



The role of macrophages in fracture healing: a narrative review of the recent updates and therapeutic perspectives

Bianca Braga Frade^{1,2}, Rhayra Braga Dias^{1,3}, Sara Gemini Piperni⁴, Danielle Cabral Bonfim¹

¹Laboratory of Stem Cells and Bone Regeneration, Institute of Biomedical Sciences, Federal University of Rio de Janeiro, Rio de Janeiro, Brazil;

²Postgraduation Program in Biological Sciences–Biophysics, Institute of Biophysics Carlos Chagas Filho, Federal University of Rio de Janeiro, Rio de Janeiro, Brazil;

³Postgraduation Program in Morphological Sciences, Institute of Biomedical Sciences, Federal University of Rio de Janeiro, Rio de Janeiro, Brazil;

⁴Laboratory of Biotechnology, Bioengineering and Nanostructured Biomaterials, Institute of Biomedical Sciences, Federal University of Rio de Janeiro, Rio de Janeiro, Brazil

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Correspondence to: Danielle Cabral Bonfim. Instituto de Ciências Biomédicas, Av. Carlos Chagas Filho, 373-sala F2-030, Cidade Universitária, Ilha do Fundão, 21941-590 Rio de Janeiro, Brazil. Email: bonfimdc@icb.ufrj.br.

Objective: This review addresses the latest advances in research on the role of macrophages in fracture healing, exploring their relationship with failures in bone consolidation and the perspectives for the development of advanced and innovative therapies to promote bone regeneration.

Background: The bone can fully restore its form and function after a fracture. However, the regenerative process of fracture healing is complex and is influenced by several factors, including macrophage activity. These cells have been found in the fracture site at all stages of bone regeneration, and their general depletion or the knockdown of receptors that mediate their differentiation, polarization, and/or function result in impaired fracture healing.

Methods: The literature search was carried out in the PubMed database, using combinations of the keywords “macrophage”, “fracture healing”, “bone regeneration”, and “bone repair”. Articles published within the last years (2017–2022) reporting evidence from *in vivo* long bone fracture healing experiments were included.

Conclusions: Studies published in the last five years on the role of macrophages in fracture healing strengthened the idea that what appears to be essential when it comes to a successful consolidation is the right balance between the M1/M2 populations, which have different but complementary roles in the process. These findings opened promising new avenues for the development of several macrophage-targeted therapies, including the administration of molecules and/or biomaterials intended to regulate macrophage differentiation and polarization, the local transplantation of macrophage precursors, and the use of exosomes to deliver signaling molecules that influence macrophage activities. However, more research is still warranted to better understand the diversity of macrophage phenotypes and their specific roles in each step of fracture healing and to decipher the key molecular mechanisms involved in the *in vivo* crosstalk between macrophages and other microenvironmental cell types, such as endothelial and skeletal stem/progenitor cells.

Keywords: Fracture healing; macrophage; bone repair; bone regeneration

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Introduction

Bone has a significant regenerative capacity after fracture, being able to completely restore its pre-injury form and function. However, consolidation failures are frequent in patients with long bone fractures resulting from high-energy trauma (traffic accidents, falls from heights, and gunshots) and with fragility of the bone mass, resulting in an inability to heal the fracture (nonunion) (1-4). Treatment options for this problem usually involve surgical removal of the nonunion fibrotic tissue, followed by replacement of the fixation device and grafting at the fracture site with autologous and/or allogeneic trabecular bone containing red marrow (3,5). Nevertheless, if the cause of the nonunion is more a biological, rather than a mechanical impairment, this strategy does not actually fix the source of the problem, and not surprisingly, re-failures are common. The burden is significant for patients who undergo multiple invasive surgeries, prolonged time of chronic pain, physical incapacity, and psychosocial disability. Furthermore, nonunion treatment is expensive, requires permanent medical assistance, multiple hospitalizations, and the use of many orthopedic devices (6,7). Thus, nonunion is already a clinical challenge, and considering the lifestyle in urban centers and the increasing populational aging rates (8), this scenario is expected to worsen, requiring the development of innovative therapeutic strategies to promote bone repair. To this end, a better understanding of the mechanisms that determine the success of fracture healing is a fundamental step.

