



Mesenchymal stem cells: From regeneration to cancer

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ABSTRACT

Mesenchymal stem cells (MSCs) are multipotent tissue stem cells that differentiate into a number of mesodermal tissue types, including osteoblasts, adipocytes, chondrocytes and myofibroblasts. MSCs were originally identified in the bone marrow (BM) of humans and other mammals, but recent studies have shown that they are multilineage progenitors in various adult organs and tissues. MSCs that localize at perivascular sites function to rapidly respond to external stimuli and coordinate with the vascular and immune systems to accomplish the wound healing process. Cancer, considered as wounds that never heal, is also accompanied by changes in MSCs that parallels the wound healing response. MSCs are now recognized as key players at distinct steps of tumorigenesis. In this review, we provide an overview of the function of MSCs in wound healing and cancer progression with the goal of providing insight into the development of novel MSC-manipulating strategies for clinical cancer treatment.

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Contents

1. Introduction	42
2. Tumors are wounds that do not heal: comparison of wound healing to injury with wound healing in tumorigenesis	43
3. In vitro and in vivo identity of mesenchymal stem cells	43
4. MSCs participate in all phases of the wound healing response to injury	44
5. MSCs in wound healing response to cancer	46
6. Concluding remarks	49
Conflict of interest statement	50
Acknowledgements	50
References	51

1. Introduction

Mesenchymal stem cells (MSCs) are multipotent progenitor cells with the potential to differentiate into diverse types of tissue cells,

Abbreviations: BM, Bone marrow; BMP, Bone morphogenetic protein; CSC, Cancer stem cell; ECM, Extracellular matrix; EMT, Epithelial-mesenchymal transition; HGF, Hepatocyte growth factor; IDO, Indoleamine 2,3-dioxygenase; IL-1, Interleukin-1; iNOS, Inducible nitric oxide synthase; MIF, Macrophage migration inhibitory factor; MMP, Matrix metalloproteinase; MSCs, Mesenchymal stem cells; PDGF, Platelet-derived growth factor; PGE₂, Prostaglandin E₂; TAF, Tumor-associated fibroblasts; TA-MSCs, Tumor-associated MSCs; TGFβ, Transforming growth factor beta; TIMP, Tissue inhibitors of metalloproteinase; TNFα, Tumor necrosis factor alpha; TSG-6, TNFα-stimulated gene-6; VEGF, Vascular endothelial growth factor.

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including osteoblasts, adipocytes, chondrocytes and myofibroblasts (Keating, 2012; Singer & Caplan, 2011). This type of tissue stem cell plays an essential role in tissue regeneration and closely interacts with cells of the immune system in the tissue microenvironment during repair from tissue damage (Le Blanc & Davies, 2015; Shi et al., 2012; Uccelli, Moretta, & Pistoia, 2008). Recently, MSCs have also emerged as a new player in the tumor microenvironment, contributing to tumor growth, metastasis and therapeutic resistance (Ridge, Sullivan, & Glynn, 2017; Shi, Du, Lin, & Wang, 2017).

Cancer, regarded as “non-healing wounds” (Dvorak, 1986, 2015), is believed to take advantage of the regenerative functions of host cells to facilitate local cancer growth, resistance to therapy, and metastases to distant organs (Calvo et al., 2015; Krall et al., 2018; Sundaram et al., 2017). In this review, we discuss how MSCs participate in distinct stages of wound healing and tumor “wound” progression, and compare the

functions and mechanisms of MSCs in these two pathological processes. Understanding how the tumor microenvironmental cues drive MSCs to regenerate tumor “wounds” will facilitate our deeper understanding of MSC biology in distinct steps of cancer progression, thereby supporting development of new cancer treatments that target MSCs.

2. Tumors are wounds that do not heal: comparison of wound healing to injury with wound healing in tumorigenesis

2.1. Wound healing response to injury

Upon tissue injury, a series of wound healing steps including inflammation, tissue proliferation and remodeling are successively initiated and highly coordinated by the adult tissue cells (Maxson, Lopez, Yoo, Danilkovitch-Miagkova, & Leroux, 2012). Blood clotting, or coagulation, occurs immediately after injury. This is achieved by platelet aggregation and clot formation from fibrinogen-converted fibrin and extracellular matrix (ECM) proteins. Blood clotting serves as the first barrier against blood and water loss as well as invading pathogens. After clot formation, an inflammation stage is initiated by neutrophil infiltration into the injury sites usually within a few hours but up to one day post injury, and is followed by the arrival of monocytes and mast cells one to two days later. The monocytes differentiate into macrophages, which persist long-term at the tissue sites until wound healing is complete. Myeloid cell migration is guided by chemotactic substances such as damage- or pathogen-associated molecular patterns (DAMPs or PAMPs), hydrogen peroxide (H₂O₂) and chemokines, which are released from the damaged tissues and surrounding stroma in response to injury and invading pathogens (de Oliveira, Rosowski, & Huttenlocher, 2016).

Migration of myeloid cells marks the beginning of the inflammation stage in wound healing from injury. Myeloid cells function to not only phagocytize the dead tissue cells or invading pathogens during the inflammation stage, but also further modulate the next stages of tissue proliferation and remodeling by releasing cytokines, chemokines and other trophic factors (Minutti, Knipper, Allen, & Zaiss, 2017). Along with myeloid cell infiltration, adaptive immune cells such as T lymphocytes are also recruited to the injury sites where they specifically target pathogens and release cytokines to further regulate inflammation, proliferation and tissue remodeling (Havran & Jameson, 2010; Keyes et al., 2016). In the absence of major infection, the inflammatory phase peaks at about two to three days post-injury and gives way to the proliferative phase that lasts about two weeks. Proliferation of three major cell types including epithelial cells (re-epithelialization), mesenchymal cells (fibroplasia) and endothelial cells (revascularization or angiogenesis), occurs concomitantly in the proliferation phase. Through both autocrine and paracrine mechanisms, a plethora of cytokines and trophic factors, such as epidermal growth factor (EGF), transforming growth factor beta (TGFβ), platelet-derived growth factor (PDGF), fibroblast growth factor (FGF) and vascular endothelial growth factor (VEGF), play essential roles in the acceleration of epithelial tissue re-growth, production of collagen and other ECM proteins and formation of new blood vessels (Guo & Dipietro, 2010). Following the proliferative phase, tissue remodeling continues from weeks to years depending on the wound type and size. During tissue remodeling, the disorganized collagen fibers formed during the proliferative phase will rearrange and align along tension lines. Wound contraction occurs in parallel with a reduction in the numbers of macrophages and fibroblasts and a reduction in blood vessels by apoptosis, which together achieve a successful healing process (Hinz et al., 2012).

2.2. Wound healing response to tumorigenesis

Tumors have long been regarded as wounds that fail to complete the normal three stages of wound healing (Dvorak, 1986). During early tumorigenesis, cancer cells are recognized by innate and adaptive immune cell-mediated host surveillance leading to apoptosis and/or

necrosis. The dead tumor cells in turn release various DAMPs such as adenosine, high-mobility group box 1 protein (HMGB1), annexins and calreticulin, which serve to initiate the inflammatory response followed by a cascade of events within the wound healing process (Hernandez, Huebener, & Schwabe, 2016). With tumor progression, the surrounding blood vessels become permeable which further triggers platelet and fibrin deposition, and, in turn, starts the program of inflammation, proliferation and ECM remodeling (Kreuger & Phillipson, 2016). As long as tumors are not eradicated by the host immune system or external therapies, the healing steps continue until the tumor burden exceeds the host's capability to survive. An apparent difference between normal physiological wound healing and tumorigenesis is that the latter possesses one or multiple prolonged, uncompleted phases, and tumors can therefore be considered as an “overhealing wound” (Schafer & Werner, 2008).

