

The pro-metastatic role of bone marrow-derived cells: a focus on MSCs and regulatory T cells

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Several bone marrow-derived cells have been shown to promote tumour growth and progression. These cells can home to the primary tumour and become active components of the tumour microenvironment. Recent studies have also identified bone marrow-derived cells—such as mesenchymal stem cells and regulatory T cells—as contributors to cancer metastasis. The innate versatility of these cells provides diverse functional aid to promote malignancy, ranging from structural support to signal-mediated suppression of the host immune response. Here, we review the role of mesenchymal stem cells and regulatory T cells in cancer metastasis. A better understanding of the bipolar nature of these bone marrow-derived cells in physiological and malignant contexts could pave the way for new therapeutics against metastatic disease.

Keywords: bone marrow-derived cells; cancer metastasis; mesenchymal stem cells; regulatory T cells

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See the Glossary for abbreviations used in this article.

Introduction

Cancer metastasis accounts for over 90% of mortality from solid malignancies and is a multi-step process that allows primary tumour cells to escape from the site of origin and colonize distant organs [1,2]. The metastatic process has traditionally been classified into several distinct stages—local invasion, intravasation, survival in the circulation, extravasation and colonization—each of which is regulated intrinsically by the tumour cell and extrinsically by the surrounding stroma [3–5]. The complexity of this multi-stage process has enraptured many generations of cancer researchers, starting with Stephen Paget's century-old 'seed and soil' hypothesis that compared tumour cells to 'seeds' that are systematically distributed, but only inhabit particular environments, or 'soils,' which are supportive of their sustained growth [6]. Although distinct genetic profiles of tumour cells with particular proclivities to colonize specific distant organs have begun to be elucidated, how such metastasis genes exert

their functions in the context of tumour–stroma interactions remains a major topic of ongoing research [3,7–9].

Throughout the expansion of the primary tumour, a vast array of host cells—ranging from macrophages to fibroblasts—create and sustain a favourable microenvironment for malignant growth [4,10]. Bone marrow-derived cells, including mesenchymal stem cells (MSCs) [11–14] and immunosuppressive cells, such as regulatory T cells (T_{regs}) [15–17], have been identified as major components of the primary tumour microenvironment. More recently, new evidence corroborates the contribution of these populations in the metastatic process [18–20], largely by providing cell motility-inducing factors and promoting a protective microenvironment for tumour cells throughout their journey to distant organs. These studies therefore reinforce the idea that the stromal components of the tumour microenvironment can play an active role in promoting cancer metastasis. However, given the complexity and multi-faceted progression of tumour metastasis, an in-depth analysis of these metastasis-promoting interactions is still incomplete. In addition, the function of bone marrow-derived cells in modulating the immune system to promote metastasis must be viewed in the context of the general controversy surrounding immunosurveillance of tumour cells [21–23], both in primary tumour growth and in cancer metastasis. Despite such challenges, numerous studies have implicated several immune cells, such as immature myeloid cells [24–26], mast cells [27,28], macrophages [29,30], platelets [31,32], neutrophils [33], and haematopoietic and endothelial progenitor cells [34–36] to promote cancer metastasis, and have already been addressed in other reviews [37–40]. Here, we review our understanding of the contribution of MSCs and T_{regs} to cancer metastasis and discuss their potential as therapeutic targets in the treatment of metastatic disease.

Tumour–MSC crosstalk in metastatic progression

The bone marrow stroma, which is the master haematopoietic compartment, is composed of various cell types and crucial regulators required for creating an ideal niche for the maintenance of haematopoietic stem and progenitor cells [41–43]. A dominant subpopulation in the bone marrow stroma is believed to be mesenchymal in nature [44–46], with a primitive population of MSCs able to differentiate into osteoblasts, adipocytes and chondrocytes [47,48]. MSCs have also been reported to differentiate into fibroblasts [49–51] and pericytes [52,53], although studies showing similar differentiation potential of dermal pericytes [54], retinal pericytes [55] and primary

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fibroblast-like populations [56] suggest that both fibroblasts and pericytes might not be terminally differentiated progenies.