Long bone fractures treated with fixation strategies that allow a certain degree of movement between the cortical bone ends, such as the gold-standard intramedullary nailing (5,9), heal through callus formation, a process that encompasses three major consecutive phases: the inflammatory reaction, the repair phase, and the final bone remodeling (10-12) (*Figure 1*). Following the breakage of bone and the rupture of blood vessels, a coagulation cascade is triggered, forming a hematoma that fills the space between bone fragments and connects the bone marrow, the periosteum, the endosteum, and the surrounding muscle. Within this unique microenvironment, immune cells, including neutrophils, macrophages, and platelets, begin an inflammatory reaction with the secretion of cytokines, chemokines and growth factors, such as interleukin-1 β (IL-1 β), IL-6, tumor necrosis factor- α (TNF- α), monocyte chemoattractant protein-1 (MCP-1), macrophage colony-stimulating factor (M-CSF), platelet-derived growth factor (PDGF), and vascular endothelial growth factor

(VEGF) (6,10). While this response is amplified within the hematoma, the surrounding periosteum expands, and a new chemotaxis axis mediated by fibroblast growth factors (FGF), bone morphogenetic proteins (BMP), Indian hedgehogs (Ihh), Wnts, placental growth factors (PIGF), and VEGFs activate and recruit endothelial and skeletal stem/progenitor cells, which migrate to specific areas of the fracture to promote revascularization and bone repair (13,14). In mice, at day 7, osteochondroprogenitors expressing runt-related transcription factor 2 (Runx-2) and SRY-Box Transcription Factor 9 (Sox-9) locate at the periphery of the fracture line adjacent to the periosteum (15). On day 14, committed osteoprogenitors expressing Runx2 and Osterix (Osx) are found on both sides of the gap, where new bone formation occurs through intramembranous ossification (15). On the other hand, Sox9⁺ chondroprogenitors are located exclusively in the central area of the fracture, where they generate chondrocytes and establish a soft callus composed of hyaline cartilage (6,12,13,15). Later, terminal differentiation of chondrocytes into the hypertrophic state stimulates matrix calcification and vascular invasion. Following the path of type H vessels formed by endothelial cells with high expression of Endomucin and CD31 (15), Runx2⁺ Osx⁺ osteoprogenitors reach the area of the calcified cartilage matrix and start depositing immature trabecular bone. This hard callus then bridges the old cortical bone fragments, and finally the bone segment is remodeled into the mature lamellar osseous structure by the activity of osteoclasts and osteoblasts, restoring its original shape and mechanical properties (12,16).

Although this general sequence of fracture healing is well described (6,12), several knowledge gaps still exist in relation to cellular interactions and molecular signaling leading to bone consolidation or nonunion. However, compelling evidence indicates that the initial inflammatory phase is the most critical for the outcome (17-19). At this stage, many types of immune cells are found within the early fracture hematoma, including lymphocytes, which are cells of the adaptive response (20-25). However, the role of macrophages in fracture healing has become a central topic in osteoimmunology research because: (I) macrophages are the main drivers of inflammatory responses in the general process of wound healing, being the source of several chemotactic molecules that activate and recruit both specialized and progenitor/stem cells to engage in tissue repair (26,27); (II) during homeostasis, they are found on the endosteal surface of bones, covering osteoblasts in a canopy-like structure (28) and supporting bone formation in