Tumor progression is closely associated with the three phases of wound healing. The first phase, inflammation, a hallmark of cancer, is in fact a double-edged sword in tumor development and therapy responses. The innate and adaptive immune cells are phenotypically and functionally plastic depending on their resident microenvironment. Type 1 immune cells are conventionally activated cells, such as “N1” neutrophils, “M1” macrophages and “Th1” or “Tc1” T cells. These cells are mainly considered to be tumoricidal and exert direct cytotoxic effects against tumor cells. In contrast, the alternatively activated type 2 immune cells function to promote distinct steps in cancer progression through release of multiple cytokines and chemokines (Gabrilovich, Ostrand-Rosenberg, & Bronte, 2012; Nowarski, Gagliani, Huber, & Flavell, 2013). During the second phase, the proliferation phase, angiogenesis (revascularization) and desmoplasia (fibroplasia) coordinately support epithelial tumor growth (re-epithelialization) via a collection of growth factors, which are equally essential for wound healing. The third phase, tissue remodeling, completes the wound healing response but fails to be completed in the context of a tumor. However, many of the tissue remodeling activities that occur during wound healing, such as lysyl oxidase (LOX)-induced collagen crosslinking and matrix metalloproteinase (MMP)-mediated collagen rearrangement and alignment, have been shown to be pivotal in tumor cell migration, invasion and metastasis, as well as in resistance to therapy (Cox & Erler, 2011; Lu, Weaver, & Werb, 2012).

3. In vitro and in vivo identity of mesenchymal stem cells

MSCs are multipotent stem cells that can differentiate into multiple cell lineages including adipocytes, osteoblasts, chondrocytes, tenocytes and myofibroblasts. In the 1960s and 1970s, this type of adult stem cell was initially identified as a population of bone marrow (BM) cells with colony forming capabilities in vitro and osteogenic cell differentiation potential in vivo (Friedenstein, Chailakhjan, & Lalykina, 1970; Friedenstein, Petrakova, Kurolesova, & Frolova, 1968). In the 1980s and 1990s, these cells were defined as “Mesenchymal Stem Cells” possessing multiple non-hematopoietic mesenchymal lineage cell differentiation capabilities in vitro and in vivo (Caplan, 1991; Prockop, 1997). Afterwards, MSCs were isolated from almost every tissue in the body (umbilical cord, Wharton's jelly, skin, lung, liver, adipose tissue, muscle, dental pulp, etc.) in addition to the BM (da Silva Meirelles, Chagastelles, & Nardi, 2006). In recent years, MSCs were also isolated from various tumor tissues as “tumor wounds” could stimulate resident MSC proliferation or recruit circulating MSCs (Karnoub et al., 2007; Ren et al., 2012; Shi et al., 2017).

According to criteria by the International Society for Cellular Therapy (ISCT) published in 2006, cultured MSCs should be adherent, fibroblast-like cells with osteogenic, adipogenic and chondrogenic differentiative capacity in vitro. Moreover MSCs must express the surface markers CD105, CD73 and CD90, but not express CD45, CD34, CD14, CD11b, CD79α, CD19 or human leukocyte antigen-DR isotype (HLA-DR) (Dominici et al., 2006). In 2013, ISCT further shared guidelines for the

immunological characterization of MSCs, highlighting the functional plasticity of MSCs in the context of different inflammatory milieus (Krampera et al., 2013). In spite of all these efforts in defining MSCs, the boundaries for delineating MSCs, fibroblasts and pericytes still remain largely illusive due to a lack of specific markers for these cell populations (Caplan, 2008; Haniffa, Collin, Buckley, & Dazzi, 2009; Keating, 2012; Nombela-Arrieta, Ritz, & Silberstein, 2011). Nevertheless, MSCs were recently described as “Medicinal Signaling Cells” to reflect their multifaceted functions in wound repair rather than considering their stemness properties (Caplan, 2017). Based upon their robust capacity in regulating every step of wound healing, cultured MSCs have been applied in over 450 clinical trials to treat degenerative diseases and immune disorders (Maxson et al., 2012; Squillaro, Peluso, & Galderisi, 2016).

While the clinical application of ex vivo-expanded MSCs in treating diseases has been established, in vivo exploration of endogenous MSCs has also progressed. Pioneering work by Crisan et al., 2008 demonstrated a perivascular origin of MSCs in multiple human organs. They purified perivascular cells from human skeletal muscle, pancreas, adipose tissue and placenta and showed after long-term culture that these cells maintained the MSC phenotype as well as the trilineage differentiation potential (Crisan et al., 2008). Since then, the endogenous MSCs were believed to reside at perivascular sites in various tissues which may be regarded as a subset of pericytes in vivo (Caplan, 2008, 2017). Genetic lineage tracing assays were also widely applied to identify the differentiation fates of perivascular MSC-like cells in the context of fibrotic diseases and tissue injury models (El Agha et al., 2017). A series of transgenic mice with Cre/loxP technology have recently been created and applied to explore the cellular hierarchies of the perivascular MSCs, such as Nestin-cre (Mendez-Ferrer et al., 2010; Tronche et al., 1999), myxovirus resistant 1 (Mx1)-cre (Kuhn, Schwenk, Aguet, & Rajewsky, 1995; Park et al., 2012), Leptin-receptor (Lepr)-cre (Decker et al., 2017; DeFalco et al., 2001; Zhou, Yue, Murphy, Peyer, & Morrison, 2014), and glioma-associated oncogene homolog 1 (Gli1)-cre (Ahn & Joyner, 2004; Kramann et al., 2015; Schneider et al., 2017). Although the differentiation specificity and efficiency of the perivascular MSCs still remain controversial, most reports support the notion that endogenous MSCs are highly plastic upon tissue injury and give rise to myofibroblasts, osteoblasts or adipocytes in bone, lung, liver, kidney, heart, spinal cord, muscle, and skin depending on the injury types or duration. It is also well accepted that cells differentiating from endogenous tissue MSCs play a major role in induction of organ or tissue fibrosis (El Agha et al., 2017). In contrast to the extensive studies that recognize the in vivo identity of MSCs in tissue regeneration, far fewer studies have been carried out to determine the role of endogenous MSCs in primary tumor tissues and metastatic sites.

4. MSCs participate in all phases of the wound healing response to injury

Perivascular mesenchymal cells including MSCs play crucial roles through every step of wound healing. Immediately after tissue injury, MSCs express tissue factor (TF) and Factor VIII:c (FVIII) which activate both extrinsic and intrinsic coagulation pathways to facilitate blood clotting (Christy et al., 2017; Sanada et al., 2013). In addition to driving the inflammation stage (described below) MSCs themselves also secrete antimicrobial peptides such as LL-37, hepcidin, β -defensin 2 and lipocalin-2 serving to restrain the invading pathogens (Alcayaga-Miranda et al., 2015; Gupta et al., 2012; Krasnodembskaya et al., 2010; Sung et al., 2016). We next provide an overview of the participation of MSCs in the three stages of wound healing: inflammation, proliferation and tissue remodeling (outlined in Fig. 1).

4.1. Inflammation

In the inflammatory stage, MSCs play dual roles serving as both facilitators and terminators of the inflammation process. At the beginning of inflammation or infection, wound signals such as toll-like receptor (TLR) ligands stimulate bone marrow MSCs to secrete the chemokine CCL-2 which triggers the emigration of monocytes from the BM to the circulation (C. Shi et al., 2011). At the wound sites, paracrine chemokines such as CCL-2, CCL-3, CCL-4, CXCL1, IL-8 and macrophage migration inhibitory factor (MIF) expressed by naive or TLR ligand-stimulated MSCs serve to recruit monocytes, macrophages and neutrophils (Brandau et al., 2010; Chen, Tredget, Wu, & Wu, 2008; Romieu-Mourez et al., 2009). Intravital microscopy, a technique that allows visualizing the biological events in live animals, has enabled us to understand how perivascular mesenchymal cells, including endogenous MSCs, regulate myeloid cell infiltration at the early stage of inflammation in vivo (Proebstl et al., 2012). Upon stimulation by lipopolysaccharide (LPS) or inflammatory cytokines, endogenous perivascular cells (or pericytes) upregulate their expression of the chemokine CXCL1 or MIF, as well as the adhesion molecule intercellular adhesion molecule 1 (ICAM-1). Chemokines serve to recruit neutrophils and monocytes for extravasation whereas adhesion molecules in turn guide these emigrating myeloid cells as they crawl on perivascular cells and move towards the inflammatory foci (Proebstl et al., 2012; Stark et al., 2013).

