extensively studied, the simultaneous occurrence of bone fragility and muscle weakness in osteogenesis imperfecta (OI) patients suggests a potential link between these two tissues (Phillips and Jeong, 2018). It has been reported that the mechanotransduction and functionality of the muscle-bone unit was repaired in OI (Veilleux et al., 2015). Whether there could be defects in the direct intercellular communication between bone and muscle stem cells still warrants further investigation. Exploring the underlying mechanisms responsible for the decline of both bone and muscle could lead to the development of novel physiotherapeutic and pharmacological interventions for OI and other musculoskeletal disorders.

5 Discussion

5.1 FAPs-the critical cell connecting muscle and bone

Ever since its verified role as an essential progenitor cell in skeletal muscle in 2010, the FAP cell has been under energetic investigation. And the discovery that FAPs can also serve as SSPCs enhanced our comprehension of the musculoskeletal system. Evidence from studies of FAPs in skeletal muscle homeostasis maintenance and regeneration may provide inspiration for the roles of FAPs and other SSPCs in the bone, and *vice versa*.

FAPs are quiescent mesenchymal stromal cells with multipotency to differentiate into all the mesenchymal lineages, depending on the context of tissue damage (Contreras et al., 2021). The systemic protease, hepatocyte growth factor activator, which was induced by tissue injury, could prime FAPs to transitions from quiescence to G alert state (Rodgers et al., 2017). TGF_β signaling remained the most studied signaling pathway regulating FAPs' fate and behavior. Ligands of the TGF^β super family, including TGF^β, myostatin and bone morphogenetic proteins (BMPs) could induce cell proliferation, myofibroblast differentiation and extracellular matrix (ECM) deposition (Juban et al., 2018). Wnt/β-catenin signaling and platelet-derived growth factor signaling could also activate FAPs and induce the expression of several ECM genes (Akhmetshina et al., 2012; Contreras et al., 2019). On the other hand, tumor necrosis factor-alpha (TNFa) induced the apoptosis of FAPs (Lemos et al., 2015). As for the adipogenic differentiation of FAPs, the cellular communication network (CCN) family members and dexamethasone could play a stimulative role, while IL4 and histone deacetylase inhibitors could play a suppressive role (Mozzetta et al., 2013; Dong et al., 2014; Hu et al., 2019). In addition, TGFB signaling can inhibit the adipogenic differentiation of FAPs (Contreras et al., 2021). It has also been reported that FAPs were the main cell responsible for intramuscular ossification. Both BMP2 and BMP9 could promote FAP osteogenic differentiation (Shore, 2011). Moreover, targeted expression of an activin receptor in FAPs could recapitulate full spectrum of heterotopic ossification in muscle, suggesting the key role of activin signaling in regulating FAP osteogenic differentiation (Lees-Shepard et al., 2018). When applied at a low concentration, TNFa could promote FAP osteogenic differentiation in the context of bone fracture healing (Glass et al., 2011).

Recent studies revealed that FAPs have different embryonic origins, similar to the muscle they reside in (Sefton and Kardon, 2019). While limb muscles arise from somites, craniofacial muscles originate from branchial arches. Craniofacial muscles exhibit delayed myofiber reconstitution and prolonged fibrosis during repair, in contrast to somite-derived limb muscles, where FAPs serve as the key mediator of muscle fibrosis (Cheng et al., 2021; Cheng et al., 2022). It has been demonstrated in studies of skin and mucosa that cells derived from neural crest and mesoderm have distinct fibrogenic potential (Rinkevich et al., 2015; Pratsinis et al., 2019). Thus the branchiomeric FAPs in the craniofacial muscle could be the main culprit for the impaired muscle regeneration.

It is also observed in the bone that cells from different origins demonstrated distinct behaviors. But the scenario is more complicated. When the bone is injured, the neural crest-derived mandible fully regenerated with neural crest-derived SSPCs, and the mesoderm-derived tibia heals with mesoderm-derived SSPCs (Leucht et al., 2008). Further transplantation experiment revealed that the SSPCs from different lineages were functionally interchangeable only when the host and the donor had the same hox code (Leucht et al., 2008). And this offers another possible explanation for the muscle: the mesoderm-derived myofiber and neural crest-derived FAPs are mutually exclusive, leading to impaired regeneration process. However, there is currently no

10.3389/fphys.2024.1349253

research exploring the osteogenic potential of FAPs from different embryonic origins, neither *in vitro* nor in the scenario of bone regeneration. It is interesting to figure out whether the beneficial role of FAPs in bone regeneration is dependent on their embryonic origin. It is also of great clinical importance to guide the flap transfer surgery.