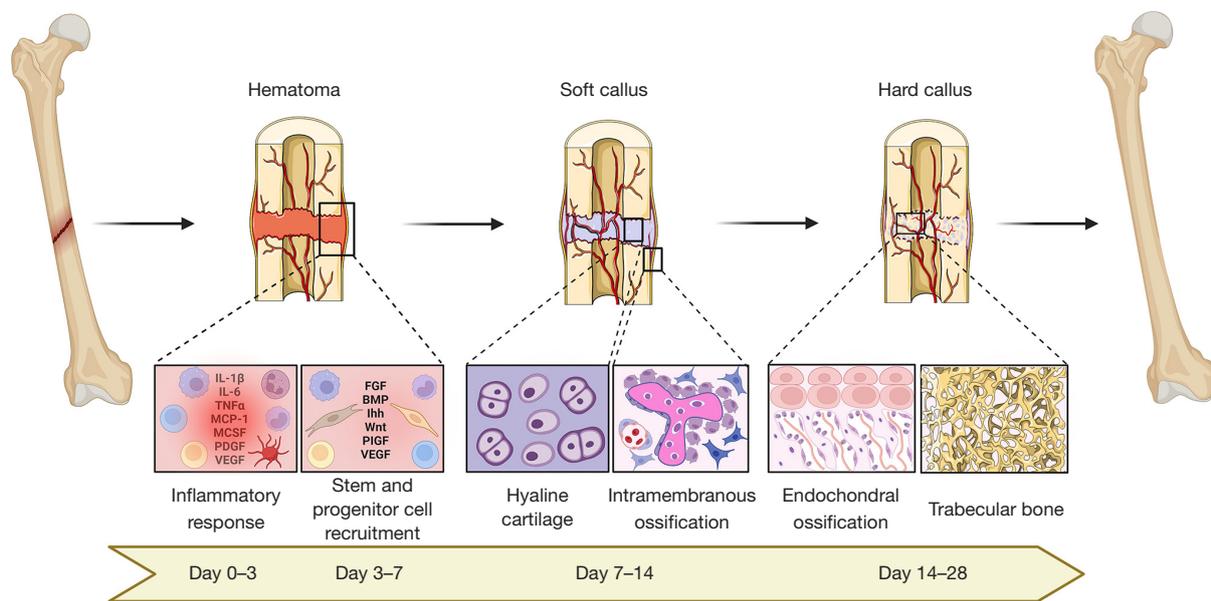


Figure 1 Overview of long bone fracture healing stages. After injury, the rupture of blood vessels forms a hematoma that fills the space between the bone fragments. Immune cells including neutrophils, macrophages, and platelets start an inflammatory reaction with the secretion of cytokines, chemokines, and growth factors. While this response is established, the periosteum expands and a new chemotaxis axis mediated by diverse signaling factors recruit endothelial and skeletal stem/progenitor cells. The process then advances to the establishment of a soft callus, whose central area is mainly composed of hyaline cartilage. At the callus border, the formation of new bone occurs through the differentiation of skeletal progenitors directly into osteoblasts (intramembranous ossification). Later, terminal differentiation of chondrocytes into the hypertrophic state stimulates matrix calcification, invasion of nascent blood vessels along with skeletal progenitors, who deposit new bone resembling the endochondral ossification process. The hard callus, composed of trabecular bone, then bridges the old cortical fragments. Finally, the bone is remodeled into its original shape. The timescale depicted in the scheme represents the approximate extension of each event in the healing of mice fractures. Created with BioRender.com. IL, interleukin; TNF, tumor necrosis factor; MCP-1, monocyte chemoattractant protein-1; M-CSF, macrophage colony-stimulating factor; PDGF, platelet-derived growth factor; VEGF, vascular endothelial growth factor; FGF, fibroblast growth factors; Ihh, Indian hedgehogs; PIGF, placental growth factors.

the context of the physiological process of bone remodeling (29-31); (III) macrophages were found to secrete inductive signals that stimulate the differentiation of skeletal (32-35) and endothelial progenitor cells (36,37); (IV) in the context of the fracture healing cascade, macrophages were found within the callus at all stages, in close association with areas of bone formation (21,34,38,39); and (V) their general depletion or the knockdown of receptors that mediate their differentiation, polarization, and/or function resulted in delayed and/or failed bone consolidation in murine fracture healing models (28,34,39-42). Collectively, these findings suggested that the contributions of macrophages to bone repair go far beyond the sole modulation of inflammation, placing these cells as central instructors of the whole healing cascade (43-45). Consequently, modulation of macrophage activity has emerged as a potentially important therapeutic

strategy to stimulate or accelerate bone repair. Nevertheless, as macrophages are a heterogeneous cell population which can adopt diverse phenotypes and functional profiles (27,46,47), further studies are still needed in order to better depict the specific contributions of the different macrophage subtypes in each stage/event of fracture healing. Therefore, the objective of this review is to address the most recent advances in research on the role of macrophages in the healing of long bone fractures, exploring their relationship with failures in bone consolidation, and the latest perspectives for the development of innovative treatments for delayed and/or failed bone consolidation, based on the modulation of macrophage activity. We present the following article in accordance with the Narrative Review reporting checklist (available at <https://sci.amegroups.com/article/view/10.21037/sci-2022-038/rc>).

Methods

The literature search was carried out in the PubMed database, using combinations of the keywords “macrophage”, “fracture healing”, “bone regeneration”, and “bone repair”. Articles published within the last years (2017–2022) reporting evidence resulting from *in vivo* long bone fracture healing experiments were included.

General concepts of macrophages and their role in fracture healing

Macrophages are a phenotypic and functionally diverse set of mononucleated phagocytic cells present in all body tissues that are involved in many physiological and pathological processes, including development, homeostasis, immunological defenses, and repair (48). This macrophage heterogeneity stems both from the origin of the macrophage population (49), as well as from the overall input signaling deriving from the tissue microenvironment at a given context, which makes macrophages polarize between two opposing spectrums: M1, which steers pro-inflammatory responses; and M2, which promotes resolution of inflammation and tissue repair (48). The M1 macrophages are also described as “classically activated macrophages” and phenotypically express high levels of the major histocompatibility complex class II (MHC II), F4/80, CD11b, and CD68 and secrete pro-inflammatory cytokines such as TNF- α , IL-1, and IL-6 (48). On the contrary, M2 macrophages, also known as “alternatively activated macrophages”, express the surface markers F4/80, CD162, and CD206 and secrete cytokines such as arginase-1, IL-4, IL-10, and transforming growth factor beta (TGF- β). These are further subdivided into M2a–M2d, with each subpopulation expressing a specific repertoire of cytokines (48) (Figure 2).