A myriad of mechanisms have evolved to impede excessive innate immune responses following the peak of inflammation. Dampening of innate immunity is particularly critical in cases of sterile inflammation. Perivascular MSCs indeed play a “gatekeeper” role to limit inflammation-caused tissue damage. Such effects are achieved through a negative feedback mechanism driven by inflammatory mediators such as tumor necrosis factor alpha (TNF α), interleukin-1 (IL-1) and reactive oxygen species (ROS). These factors stimulate MSCs to upregulate cyclooxygenase-2 (COX-2), TNF α -stimulated gene-6 (TSG-6), superoxide dismutase 3 (SOD3) and possibly other effector molecules, to suppress the phagocytic activities of type 1 myeloid cells leading to remission of inflammatory responses (Francois, Romieu-Mourez, Li, & Galipeau, 2012; Jiang et al., 2016; Lee et al., 2009; Nemeth et al., 2009). Notably, the COX-2- prostaglandin E2 (PGE2) pathway and TSG-6 have been extensively studied in MSC-mediated reprogramming of M1 macrophages to M2 regulatory macrophages. These regulatory macrophages preferentially release higher levels of cytokines and growth factors to drive the wound healing process into the proliferation stage (Mittal et al., 2016; Nemeth et al., 2009; Ylostalo, Bartosh, Coble, & Prockop, 2012). In addition to these mechanisms, a recent report suggested that engulfment of apoptotic MSCs can also reprogram the host phagocytes to be an immunosuppressive phenotype via production of indoleamine 2, 3-dioxygenase (IDO) which in turn inhibits graft-versus-host disease (GvHD) in mice (Galleu et al., 2017). In addition to modulating monocytes, macrophages and neutrophils, MSCs are also able to repress the functions of other innate immune cell populations including natural killer (NK) cells, dendritic cells and mast cells through PGE2 release and other mechanisms (Aggarwal & Pittenger, 2005; Cui et al., 2016; Jiang et al., 2005; Spaggiari et al., 2008; Spaggiari, Capobianco, Becchetti, Mingari, & Moretta, 2006; Su, Zhang, Shi, Nguyen, & Le, 2011).

Compared to the innate immune responses that participate early in the wound healing process, T and B lymphocyte-mediated adaptive immune responses participate at a later time point (Strbo, Yin, & Stojadinovic, 2014). Similar to their dual roles in regulation of innate immune responses, MSCs are capable of augmenting or inhibiting the activities of the adaptive immune cells. When concentrations of environmental stimuli or inflammatory cytokines are low, in vitro cultured MSCs can boost T-cell responses likely through

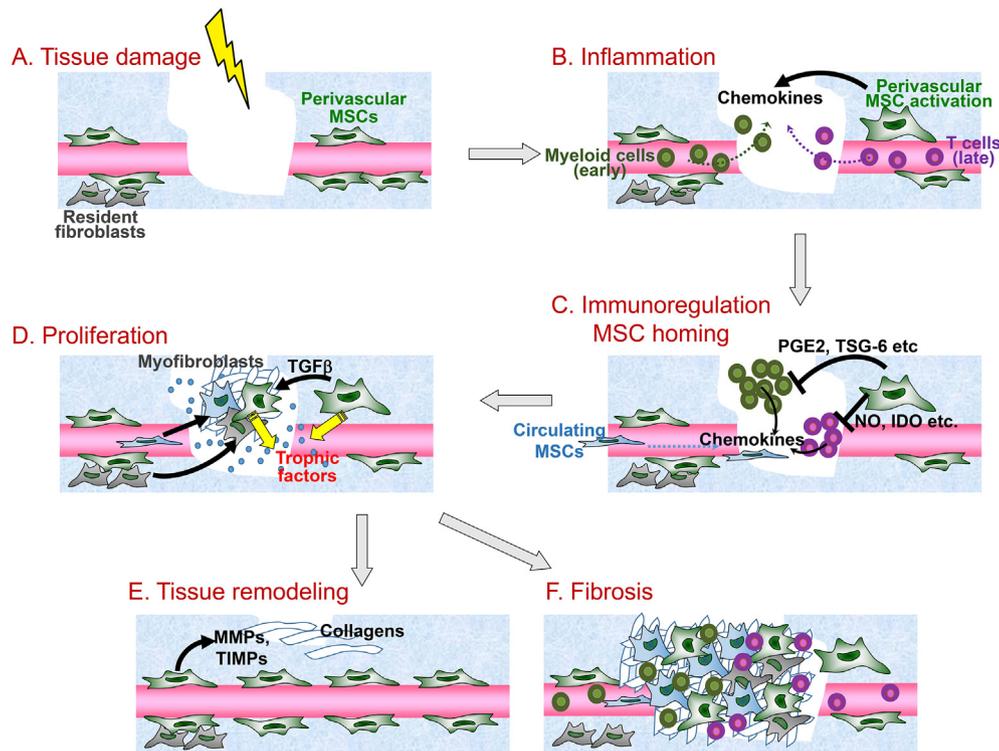


Fig. 1. Participation of MSCs in distinct steps of wound healing. MSCs are involved in all stages of wound healing. (A) After tissue damage, perivascular MSCs are activated. (B) At the inflammation stage, MSCs recruit both innate and adaptive immune cells via secretion of chemokines. (C) Through negative feedback loops, MSCs regulate the immune response (immunoregulation) by suppressing excessive innate and adaptive immune cell activities through release of prostaglandin E2 (PGE2), TSG-6, nitric oxide (NO), indoleamine 2, 3-dioxygenase (IDO) and others. In the meantime, chemokines produced by immune cells and stromal cells further recruit BM- and other tissue-derived MSCs (MSC homing). (D) MSCs then differentiate into myofibroblasts via transforming growth factor beta (TGF β) signaling and also produce a series of trophic factors to support tissue cell proliferation to accomplish re-epithelialization, fibroplasia and revascularization. (E) At the tissue remodeling stage, MSCs express matrix metalloproteinases (MMPs) as well as tissue inhibitor of metalloproteinases (TIMPs) to re-organize the extracellular matrix (ECM) structures leading to scar formation and wound repair. (F) In the cases of chronic inflammation and infection, the dysregulated MSC differentiation program can result in uncontrolled myofibroblast generation, causing tissue fibrosis.

MSC-secreted T-cell chemokines (Li et al., 2012; Renner et al., 2009). With the elevation of the T-cell responses, the high levels of T-cell cytokines subsequently turn on the negative feedback mechanisms in MSCs to dampen excessive T-cell reactivity. A key T-cell cytokine in this regulation is IFN γ , which acts independently or with inflammatory mediators including TNF α , IL-1, interleukin 17A (IL-17) and TLR ligands to functionally convert MSCs from a resting state to a highly immunosuppressive stage (Han et al., 2014; Opitz et al., 2009; Wang, Chen, Cao, & Shi, 2014). Such an immunosuppression is achieved via cytokine-induced expression of the immunosuppressive molecules inducible nitric oxide synthase (iNOS) and IDO in mouse and human MSCs, respectively (Krampera et al., 2006; Ren et al., 2009; Ren et al., 2008; Sato et al., 2007; Su et al., 2014). Other effector molecules and cell populations implicated in MSC-mediated T-cell immunoregulation include PGE2, programmed death ligand 1 (PD-L1), heme oxygenase-1 (HO-1), leukemia inhibitory factor (LIF), IL-6, galectins, Fas ligand, TGF- β and regulatory T cells (reviewed in (Y. Shi et al., 2012; Singer & Caplan, 2011)). Certain chemokines and adhesion molecules that commonly play roles in accelerating immune responses, however, serve to recruit immune cells to form an MSC-immune cell interaction that supports a more robust immunosuppression (Espagnolle, Balguerie, Arnaud, Sensebe, & Varin, 2017; Lee et al., 2017; Ren et al., 2010; Rubtsov et al., 2017). Such a cytokine-elicited immunoregulatory mechanism was also reported in other endogenous stromal cells such as fibroblastic reticular cells (FRCs) and lymphatic endothelial cells (LECs) suggesting a common negative feedback regulation employed by distinct stromal subsets in vivo (Lukacs-Kornek et al., 2011; Siebert et al., 2011).