5.2 Therapeutic possibilities involving FAPs for treating musculoskeletal diseases

The perspective of FAPs as an active stem cell participant in neighboring tissue regeneration offers more possibilities for regenerative medicine and orthobiologics for musculoskeletal disorders. Platelet-rich plasma (PRP), bone marrow and adipose tissue are the most commonly used orthobiologics (Fang and Vangsness, 2021). PRP and bone marrow aspirations have the advantage of being minimally manipulated but have a relatively low concentration of stem cell components, which can largely aid tissue regeneration (Fang and Vangsness, 2021). Since mesenchymal stem cells (MSCs) are the most prospective stem cell for regenerative medicine (Lemos and Duffield, 2018), adipose tissue and umbilical cord, which are rich in MSCs, are believed to be promising. Adipose tissue can yield the highest number of MSCs per milliliter of tissue while Whartons jelly tissue can provide the highest concentration of MSCs (Shapiro et al., 2022). Nevertheless, the use of adipose and umbilical cord tissue for treating musculoskeletal diseases is either off-shelf or still under investigation (Shapiro et al., 2023). The proportion of FAPs in skeletal muscle tissue is comparable to that of MSCs in the adipose tissue (Giordani et al., 2019), making them another possible choice for stem cell therapies, though the efficiency and security issues are to be scrutinized. Meanwhile, the secretory factors of FAPs also stand a chance of turning into potential therapeutics for promoting bone and muscle regeneration and warrants further investigation. Furthermore, since several immunomodulation agents hold greate promise in several preclinical studies to treat muscle and bone injuries, the immunomodulatory role of FAPs might also facilitate musculoskeletal tissue regeneration (Maruyama et al., 2020; Luo et al., 2022; Duda et al., 2023; Qin et al., 2023).

Potential therapeutic avenues involving FAPs for musculoskeletal disorders have been proposed by a few studies. Nilotinib, a clinically approved tyrosine kinase inhibitor, could exert antifibrotic effects on skeletal muscle by restoring FAP apoptosis (Lemos et al., 2015). Another member of the tyrosine kinase inhibitor family, Imatinib, could improve bone regeneration by decreasing the persistent callus fibrosis, which was mainly caused by FAP, in the context of bone-muscle polytrauma (Julien et al., 2021). In addition, the observation that the endothelin receptor type B (EDNRB) was highly expressed in fibrotic FAPs uncovered the critical role of endothelin in the altered crosstalk between muscle cells and FAPs (Bensalah et al., 2022). The application of Bosentan, an antagonist against EDNR could counteract fibrosis and enhance skeletal muscle regeneration (Bensalah et al., 2022).

The view of the muscle-bone dialogue has evolved far from mechanical coupling and secretory crosstalk (Figure 2). After FAPs had been identified as the substantial cell contributing to bone fracture healing, cellular exchanges between another two juxtaposed tissues started to be uncovered. Subcutaneous adipose tissue can provide regenerative cells responsible for skeletal muscle regeneration (Sastourné-Arrey et al., 2023). This not only offers more solid evidence to support the existing perspective that bone, muscle and fat work synergistically as a functional unit, but also renders a modality for studying intertissue communication in general and brings revelation for future investigations into the cellular crosstalk in multi-organ syndromes, such as cancer cachexia.

Author contributions

HS: Conceptualization, Investigation, Resources, Writing-original draft, Writing-reviewing and editing. JD: Investigation, Methodology, Writing-original draft. BS: Funding acquisition, Writing-review and editing, Resources, Validation. XC: Conceptualization, Investigation, Writing-original graft, Writing-review and editing, Funding acquisition.

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Conflict of interest

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fphys.2024.1349253/ full#supplementary-material

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