Regarding origin, studies have shown that the population of macrophages that reside within specific tissues [which are collectively called tissue-resident macrophages; TRM (50)] derives from yolk-sac erythromyeloid progenitors and can self-renew, maintaining their pools throughout life (51). Functionally, TRMs were shown to maintain tissue homeostasis and facilitate tissue repair by resolving inflammation. On the other hand, in adult life, macrophages can also originate from blood-circulating monocytes, which derive from bone marrow myeloid progenitors (49). In bones, TRMs are called osteomacs (28,52) (Figure 2). These are found in the endosteum, close to bone lining cells (28,53). On the other hand, recruited macrophages

are found to be distributed among the bone marrow and increase in number when a fracture occurs (53).

Studies published in the last five years on the role of macrophages in fracture healing strengthened the idea that what appears to be essential when it comes to a successful consolidation is the right balance between the M1/M2 populations, which have different but complementary roles in the process. The general paradigm of macrophage functional polarization in bone regeneration indicates that M1 macrophages contribute to the establishment of the initial acute inflammatory response within the hematoma, increasing its number early after fracture and secreting pro-inflammatory factors, such as IL-1 α , IL-1 β , IL-6, MCP-1, granulocyte-colony stimulating factor (G-CSF), IL-12, IL-23, TNF- α , and inducible nitric oxide synthase (iNOS), that will recruit additional immune cells to amplify the response (54). Additionally, these cells clear debris from dying and necrotic cells and produce growth factors, such as VEGF and PDGF, that activate endothelial progenitors that will start the angiogenic process. In fact, McCauley *et al.* (54) showed that F4/80⁺/MHC II⁺/CD86⁺ M1 macrophages were elevated early after injury in fractured femurs of mice and that this increase consistently progressed over time until day 7, which in mice is marked by the transition from the inflammatory to the tissue neoformation phase. Furthermore, the authors showed that while the M1 macrophage pool increased, M2 macrophages identified by the phenotype F4/80⁺ MHCII⁻/CD86⁻/CD11b⁻ decreased significantly, and this was accompanied by the surge of two additional populations of F4/80⁺ macrophages with intermediate phenotypes (F4/80⁺/MHCII⁺/CD86⁻ and F4/80⁺/MHCII⁻/CD86⁺), which were defined as macrophages transitioning from M2 to M1 (54), thus pointing to the critical role of M1 macrophages for the initiation of the healing cascade.

The importance of M1 macrophages in fracture healing was further highlighted by studies showing that the deletion of this population in mice fracture models resulted in impairment of bone consolidation (55–57). Using the saporin-conjugated Mac-1 antibody to deplete CD11b⁺ macrophages, Hozain & Cottrell observed a reduction in F4/80⁺/CD11b⁺ cells in the callus and a decrease in cartilage and bone volume, trabecular bone volume and thickness, and cortical area in the fractured limb (58). Similarly, Wasnik *et al.* showed that the administration of 1,25-Dihydroxyvitamin D [1,25(OH)₂D] at the fracture site during the early pro-inflammatory stage suppressed M1 and stimulated M2 macrophage differentiation, reducing