4.2. Proliferation

After completion of the inflammation stage, MSCs further functionally contribute to wound healing by participating in the proliferation stage mainly via secretion of trophic factors. In addition to the resident perivascular MSCs, other tissue/organ-derived MSCs are recruited into the wound sites guided by chemokines and adhesion molecules produced during the inflammation stage (Karp & Leng Teo, 2009). The major chemokine-chemokine receptors involved in MSC trafficking include CXCL12-CXCR4, CCL-2-CCR2, CCL27-CCR10, and CCL21-CCR7 (Alexeev, Donahue, Uitto, & Igoucheva, 2013; Belema-Bedada, Uchida, Martire, Kostin, & Braun, 2008; Hu et al., 2013; Kitaori et al., 2009; Sasaki et al., 2008). Coordinated with chemokine-mediated chemotaxis, CD44-hyaluronic acid (HA) and vascular cell adhesion protein 1 (VCAM-1)- α 4/ β 1 integrin interactions, as well as MMP-mediated ECM degradation, facilitate MSCs to transmigrate into the endothelium (Ries et al., 2007; Ruster et al., 2006; Zhu et al., 2006).

Both resident and newly recruited MSCs exert two main effects in the proliferation stage at the wound sites. First, MSCs are directly involved in the tissue repair process through differentiating into multiple mesenchymal lineages. In the 1990s, the exogenously implanted BM-MSCs were shown to have a potent osteogenic potential as detected by the “cube assay” in vivo (Dennis, Haynesworth, Young, & Caplan, 1992; Dennis, Kostantakos, Arm, & Caplan, 1998). Further, by fluorescent protein labeling and in vivo genetic lineage tracing assays, exogenous and endogenous MSCs were demonstrated to possess the capacity to differentiate into other mesenchymal lineages such as adipocytes and myofibroblasts in various injury conditions (Anjos-Afonso, Siapati, & Bonnet, 2004; Kramann et al., 2015; Park et al., 2012;

Uezumi, Fukada, Yamamoto, Takeda, & Tsuchida, 2010). Second, MSCs and their lineages serve as a reservoir of trophic agents continuously supplying growth factors to accelerate cell proliferation towards a successful damage repair. The major factors produced by mesenchymal lineages include EGF, PDGF, FGF, VEGF, TGF β , keratinocyte growth factor (KGF) and hepatocyte growth factor (HGF) (Caplan & Dennis, 2006; Hofer & Tuan, 2016). Most of the wound-related cell types including epithelial cells, endothelial cells, keratinocytes and resident fibroblasts respond to these MSC-derived growth factors for their survival, activation and proliferation to accomplish re-epithelialization, angiogenesis and fibroplasia (Maxson et al., 2012; Zambetti et al., 2016).

4.3. Tissue remodeling

In the tissue remodeling stage, MSCs regulate collagen deposition, degradation and rearrangement through release of MMPs and tissue inhibitors of metalloproteinases (TIMPs). Upon stimulation by TGF- β , IL-1, and TNF α , MSCs overexpress MMP-2, MMP-9, and membrane type-1 MMP (MT1-MMP) which drive ECM degradation and support MSC cell invasion in three-dimensional (3-D) cultures (Lu, Li, Hu, Rowe, & Weiss, 2010; Ries et al., 2007). In a rat model of myocardial infarction (MI), implanted MSCs promoted the expression of MMP2 by cardiac fibroblasts and reduced cardiac ventricular fibrosis after MI (Mias et al., 2009). On the other hand, MSCs secrete the MMP inhibitors-TIMPs, which may protect vascular matrix molecules and endothelial cell structures from MMP-induced disruption (Lozito & Tuan, 2011). Recent evidence also suggests that MSC-derived TIMP-1 can function as an anti-angiogenic effector molecule in the inflamed endothelium to reduce the inflammatory response (Zanotti et al., 2016). Lineage tracing of the endogenous MSCs indicated that they can directly give rise to myofibroblasts and contribute to dysregulated ECM remodeling leading to tissue fibrosis. Such roles of MSCs in fibrotic diseases

have been extensively reviewed recently (El Agha et al., 2017) and will not be discussed here.

Overall, MSCs are not only beneficial to all steps of wound healing, but also serve as a facilitator for non-healing pathological processes such as fibrosis and cancer when chronic stresses (injury, infection, etc.) are present.

5. MSCs in wound healing response to cancer

Cancer is regarded as an “overhealing wound” and MSCs recently emerged as a new player in the tumor wound microenvironment. Tumor-associated MSCs (TA-MSCs), isolated from malignant tissues, actively participate in tumor-associated inflammation, immunosuppression, tumor growth, angiogenesis and tumor metastasis in various cancer types including breast cancer, ovarian cancer, pancreatic cancer, lymphoma, melanoma and others in both human and mouse tumor models (reviewed in (Shi et al., 2017)). These isolated TA-MSCs also have multipotent differentiation capacities for mesenchymal lineages with an ability to generate fibroblastoid colony-forming units (Karnoub et al., 2007; McLean et al., 2011; Ren et al., 2012). Here we present an overview of the roles of MSCs in cancer progression from the point of view of cancer as a wound (outlined in Fig. 2).

Dysfunction of MSCs is thought to contribute to tumorigenesis. Indeed, mutation of *tyrosine-protein phosphatase non-receptor type 11* (*Ptpn11*) in MSCs was recently shown to trigger a CCL-3-monocytes-IL-1 β axis leading to BM inflammation and myeloproliferative neoplasm in a mouse model (Dong et al., 2016). Similarly, deletion of the *Shwachman-Bodian-Diamond syndrome* (*SBDS*) gene in *osterix*⁺ mesenchymal progenitor cells caused a p53-S100A8/9-TLR inflammatory response which drove the development of Shwachman-Diamond syndrome (SDS) and myelodysplastic syndrome (MDS) in mice (Zambetti et al., 2016). In contrast to these blood malignancies elicited by