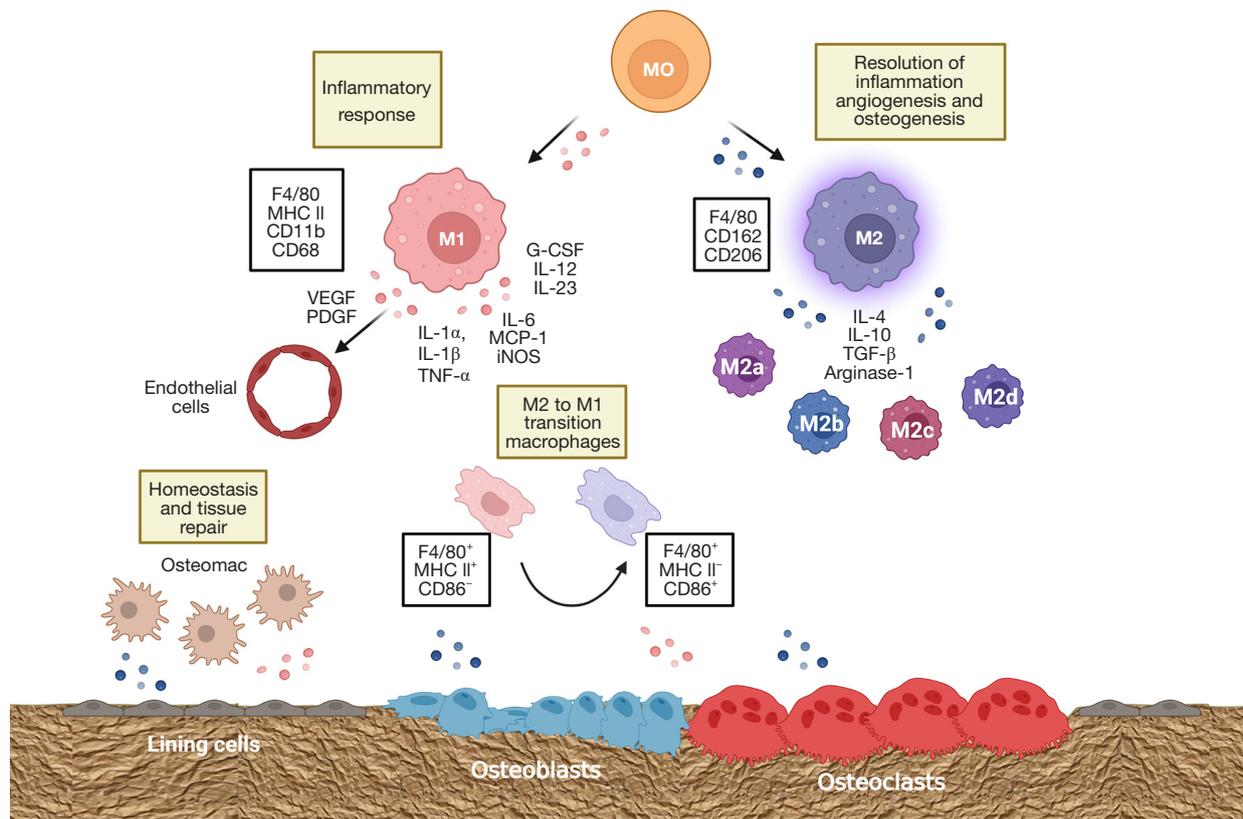


Figure 2 Macrophage diversity and their role in fracture healing. Macrophages are a phenotypic and functionally diverse set of cells whose heterogeneity stems from their origin and input signaling from the tissue microenvironment. In bones, the tissue-resident macrophages are the osteomacs, which originate from yolk-sac erythromyeloid progenitors. During homeostasis, osteomacs contribute to bone maintenance by stimulating the osteoblastic function. In fracture healing, they act to resolve inflammation. Within the recruited macrophage pool, M1 or M2 phenotypes possess different but complementary roles in the fracture healing process. F4/80⁺/MHC II⁺/CD86⁺ M1 macrophages are elevated until day 7 after injury, recruiting immune cells by the secretion of inflammatory factors, such as IL-1 α , IL-1 β , IL-6, MCP-1, G-CSF, IL-12, IL-23, TNF- α , and iNOS and activating endothelial progenitors by VEGF and PDGF production. On the contrary, M2 macrophages, also known as “alternatively activated macrophages”, are characterized by the expression of F4/80, CD162, and CD206 and secrete cytokines such as arginase-1, IL-4, IL-10, and TGF- β to promote angiogenesis and osteogenesis when the inflammatory reactions subside. These are further subdivided into M2a-M2d, with each subpopulation expressing a specific repertoire of cytokines. During the inflammatory response, M2 macrophages can transition to the M1 state, to contribute for the initiation of the healing process. Created with BioRender.com. MHC II, major histocompatibility complex class II; IL, interleukin; G-CSF, granulocyte-colony stimulating factor; MCP-1, monocyte chemoattractant protein-1; TNF, tumor necrosis factor; iNOS, inducible nitric oxide synthase; VEGF, vascular endothelial growth factor; PDGF, platelet-derived growth factor; TGF, transforming growth factor.

callus size by approximately 40% on day 14 after fracture and the final bone union rate by 65%. Importantly, these effects were not observed when 1,25(OH)₂D treatment was performed after the inflammatory phase (55), which agrees with the notion that the role of M1 macrophages is preponderant at the inflammatory stage of fracture healing, which then subsides with the transition to the tissue neof ormation stage, when M2 macrophages dominate and

are believed to instruct the mechanisms of angiogenesis and osteogenesis (56). In this regard, using the unicortical drill-hole fracture model in mice, Olmsted-Davis *et al.* identified a transient population of macrophages that express the beta-3 adrenergic receptor (ADRB3) within the callus, which peaked approximately 4 days after the injury. The authors verified, through immunophenotypic analysis, that these cells were more polarized toward the M2 spectrum, acting

to regulate oxygen tension and promote angiogenesis (59).

The relevance of the timely progression of the pro- to anti-inflammatory microenvironment for fracture healing was reported in two studies by Zhao *et al.* (41,42). In the first, the authors showed that the deficiency in the function of macrophage G-protein coupled receptor interacting protein 1 (GIT1) in a tibial monocortical fracture model resulted in persistent and enhanced M1 macrophage infiltration, exacerbation of IL-1 β production, and impairment of bone formation through intramembranous ossification (42). On the other hand, using the same mice fracture model, it was verified that the loss of function of macrophage scavenger receptor 1 (MSR1) in macrophages was correlated with a significant increase in M1 macrophages (F4/80⁺ iNOS⁺) and a marked decrease in the M2 pool (F4/80⁺ CD206⁺) on day 7 after fracture, also resulting in impaired intramembranous ossification (41).