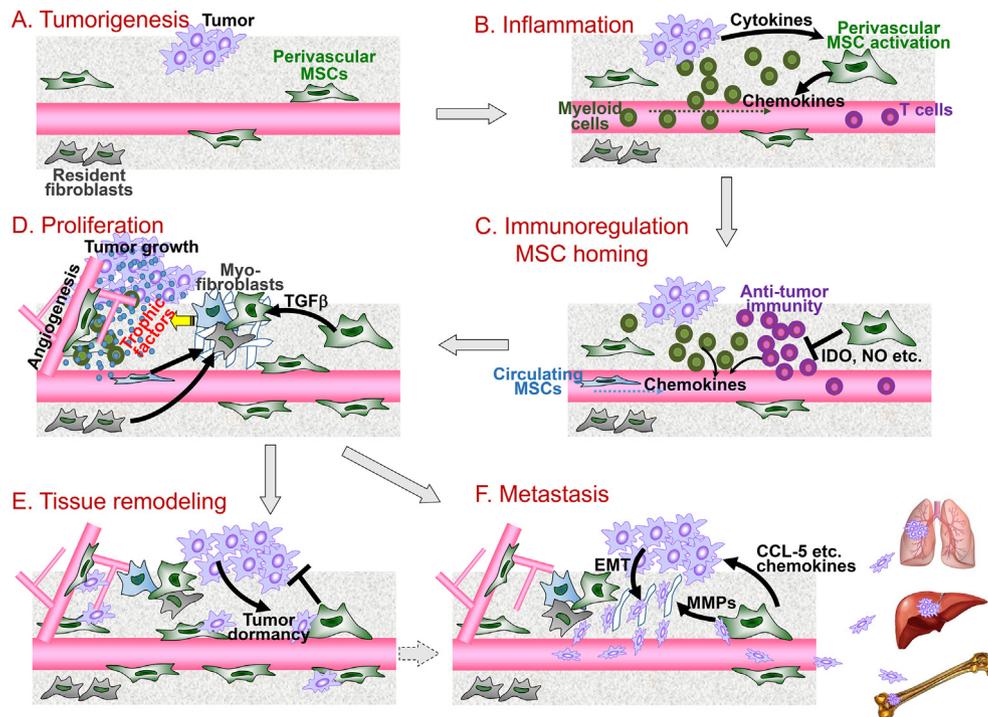


Fig. 2. Roles of tumor-associated MSCs (TA-MSCs) in the tumor “wounds”. TA-MSCs are essential for distinct steps of cancer progression from the view of cancer as “wounds”. (A) During tumorigenesis, perivascular MSCs are activated. (B) Inflammatory cytokines stimulate TA-MSCs to produce myeloid cell chemokines which recruit inflammatory cells to the tumor microenvironment causing an aggravated inflammation. (C) TA-MSCs suppress anti-tumor adaptive immunity through immunosuppressive effector molecules IDO, NO, and others. Guided by the chemotactic cues, circulating MSCs home to the tumor sites. (D) TA-MSCs are further differentiated into myofibroblasts (or CAFs), and both TA-MSCs and myofibroblasts produce trophic factors to support tumor cell proliferation and angiogenesis. (E) At the tissue remodeling stage, TA-MSCs may suppress the outgrowth of tumor cells while impelling them to enter dormancy. Such a dormant cancer stem cell status plays a key role in tumor recurrence and therapeutic resistance. (F) TA-MSCs secrete multiple chemokines and also stimulate the tumor cells to undergo epithelial-mesenchymal transition (EMT) which instigates tumor cell invasion and metastasis to distant organs such as lung, liver and bone.

abnormal BM-MSCs, it remains largely undefined how tissue MSCs regulate early tumorigenesis events in carcinomas.

If we consider cancer progression in terms of the three stages of wound healing, inflammation, proliferation and tissue remodeling, recent studies using exogenous MSC implantation models showed that MSCs exert either supportive or inhibitory effects on tumor development. While these studies were informative, future efforts should be directed toward a precise understanding of the role of endogenous MSCs in distinct steps of cancer progression. Below we discuss the contribution of MSCs to distinct steps of tumor “wound healing”, although some processes are intermingled.

5.1. Inflammation

5.1.1. TA-MSCs facilitate tumor-associated inflammation

Similar to their effects in wound healing, MSCs play a regulatory role in tumor-associated immune responses. First, TA-MSCs, thought to be derived from healthy MSCs at the tumorigenesis sites, are educated by the tumor inflammatory microenvironment (Ren et al., 2012; Ren et al., 2014), after which they have the capacity to elicit tumor-associated inflammation via secretion of cytokines and chemokines. The tumor and stroma-derived factors such as TNF α and IL-1 stimulate the TA-MSCs to specifically elevate their expression of chemokines, which in turn induce myeloid cell infiltration to exaggerate tumor-associated inflammation (Escobar et al., 2015; Ren et al., 2012; Yu et al., 2017). In mouse and human models of lymphoma, TA-MSCs were shown to express CCL-2, which serves to recruit monocytes and macrophages into the tumor microenvironment to sustain tumor growth (Guilloton et al., 2012; Ren et al., 2012). TA-MSCs also potently recruit neutrophils through overexpression of chemokines CXCL1 and CXCL2, and the neutrophils in turn stimulate primary tumor cell invasion and metastasis to distant organs in mice (Yu et al., 2017). In a human breast cancer xenograft model, the MSC-breast cancer cell interaction provoked recruitment of both macrophages and neutrophils through the colony stimulating factor 1 (CSF1)-CSF1 receptor signaling pathway, resulting in enhanced cancer cell metastases (Chaturvedi, Gilkes, Takano, & Semenza, 2014).

5.1.2. TA-MSCs reprogram innate immune cells

In the tumor microenvironment, TA-MSCs further convert the recruited myeloid cells from a type 1 to a type 2 phenotype by abolishing their phagocytic abilities while activating their “healing” potentials (Biswas & Mantovani, 2010; Coffelt, Wellenstein, & de Visser, 2016). In both in vivo and in vitro models, TA-MSCs isolated from lymphoma or pancreatic carcinoma were shown to polarize macrophages into an M2-like phenotype with increased expression of the alternatively activated macrophage markers (Guilloton et al., 2012; Mathew et al., 2016; Ren et al., 2012). These M2 macrophages function to support tumor growth and depletion of these macrophages substantially reduced TA-MSC-mediated tumor promoting effect in vivo (Mathew et al., 2016; Ren et al., 2012). Similar to the TA-MSC-mediated education of macrophages, TA-MSCs also accelerate a preferential differentiation of leukocytes into immunosuppressive myeloid-derived suppressor cells (MDSCs) (Chen et al., 2013; Giallongo et al., 2016; Yen et al., 2013). In vitro, human MSCs induced a differentiation of human peripheral blood leukocytes (PBLs) into MDSCs through secretion of HGF and CXCL3 (Chen et al., 2013; Yen et al., 2013). In vivo, knockdown of HGF in co-injected MSCs caused a reduction of tumor-infiltrating MDSCs in a human colon cancer xenograft model (Yen et al., 2013). In multiple myeloma and chronic myeloid leukemia, TA-MSCs isolated from the patients' BM had an elevated ability to induce MDSC expansion compared to healthy donor-derived BM-MSCs (Giallongo et al., 2016; Giallongo, Tibullo, et al., 2016). Together, these studies showed that TA-MSCs are a key regulator of the innate immune cell plasticity.

5.1.3. TA-MSCs suppress adaptive immunity

In addition to their capacity to elicit tumor-associated inflammation, TA-MSCs also suppress adaptive immunity in the tumor microenvironment. The role of MSC-mediated immunosuppression in tumor progression was first indicated by the finding that subcutaneous injection of B16 mouse melanoma cells led to tumor growth in allogeneic recipients only when MSCs were co-injected (Djouad et al., 2003). Subsequently, MSC-derived immunosuppressive effector molecules such as NO andIDO were found to ablate anti-tumor T cell and NK cell immunity and facilitate tumor growth (Gazdic et al., 2017; Han et al., 2011; Huang et al., 2014; Ling et al., 2014; Liotta et al., 2015). Besides the production of immunosuppressive factors, TA-MSCs also abrogate adaptive immunity by inducing immunoregulatory cells such as T regulatory cells (Treg) and regulatory CD8⁺ T cells in vitro and in vivo (Hof-Nahor et al., 2012; Kudo-Saito, Fuwa, Murakami, & Kawakami, 2013; Patel et al., 2010).

Overall, TA-MSCs accelerate tumor-associated inflammation but suppress anti-tumor adaptive immunity. These two mechanisms, though beneficial in a conventional wound healing response, are utilized by the tumor cells to enhance their progression.

5.2. Proliferation

After completion of the inflammation stage of wound healing, there is a transition into the proliferation stage characterized by exogenous tissue precursor cell recruitment and resident tissue cell differentiation and proliferation. TA-MSCs have been shown to actively participate in the proliferative stage in various cancers.

5.2.1. MSC homing to tumors

A preferential homing of MSCs to the tumor, a process equivalent to the homing of MSCs to the wound site in wound healing, is a hallmark of MSC biology in cancer. In both syngeneic and xenogeneic tumor models, systemically delivered luciferase-labeled or GFP-labeled MSCs showed a persistent and specific co-localization at the sites of tumor progression (Huang et al., 2014; Kidd et al., 2009). Such tumor-oriented MSC trafficking is coordinated by tumor-secreted paracrine factors and the autocrine BM-MSC-expressed chemoattractants. Many tumor-derived factors such as CCL-2, CCL-25, CXCL16, MIF, IL-6 and antimicrobial peptide LL-37 have been reported to be functional in recruiting MSCs in vitro and in vivo (Coffelt et al., 2009; Dwyer et al., 2007; Jung et al., 2013; Lourenco et al., 2015; Rattigan, Hsu, Mishra, Glod, & Banerjee, 2010; Xu et al., 2012). Complementary to these paracrine mechanisms, BM-MSCs also utilize the CXCL12/CXCR4 autocrine loop to impel their own migration towards tumors as shown in human colorectal cancer and mouse gastric cancer cell models (Menon et al., 2007; Quante et al., 2011).