The role of osteomacs, the TRMs of bones, was also investigated. Using the CD169-diphtheria toxin receptor (DTR) knock-in model, Batoon *et al.* evaluated the effects of osteomac depletion on the regeneration of bone injuries produced by the drill-hole method (which heals by intramembranous ossification) and by complete femoral osteotomy (which heals by endochondral ossification) (60). In vehicle-treated CD169-DTR mice, F4/80⁺ macrophages were abundantly seen in the adjacent bone marrow area, dispersed throughout the site of the injury associated with the granulation tissue, and accumulated in the peripheral injury zone. However, in CD169-DTR mice treated with diphtheria toxin and subjected to the drill hole lesion, F4/80⁺ macrophages were greatly reduced in the adjacent bone marrow, rare within the granulation tissue, and present, but with a reduced frequency in the peripheral injury zone. This observation was correlated with a reduction in bone formation and an increase in fibrotic tissue at the fracture site. In the endochondral fracture healing model, a general decrease in callus size was observed, whose magnitude was correlated with the number of residual F4/80⁺ macrophages that remained in the fracture site, suggesting that osteomacs are also important for the success of bone consolidation (60).

Relationship of macrophages and fracture impairment in the context of bone diseases

The disturbed frequency of macrophages and/or of their activity has been reported in various skeletal disease conditions and has been further investigated in the last five years. In a mice model of glucocorticoid-induced delayed

bone repair, a common condition in fractured patients who use glucocorticoids to treat chronic inflammatory diseases, Okada *et al.* verified that dexamethasone treatment significantly decreased the number of F4/80⁺ cells at the femoral fracture site two days after injury, as well as the mRNA levels of M-CSF, MCP-1, IL-1 β , and stromal cell-derived factor-1 (SDF-1, also known as CXCL-12), cytokines involved with macrophage activation and differentiation (61).

Similar findings were reported in a study by Chen *et al.* (62), in a mouse model of postmenopausal osteoporosis, a condition in which delayed fracture healing is also frequently observed. The authors found that both endochondral ossification and callus remodeling were impaired in ovariectomized mice (OVX), which possessed a decreased expression of TNF- α and IL-6 and a lower frequency of M1 and M2 macrophages in the fracture hematoma, respectively, on days 1 and 14 after the fracture (62).

Diabetes is another recognized condition that adversely affects fracture healing, with patients taking twice the time to heal a fracture and at a higher risk of progressing to nonunion. Using a mouse model of streptozotocin-induced diabetes and the drill hole model, Shimoi *et al.* showed that the number of macrophages at the fracture site was significantly lower on day 2 after fracture and the mRNA levels of M-CSF, iNOS, IL-6 and CD206 were significantly decreased (63).

In skeletal fluorosis, a clinical manifestation caused by the excessive ingestion of fluoride and its incorporation into hydroxyapatite crystals, delayed fracture healing is also reported. In a rat fracture healing model, Du *et al.* found that the number of CD86⁺ M1 macrophages increased at the fracture site on day 7 after the fracture, while the number of CD206⁺ M2 macrophages decreased. At 21 days, the number of M1 macrophages was still high in the fluoride treated group, indicating that fluoride results in prolongation of the pro-inflammatory stage, and therefore inhibition of the tissue neoformation phase (64).

A prolonged initial inflammatory phase is also reported in fracture healing in aged individuals and is considered to be part of the systemic “inflammaging”, that is, the chronic and increased pro-inflammatory status associated with aging. Clark *et al.* showed, through bulk mRNA sequencing, that macrophages from old mice have a more pronounced M1 gene signature than macrophages from young animals, and that these M1 macrophages persistently accumulate at the fracture site, thus affecting the resolution of the inflammatory response and callus formation (65).

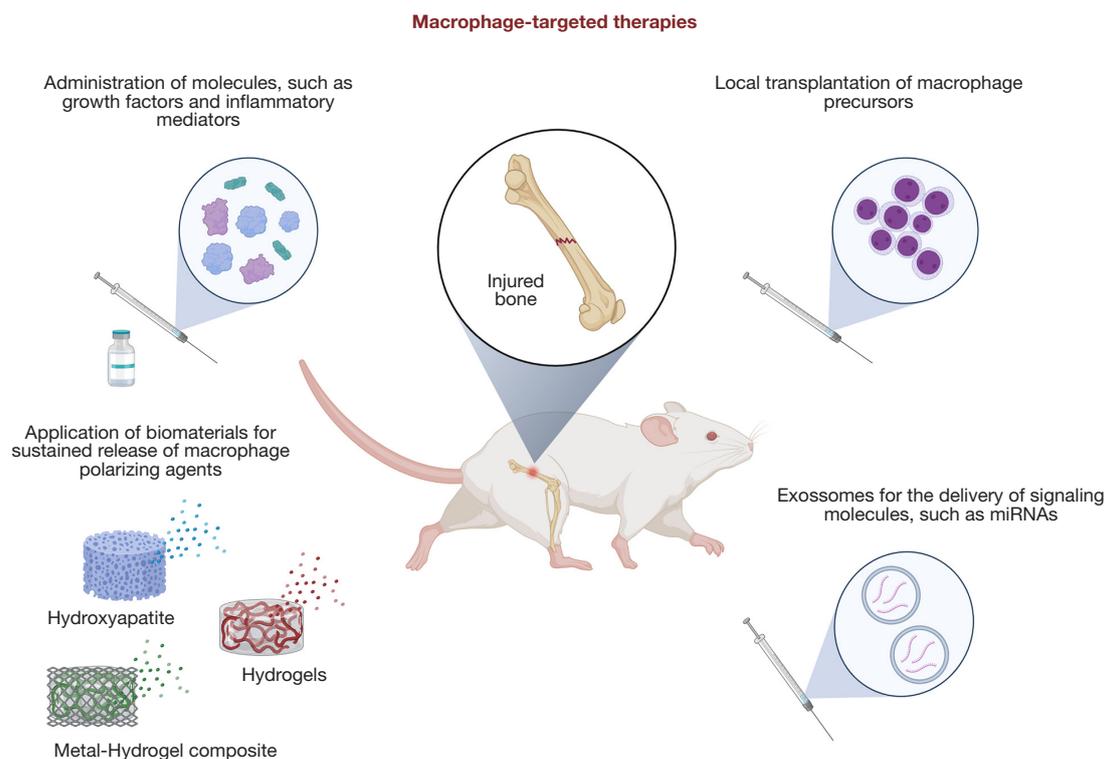


Figure 3 Main macrophage-oriented therapeutic strategies explored for the development of treatments to stimulate fracture healing. The latest perspectives for the development of innovative treatments for delayed and/or failed bone consolidation based on macrophage activity modulation include the local or systemic administration of molecules (mainly growth factors and inflammatory modulators, such as M-CSF and MaR1); the usage of biomaterials intended to regulate macrophage differentiation and polarization through sustained release of polarizing agents, such as IL-4 and copper ions; the local transplantation of macrophage precursors; and the use of exosomes to deliver signaling molecules that influence macrophage activities, such as miRNAs. Created with BioRender.com. M-CSF, macrophage colony-stimulating factor; MaR1, macrophage mediator in resolving inflammation; IL, interleukin.

Collectively, these studies strengthened the concept of the major influence of the inflammatory phase on the outcome of fracture healing and of macrophages as central players in the regulation of its first initiation and, later, of its resolution, opening new avenues for the development of treatments to stimulate bone repair.

Perspectives for the development of advanced therapies

Due to its preponderant role in fracture healing, several macrophage-targeted therapies have recently been tested in animal models. The newest research efforts include the following strategies: local or systemic administration of molecules (mainly growth factors and inflammatory modulators) and biomaterials intended to regulate

macrophage differentiation and polarization; local transplantation of macrophage precursors; and the use of exosomes to deliver signaling molecules that influence macrophage activities, such as miRNAs (Figure 3).

Batoon *et al.* investigated the effects of intermittent systemic administration of a chimeric M-CSF-1 molecule, which was developed to have an extended circulating half-life compared to native M-CSF-1, in healthy and osteoporotic femur diaphyseal fractures (66). The authors reported an improvement in callus development in healthy mice, which was correlated with an increase in the number of macrophages at the fracture site (66). These findings agreed with the report by Starlinger *et al.*, who showed that systemic application of M-CSF in fractured mice resulted in larger calluses with increased trabecular thickness (67). Most importantly, Batoon *et al.* showed that M-CSF-1 treatment

was also beneficial for the treatment of osteoporotic fractures, in which bone volume within the callus and cortical bridging were increased significantly, culminating in improved fracture biomechanical strength (66).

Similar results were observed in the study by Huang *et al.*, that investigated the impact of Maresin 1 (macrophage mediator in resolving inflammation, MaR1), a macrophage secreted molecule that signals both in autocrine and paracrine ways to decrease macrophage-associated inflammation, for the treatment of tibial fractures in aged mice (68). Upon systemic administration of MaR1, the authors verified that the fracture calluses of aged mice had increased bone volume and better structural stiffness, consistent with overall improved healing. These findings were correlated with a lower percentage of pro-inflammatory macrophages within the fracture callus and a decreased plasmatic level of inflammatory cytokines. *In vitro*, MaR1 treatment induced the expression of anti-inflammatory markers in macrophages, indicating its potential to act as a polarizing agent, shifting macrophage fate toward an anti-inflammatory phenotype, and contributing to the resolution of inflammation, which is naturally amplified in the elderly (68). Another study, published by Clark *et al.*, showed beneficial effects on the healing of fractures in old mice after the use of a pharmacological agent that antagonizes the M-CSF-1 receptor, thus inhibiting monocyte to macrophage differentiation (65).