When arriving at the tumor sites newly recruited MSCs, together with resident MSCs, cooperatively accelerate tumor “wound” healing. The TA-MSCs themselves undergo proliferation and differentiation into myofibroblasts to build a collagen network. Moreover, TA-MSCs support the survival and growth of cancer cells as well as cancer stem cells, and supply angiogenic factors for neovascularization. Again, these natural mechanisms in wound healing are well exploited by malignant cells to support their own progression.

5.2.2. TA-MSC transdifferentiation to myofibroblasts

Myofibroblasts, or tumor-associated fibroblasts (TAFs), are one of the major populations of the tumor stromal cells. They exert diverse effects at distinct steps during cancer progression (Kalluri, 2016). Resident and distant tissue (such as BM)-derived MSCs have been reported as precursors of TAFs. In a syngeneic mouse model of ovarian cancer, the origins of TAFs in the tumor microenvironment have been quantitatively assessed in vivo (Kidd et al., 2012). It was shown that ~40% of tumor stromal cells are BM-derived. Among the TAFs, most fibroblast specific protein (FSP) positive and fibroblast activation protein (FAP) positive TAFs originate from BM-MSCs, whereas α -SMA⁺ TAFs

and perivascular stromal cells (pericytes) are mainly derived from the adipose tissue adjacent to the tumor (Kidd et al., 2012). In various syngeneic and xenograft tumors such as gastric cancer, breast cancer, glioma, pancreatic cancer, ovarian cancer and prostate cancer, exogenously implanted BM-MSCs were shown to differentiate into α -SMA⁺ vimentin⁺ myofibroblasts in the tumor microenvironment. This process was largely dependent on the TGF β /Smad signaling axis (Barcellos-de-Souza et al., 2016; Mishra et al., 2008; Quante et al., 2011; Shangguan et al., 2012; Spaeth et al., 2009; Spaeth et al., 2013). Together, these experiments demonstrate that myofibroblasts are a major MSC-derived cell lineage in the tumor microenvironment.

5.2.3. Trophic effects of TA-MSCs on tumor cells and cancer stem cells

In addition to transitioning into myofibroblasts to sustain cancer progression, TA-MSCs also directly release trophic factors to epithelial and hematologic malignant cells. In particular, TA-MSCs provide survival and pro-proliferative signals to cancer stem cells (CSCs) and tumor-initiating cells (TICs), leading to therapeutic resistance and early relapse (Y. Shi et al., 2017). In the tumor microenvironment, TA-MSCs facilitate tumor initiation and cause an increase in the numbers of aldehyde dehydrogenase (ALDH) positive CSCs in xenograft models of breast cancer, ovarian cancer, colorectal cancer and glioma. These effects were mainly exerted through TA-MSC-derived pro-survival signals such as IL-6, IL-8, CXCL1 CXCL7, and bone morphogenetic protein (BMP) (Hossain et al., 2015; Li, Reinhardt, Herschman, & Weinberg, 2012; Liu et al., 2011; McLean et al., 2011; Tsai et al., 2011). In a breast carcinoma xenograft model, cell-cell contact between TA-MSCs and tumor cells led to induction of miR-199a and subsequent repression of FOXP2, a transcriptional regulator inhibiting CSC associated factor. Thereby, TA-MSCs elicited a propagation of breast CSCs through the miR-199a-FOXP2 axis (Cuiffo et al., 2014). In accordance with these pre-clinical results, extensive bioinformatic and immunohistochemical analyses of human colorectal cancer patient specimens suggested that stromal gene expression is associated with a high frequency of CSCs and disease relapse (Calon et al., 2015). Thus, TA-MSCs directly function to nourish the tumor cells and CSCs in the tumor microenvironment.

5.2.4. TA-MSCs support angiogenesis

In normal wound healing, MSCs release angiogenic factors such as VEGF to foster growth of neovascular vessels for angiogenesis. This mechanism is also executed by TA-MSCs. In syngeneic models of melanoma, lung cancer and colorectal cancer, administration of mouse BM-MSCs promotes tumor growth through increased angiogenesis via the hypoxia-inducible factor 1 (HIF1)-VEGF signaling pathway (Liu et al., 2011; Suzuki et al., 2011). In a human pancreatic cancer xenograft model, GFP-labeled human BM-MSCs, when systemically administered, were found to attach onto the tumor vessel endothelium to induce neovascular sprouting without directly differentiating into endothelial cells in vivo (Beckermann et al., 2008). In addition to MSC-derived VEGF, in a human colorectal cancer xenograft model, TA-MSCs also produce IL-6 which in turn stimulates cancer cells to express endothelin 1 (ET-1) for angiogenesis (Huang et al., 2013).

Therefore, at the proliferation stage, TA-MSCs exert multifaceted functions to sustain tumor cell survival and proliferation. Several key MSC-regulating signaling pathways such as CXCL12/CXCR4, TGF β /Smad, IL-6 and VEGF could be potential candidates for developing novel MSC-targeting adjuvant therapeutics.

5.3. TA-MSCs in tissue remodeling

In the last stage of wound healing, host tissues including MSCs evolve self-restricting mechanisms to prevent over-proliferation of different types of tissue cells. Although such capabilities are largely dampened in the tumor microenvironment, MSC-mediated inhibition of epithelial tumor cell outgrowth, driving tumor cells to enter dormancy, as well as suppression of tumor angiogenesis, were indeed observed in

many types of cancers. In addition, in response to a hypoxic tumor environment, TA-MSCs modulate the primary tumor ECM, and support tumor cell invasion and metastasis into distant organs. Although such mechanisms are beneficial for the host to accomplish tissue remodeling in wound healing, they instigate cancer progression and metastasis.

5.3.1. TA-MSC-mediated suppression of tumor growth and angiogenesis

It is unclear whether TA-MSC-elicited tumor suppression is a reflection of the self-defending (antibacterial) capacity of MSCs in the early inflammation stage, or a self-restraining mechanism exerted in the tissue remodeling stage of wound healing. Studies of various hematopoietic and non-hematopoietic cancers suggest that TA-MSCs potentially inhibit tumor growth. Such inhibition was mediated by MSC-expressed TNF-related apoptosis-inducing ligand (TRAIL), dickkopf-related protein 3 (DKK-3), or cell-cell contact-dependent mechanisms causing tumor cell apoptosis or cell cycle arrest (Khakoo et al., 2006; Lee, Yoon, Reneau, & Prockop, 2012; Qiao et al., 2008; Ramasamy et al., 2007; Sun et al., 2009).

In human breast carcinoma xenograft models, TA-MSCs also serve to protect tumor cells from stress-induced cell apoptosis and facilitate proliferating tumor cells to enter dormancy and acquire cancer stem cell capabilities in vitro and in vivo. Such a cancer dormancy status is mainly achieved by MSC-released exosomes containing microRNAs such as miR-23b, 127, 197, 222, and 223 which repress the cell cycling regulatory genes, myristoylated alanine-rich C-kinase substrate (MARCKS) and CXCL12, as well as the cell cycle genes, cyclin-dependent kinase 4 (CDK4), cyclin D1 and p21WAF1 (Bliss et al., 2016; Lim et al., 2011; Ono et al., 2014). In an elegant three-dimensional MSC-human breast tumor cell co-culture system, MSCs were observed to be gradually internalized (cannibalized) by cancer cells and such a cannibalism led to the functional alteration of the cancer cells from active to dormant status in vitro and in vivo (Bartosh, Ullah, Zeitouni, Beaver, & Prockop, 2016). In addition to directly restraining the growth of epithelial tumor cells, TA-MSCs were also found to inhibit angiogenesis. In a syngeneic mouse melanoma model, co-administration of MSCs induced endothelial cell apoptosis through MSC-released ROS (Otsu et al., 2009). In a human glioma xenograft model, TA-MSCs suppressed endothelial cell growth by inhibiting the PDGF/PDGFR signaling axis (Ho et al., 2013). Taken together, the role of MSCs in limiting tumor growth could be a reflection of their modulatory effects to terminate the excessive tissue proliferation occurred in a regular wound healing process.

5.3.2. TA-MSCs accelerate tumor cell invasion and metastasis

At the remodeling stage, the final stage in wound healing, the originally disorganized collagen fibers synthesized during the proliferation stage undergo rearrangement, cross-linking and alignment, which are regulated by MMPs, TIMP3, TGF β and other cytokines (Xue & Jackson, 2015). In the tumor microenvironment, such a remodeling process is more complicated as the driving forces of remodeling are always affected by the earlier inflammation and proliferation processes, resulting in non-healing or over-healing wounds. TA-MSCs have been widely investigated for their potency to accelerate tumor cell invasion and metastasis through remodeling of the ECM (Kuhn & Tuan, 2010). Three mechanisms have been proposed in TA-MSC-elicited tumor cell invasion and metastasis, and are outlined below.

First, TA-MSCs secrete chemokines to direct tumor cell invasion. In syngeneic mouse melanoma and human breast carcinoma xenograft models, CCL-2, CCL-5, CCL-9 and CXCL10 are released from TA-MSCs to activate tumor cell migration and invasion which facilitate primary tumor cell metastases to lung, bone and lymph nodes (Chaturvedi, Gilkes, Wong, Zhang, et al., 2013; Karnoub et al., 2007; Kudo-Saito et al., 2013; Luo et al., 2014; Swamydas, Ricci, Rego, & Dreau, 2013). As certain chemokines were shown to activate MMPs, the MSC-derived chemokines may also play a role in MMP-mediated ECM remodeling (Swamydas et al., 2013).

Second, TA-MSCs serve as potent drivers for tumor cells to undergo an epithelial-mesenchymal transition (EMT) which in turn stimulates the invasion and metastasis of tumor cells. In a human breast cancer xenograft model, MSCs induced breast tumor cells to undergo EMT via a LOX-Twist signaling pathway. This, in turn, resulted in an elevated capability for tumor cells to metastasize to lung and bone (El-Haibi et al., 2012). Direct contact between MSCs and cancer cells is regarded as a mechanism for upregulation of the EMT-related genes (Martin et al., 2010; Takigawa et al., 2017). In human cancer specimens, the typical EMT markers were found to be expressed at higher levels at the stroma-epithelial invasive edge (Takigawa et al., 2017).

Third, in human breast cancer xenograft and tongue cancer models, human MSCs were shown to modulate collagen deposition through their expression of collagen receptor discoidin domain receptor 2 (DDR2), or to increase collagen I expression in cancer cells. Such ECM modulation further favors tumor cell migration and invasion thus supporting metastases (Gonzalez et al., 2017; Salo et al., 2013).

Therefore, TA-MSCs impede the outgrowth of the primary tumors in the tumor “wound” remodeling stage. On the other hand, through secretion of chemokines and cytokines, TA-MSCs potently drive the primary tumor cells to undergo EMT and metastasize into distant organs.

5.4. TA-MSCs and cancer therapeutic resistance

MSCs were recently identified as a key tumor microenvironmental component that elicits tumor cell resistance to various therapies (Houthuijzen, Daenen, Roodhart, & Voest, 2012). Such a capability is mainly attributed to the pro-survival and cancer stem cell-promoting effects of TA-MSCs, as mentioned above. Following chemotherapy, MSCs are capable of retaining their “stemness” characteristics, proliferative rate and differentiation potential, in part because of their elevated apoptotic threshold (Mueller et al., 2006). Similarly, MSCs were also shown to be radioresistant since they could develop multiple, atypical DNA damage response mechanisms to offset effects of radiation-induced DNA damage (Sugrue, Brown, Lowndes, & Ceredig, 2013).

MSCs themselves are relatively resistant to conventional cancer therapies compared to tumor cells, and more importantly, they can further protect tumor cells from therapy-induced cytotoxicity. In a chronic lymphocytic leukemia model, TA-MSCs were shown to directly interact with leukemia cells to prevent drug-induced cleavage of myeloid cell leukemia 1 (Mcl-1) and poly ADP ribose polymerase (Parp) resulting in drug resistance (Kurtova et al., 2009). In a human ovarian cancer model, a paracrine hedgehog-BMP4 positive feedback loop between ovarian cancer cells and TA-MSCs was defined as the mechanism for drug resistance (Coffman et al., 2016). In response to treatment with the chemotherapeutic drug cisplatin, TA-MSCs are induced to produce certain fatty acids and the cytokines IL-6 and IL-8 leading to drug resistance in a mouse colon cancer model and a human breast cancer model, respectively (Roodhart et al., 2011; Skolekova et al., 2016). Although there is little experimental evidence showing MSCs directly participate in resistance to the currently promising immunotherapeutics particularly the immune checkpoint blockade, the mesenchymal lineage cells such as myofibroblasts have been revealed to mediate immunotherapy failure via autocrine and paracrine TGF β signaling and their secretion of CXCL12, MMP9 and ECM proteins in multiple pre-clinical tumor models (Chakravarthy, Khan, Bensler, Bose, & De Carvalho, 2018; Feig et al., 2013; Mariathasan et al., 2018; Zhao et al., 2018). Moreover, when epithelial tumor cells undergo EMT, their “mesenchymal” status favors the development of resistance to immune checkpoint blockade in treatment of breast tumors in mice (Dongre et al., 2017). In human cancer patients, the EMT signature is associated with unique tumor microenvironment and therefore could be a potential biomarker for selecting patients who will benefit from immunotherapeutics (Lou et al., 2016; Mak et al., 2016; Wang et al., 2018).

Therefore, TA-MSCs are highly anti-apoptotic and can further help tumor cells to evade different types of cancer therapeutics. A deeper

understanding of the robust pro-survival potency of TA-MSCs, as well as the mechanisms underlying MSC-promoted tumor cell dormancy, will benefit development of MSC-targeting approaches to overcome therapeutic resistance and prevent cancer recurrence.

6. Concluding remarks

Considering tumors as an unresolvable wound, the diverse functions of MSCs in the three stages of regular wound healing and cancer progression are compared (Table 1). MSCs indeed play similar roles in cancer progression, as they do in wound healing, however, such beneficial effects to repair a wound are fully utilized by the tumor cells for further progression and evolution of resistance towards different therapeutics. Increased knowledge of the biology of MSCs in tissue regeneration would highly accelerate a deeper and more comprehensive understanding of the contribution of MSCs to cancer progression. Several questions remain to be answered to achieve the goal of targeting MSCs and mesenchymal lineages as new strategies in clinical cancer research.

First, the tumor modulating effects mediated by TA-MSCs are largely controversial. The conflicting results could be due to the variance in MSC cell isolation and culture maintenance methods, MSC cell passages, species, cancer models utilized, adoptive cell transplantation time, and doses or duration. From the analysis of the function of MSCs in both wound healing and cancer contexts, it is clear that MSCs are a type of regulatory cell and can function by either enhancing or suppressing distinct steps of the healing process. For example, MSCs can be potentially immunosuppressive, but can also enhance inflammation depending on the immune environment in which they reside. Moreover, MSCs are able to both accelerate and inhibit epithelial cell and endothelial cell growth at the proliferation stage and tissue remodeling stage, respectively. In accordance, many feedback loops have been identified in studies of MSCs in both wound healing and cancer (Caplan, 2017; Le Blanc & Davies, 2015; Prockop, 2013; Wang et al., 2014). The functions of MSCs are usually “licensed” upon activation by external stimuli (Krampera, 2011), and MSCs have been proposed to possess an MSC1 or MSC2 status based upon their activation by different TLR ligands (Waterman, Tomchuck, Henkle, & Betancourt, 2010). Such concepts would inspire us to further deliberate the endogenous functions of MSCs in a systematic and dynamic view in the growth, metastasis and drug resistance of various cancers and in other disease processes.