Modification of biomaterials as a strategy to immunomodulate the fracture microenvironment has also been recently explored. Applying a biomimetic polysaccharide hydrogel-metal scaffold composite loaded with IL-4 and BMP-2 within femoral defects in rats, Wang *et al.* observed an increase in the number of M2 macrophages and in the proliferation and differentiation of skeletal progenitors into osteoblasts, which significantly improved bone regeneration (69). In another study, Xu *et al.* (70) developed an intramedullary nail composed of a copper-containing stainless steel as a strategy to deliver copper ions at the injury site, as this ion had been previously shown to considerably induce M2 macrophage polarization *in vitro* (71). Using the drill-hole injury model in the tibia of mice, the authors reported an accelerated formation of new cortical bone in the animals that received the copper-enriched intramedullary nail. When the callus was evaluated, it was observed an increased infiltration of CD206⁺ M2a macrophages, which located close to the newly formed type-H vessels and around the surface of the

neoformed osseous tissue (70).

The strategy based on local transplantation of macrophage precursors as an attempt to correct the imbalance between the M1/M2 macrophage pools at the fracture site of aged rats was explored in the study by Löffler *et al.* (72). The authors reported a partial rescue of bone regeneration after 6 weeks of injury, with increased bone deposition, reduced areas of fibrosis, and improved neovascularization within the calluses, presumably through induction of M2 macrophage differentiation (72). Vi *et al.* also investigated the possibility of modulating macrophage content within fractured bones through bone marrow transplantation and parabiosis strategies (73). In the study, when old animals received F4/80⁺ macrophages from young animals, their fracture calluses showed increased bone formation relative to animals that received old F4/80⁺ cells, indicating the potential of young macrophages to rejuvenate the repair process in aged animals. Among the factors produced by young macrophage cells that could be associated with this rejuvenation effect, the authors highlighted the role of LRP-1 (low-density lipoprotein receptor-related protein 1), which is involved in osteogenic differentiation of skeletal progenitors (73).

At the frontier of knowledge are the strategies that envision the use of exosomes, small vesicles secreted into the circulation by diverse cell types that contain various signaling molecules that once internalized by neighboring or distal cells, can alter the recipient function. Using a femur diaphyseal mice fracture model, Xiong *et al.* showed that the injection of M2 macrophage-derived exosomes, containing the microRNA miR-5106, at the injury site resulted in calluses with increased bone volume, reduced cartilage area and smaller fracture gaps, indicative of an accelerated fracture healing process (74). *In vitro* data showed that miR-5106 was able to stimulate osteogenic differentiation of skeletal progenitors, suggesting that the improvements in fracture healing observed *in vivo* may be related to the stimulation of skeletal cells (74). In a study by Zhang *et al.*, it was verified that the injection of macrophage-derived exosomes isolated from diabetic mice at the fracture site of healthy mice resulted in the development of calluses with significantly lower bone volume and larger fracture gaps (75). The authors identified the microRNA miR-144-5p as responsible for the observed impairments in fracture healing, possibly acting by suppressing osteogenic differentiation of skeletal progenitors (75). As the signaling activities of microRNAs can be molecularly targeted with antagonists, these findings point to a promising new

avenue of research that can be further explored to develop innovative therapies to improve fracture healing.

Conclusions

Research on fracture healing has significantly advanced the knowledge about the central cellular and molecular mechanisms that drive bone regeneration and lead to a successful clinical outcome, which ultimately is patient rehabilitation. While the first studies in the field focused on depicting the role of skeletal stem/progenitor cells in the process and how these could be explored to devise therapeutic strategies to promote bone regeneration (13,76-80), we now observe an important paradigm shift, with studies now focusing on the fact that fracture healing is a complex event, influenced by many factors beyond stem/progenitor cells, and that these factors deserve better attention. The role of macrophages in fracture healing has long been recognized (34,39,40,43,45,81-83), and recent literature consolidates this notion, opening promising new avenues for the development of advanced and innovative therapies based on targeted macrophage activities. However, considering that macrophages are highly heterogeneous and their activities must be precisely in concert with the spatio-temporal context of the fracture healing process, more research is still warranted to better understand their diversity (48,84), their specific role in each step of fracture healing, and to decipher the key molecular mechanisms involved in their *in vivo* crosstalk with other microenvironmental cells, especially with endothelial and skeletal stem/progenitor cells. Although many studies in the literature bring evidence of putative signaling pathways involved in the communication between macrophages and endothelial and skeletal cells, the majority come from *in vitro* studies, which do not fully recapitulate the complexity of the fracture healing microenvironment. Nevertheless, with the recent advancements in molecular biology tools, such as the diverse omics performed at the single cell level and next-generation sequencing, novel possibilities of investigation are open, increasing the possibility of effectively answering these questions and translating new and effective therapies for those who suffer with incapacitating bone fractures.

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Footnote

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Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

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