Second, there is currently a lack of well-accepted standards for MSC identification particularly for the identification of endogenous MSCs (da Silva Meirelles, Caplan, & Nardi, 2008; Sacchetti et al., 2016). The distinctions among MSCs, perivascular cells (pericytes) and fibroblasts are still ambiguous (Guimaraes-Camboa et al., 2017; Murphy, Moncivais, & Caplan, 2013). What has been observed or concluded thus far may be from studies of mixed types of mesenchymal cells in vitro and in vivo. Furthermore, the research in the field has mainly relied on exogenous MSC adoptive transfer or ex vivo systems in which conclusions may be less physiologically relevant. Recent efforts have been made to explore endogenous MSCs in pathological contexts, such as fibrotic diseases (El Agha et al., 2017). In the future, well-recognized endogenous markers and transcriptional factors for endogenous MSCs need to be tested and established in both human and experimental animals.

Lastly, it is time to consider harnessing the properties of MSCs for cancer therapy (Marofi, Vahedi, Biglari, Esmailzadeh, & Athari, 2017). There are already a few pre-clinical studies in which the homing capacity MSCs to tumors is exploited to utilize MSCs as an excellent in vivo vehicle for delivering tumoricidal agents such as IFN α , IFN β and TRAIL (Shah, 2012; Stuckey & Shah, 2014). In consideration of the complicated roles of MSCs in cancer progression, especially their function to support tumor dormancy (Bartosh et al., 2016), a more careful and specific tailoring of engineered MSCs need to be conducted for designing future clinical application. Approaches to insert certain suicide genes or genetic modulation to deplete known tumor-promoting genes in engineered MSCs would help develop safer and more efficacious cell-

Table 1
MSCs in wound healing responses to injury and cancer

		Wound healing response to injury		Wound healing response to Cancer	
		Functions	Mechanisms	Functions	Mechanisms
Blood clotting		MSCs promote blood clotting	TFs; FVIII	Not known	
Anti-bacterial		MSCs are self-defending by elimination of the invading bacteria	Secretion of anti-microbial peptides LL-37, hepcidin, β -defensin 2 and lipocalin-2 etc.	It is unclear whether the tumor-suppressive effects of MSCs are related to such a self-defending capacity by MSCs	
Inflammation	Myeloid cell recruitment	MSCs drive myeloid cell migration from BM to inflamed tissues	Chemokines CCL-2, CCL-3, CCL-4, CXCL1, IL-8, MIF; adhesion molecules ICAM-1, etc.	TA-MSCs recruit monocytes, macrophages and neutrophils to the tumor microenvironment	Chemokines and cytokines CCL-2, CCL-7, CCL-12, CXCL1, CXCL2, CSF1 etc.
	Myeloid cell and other innate immune cell suppression	MSCs suppress myeloid cell functions upon stimulation by TNF α , IL-1 and ROS	COX-2, PGE2, TSG-6, SOD3, etc.	TA-MSCs polarize the M1 macrophages to an M2 phenotype; TA-MSCs also stimulate MDSC differentiation	CXCL3, HGF etc.
	Myeloid cell reprogramming	MSCs convert M1 macrophages to an M2-like type	PGE2, TSG-6, etc.		
	T cell recruitment	MSCs augment T-cell infiltration and activities in a low-level inflammatory environment	Chemokines CXCL9, 10, 11; adhesion molecules ICAM-1, VCAM-1	Not known	
	T cell suppression	MSCs exert a robust suppression of T-cells upon stimulation by IFN γ together with other inflammatory cytokines	NO, IDO, PGE2, PD-L1, HO-1, LIF, IL-6, galectin 1, FasL, TGF β , Treg, etc.	MSCs suppress anti-tumor immunity	NO, IDO, Treg, regulatory CD8 ⁺ T cells, etc.
		Wound healing response to injury		Wound healing response to Cancer	
		Functions	Mechanisms	Functions	Mechanisms
Proliferation	MSC homing	MSCs efficiently home to tissue injury sites	Chemokines CXCL12, CCL-2, CCL27 and CCL21; VCAM-1; MMPs; etc.	Exogenously implanted and endogenous MSCs efficiently migrate into the tumor environment	CXCL12, CCL-2, CCL-25, CXCL16, MIF, IL-6, LL-37 etc.
	MSC differentiation and fibroplasia	MSCs can be differentiated into osteoblasts, adipocytes and myofibroblasts at the injury sites	TGF β , BMPs, Notch, Wnt, Hedgehogs, etc.	BM-MSCs or adjacent tissue-derived MSCs give rise to distinct types of tumor-associated myofibroblasts	TGF β etc.
	Re-epithelialization	MSCs promote epithelial cell growth	Growth factors EGF, HGF, KGF, PDGF, IGF etc.	TA-MSCs support survival of cancer stem cells and proliferation of tumor cells	IL-6, IL-8, CXCL1, CXCL7, BMPs, miRNAs
	Angiogenesis	MSCs enhance angiogenesis	Pro-angiogenic factors VEGF, FGF, PDGF, TGF β , etc.	TA-MSCs facilitate tumor-associated angiogenesis	VEGF, IL-6, etc.
Tissue remodeling	Inhibition of proliferation	MSCs inhibit tissue cell proliferation	TGF β , cell-cell contact, MMPs	TA-MSCs suppress epithelial tumor growth but drive them to enter dormancy	TRAIL, DKK-3, cell-cell contact, cannibalization, miR-23b, 127, 197, 222, 223
	Inhibition of angiogenesis	MSCs suppress angiogenesis during tissue remodeling	TIMPs, MMPs	TA-MSCs can inhibit angiogenesis	ROS, inhibition of PDGF signaling
	Collagen rearrangement	MSCs promote ECM remodeling	MMPs, TIMPs, TGF β	TA-MSCs modulate collagen organization	DDR2, MMPs
	Tissue cell invasion	MSCs stimulate tissue cell invasion	MMPs, collagen rearrangement	TA-MSCs accelerate tumor cell invasion and metastasis	Chemokines CCL-2, CCL-5, CCL-9, CXCL10 etc.; TA-MSCs-induced EMT

Abbreviations: TFs, Tissue factors; FVIII, Factor VIII; MIF, Macrophage migration inhibitory factor; ICAM-1, Intercellular adhesion molecule 1; CCL, CC motif chemokine ligand; CXCL, C-X-C motif chemokine ligand; IL, Interleukin; CSF1, Colony stimulating factor 1; COX-2, Cyclooxygenase-2; PGE2, Prostaglandin E2; TSG-6, Tumor necrosis factor (TNF)-stimulated gene-6; SOD3, Superoxide dismutase 3; HGF, Hepatocyte growth factor; VCAM-1, Vascular cell adhesion molecule 1; NO, Nitric oxide; IDO, Indoleamine 2,3-dioxygenase; PD-L1, Programmed death-ligand 1; HO-1, Heme oxygenase 1; LIF, Leukemia inhibitory factor; FasL, Fas ligand; Treg, Regulatory T cell; EGF, Epidermal growth factor; KGF, Keratinocyte growth factor; PDGF, Platelet-derived growth factor; IGF, Insulin-like growth factor; BMP, Bone morphogenetic protein; miRNA, microRNA; VEGF, Vascular endothelial growth factor; FGF, Fibroblast growth factor; PDGF, Platelet-derived growth factor; TGF β , Transforming growth factor beta; MMP, Matrix metalloproteinase; TRAIL, TNF-related apoptosis-inducing ligand; DKK-3, Dickkopf WNT Signaling Pathway Inhibitor 3; TIMP, Tissue inhibitor of metalloproteinase; ROS, Reactive oxygen species; DDR2, Discoidin domain receptor 2; EMT, Epithelial-mesenchymal transition.

based cancer therapeutics. Furthermore, following the concept of vascular normalization in cancer treatment (Goel et al., 2011), new avenues for conversion of malignancy-facilitating MSCs back to their normal malignancy-inhibitory status, could be a promising direction for future MSC-targeting cancer research. Such an MSC or mesenchymal normalization may also benefit the promising vascular normalization strategies owing to the essential vascular gatekeeper role of endogenous MSCs.

Overall, a deeper exploration of MSC biology from a regenerative perspective will expand our understanding of the multiple functions of MSCs in cancer progression and response to therapy, thereby supporting development of more precise MSC targeted therapy for clinical translation.

Conflict of interest statement

The authors declare that there are no conflicts of interest.

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