Despite numerous studies characterizing the versatile differentiation potential of MSCs [57–59], a general consensus is lacking on the immunophenotypic markers used to isolate this multi-potential population [49,60]. A broad array of MSC markers have been used in various studies in both human and mouse models, including CD73, CD90, CD105, CD140, CD146 and Sca1, and the clonogenic purity of the MSC population has been proposed to vary depending on which combinations of these markers are used for isolation [52,53,61–63]. Given that a multipotent cell able to differentiate into osteoblasts, adipocytes and chondrocytes meets the criteria of an MSC [61] and given the extremely heterogeneous nature of this population, it is essential to evaluate studies on MSCs in a context-dependent manner. It is also important to note that the bone marrow, although providing a significant proportion, is not the only source of MSCs. Adipose tissue [64,65] and the umbilical cord [66,67] have been shown to create a niche for MSCs, which can also be recruited to sites of wound healing and the primary tumour. These tissue-specific MSCs meet the tri-lineage criterion mentioned above, but multiple studies show varying differentiation propensity toward specific lineages and plasticity depending on the source of MSC isolation [68–71]. For example, adipose tissue-derived MSCs, otherwise referred to as adipose-derived stem cells (ADSCs), can differentiate into osteoblasts [72], adipocytes, chondrocytes [73,74], myoblasts [75] and even endothelial cells [76,77], but they have also been shown to arise from mature human adipocytes through a process known as dedifferentiation [78,79]. These reprogrammed multipotential cells, commonly referred to as DFAT cells, acquire the surface markers of ADSCs and have the capacity to differentiate into osteoblasts, chondrocytes and adipocytes *in vitro* [79,80]. However, these findings are still contested, as conflicting results can be obtained depending on the culture conditions and immunophenotypic markers used for isolation, emphasizing the need for *in vivo* lineage tracing studies of these versatile stem cells. The plasticity of MSCs to transdifferentiate in various settings is crucial for their normal physiological functions, but also allows them to have a more ominous role in cancer. For example, one of the major functions of MSCs is their differentiation into connective tissue during wound healing [81–83] and secretion of a battery of growth-modulating factors, such as IL-6 [11,84] and Ang1 [53], to promote accelerated regeneration of injured tissue. In cancer, this process can create a growth-promoting tumour microenvironment in which MSCs differentiate into cancer-associated fibroblasts (CAFs) [14,85,86], which secrete IL-6 [87] and VEGF [88] to promote tumour angiogenesis. In this context, MSCs act in sharp contrast to their known endogenous functionality by exacerbating what has been described as “a wound that never heals” [89].

Although the contribution of MSCs to primary tumour growth has been studied extensively in a variety of cancers [11], including breast cancer [13], colon cancer [90] and lymphoma [91], studies demonstrating their potential to promote tumour metastasis are relatively rare in comparison. Nevertheless, MSCs have been recently shown to facilitate tumour metastasis by secreting inflammatory cytokines to promote cell motility (Fig 1). Subcutaneous co-injection of the human breast cancer cell line MDA-MB-231 and human bone marrow-derived cells into immunodeficient mice significantly enhances lung and liver metastasis [13,18]. These studies identify

Glossary

Ang1	angiopoietin 1
Ap1	activator protein 1
β-GBP	β-galactoside-binding protein
CCL	CC chemokine ligand
CCR	CC chemokine receptor
CTL	cytotoxic T lymphocyte
CTLA4	cytotoxic T-lymphocyte antigen 4
DFAT	dedifferentiated fat
ECM	extracellular matrix
EGF	epidermal growth factor
ErbB2	v-erb-b2 erythroblastic leukaemia viral oncogene homologue 2
FOXP3	forkhead box P3
FSP1	fibroblast-specific protein 1
HGF	hepatocyte growth factor
HLA	human leukocyte antigen
HLA-G5	human leukocyte antigen-G5
IFN-γ	interferon-γ
IL	interleukin
IL-17BR	IL-17B receptor
MCA	3-methylcholanthrene
M-CSF	macrophage colony-stimulating factor
MMP	matrix metalloproteinases
NK	natural killer
NOD/SCID	nonobese diabetic/severe combined immunodeficient
PGE2	prostaglandin E2
Rag	recombination-activation gene
RANK(L)	receptor activator of nuclear factor-κB (ligand)
Sca1	stem cell antigen 1
SDF1	stromal-derived factor 1
shRNA	short hairpin RNA
SMA	smooth muscle actin
TGF-β	transforming growth factor-β
T _H 2 cell	T helper 2 cell
TNF-α	tumour necrosis factor-α
TRAIL	tumour necrosis factor-related apoptosis-inducing ligand
VEGF(R)	vascular endothelial growth factor (receptor)
VLA4	very late antigen 4

CCL5 as an MSC-derived metastasis-promoting factor, the expression of which increases after the interaction of MSCs with cancer cells (Fig 1). CCL5-induced Akt activation allows tumour cells to extravasate from the circulation to colonize distal organs, and thereby significantly increases metastatic potential [13]. Tumour-derived osteopontin (OPN) was also recently found to promote CCL5 secretion from MSCs by stimulating the binding of c-Jun homodimers to the CCL5 promoter and, thus, stimulating its transactivation (Fig 1; [18]). Aptamer-mediated neutralization of circulating OPN in MDA-MB-231 xenograft models reduces serum CCL5 levels, as well as lung and liver metastasis. In this model, the contribution of CCL5 to metastasis does not seem to depend on its effect on the primary tumour stroma, as there is no difference in infiltrating macrophages, angiogenesis or SMA-positive stromal cells between tumours initiated by CCL5 overexpressing MDA-MB-231 cells and those by control MDA-MB-231 cells [13].

In addition to promoting tumour-intrinsic metastatic properties, the tumour–MSC interaction can modulate the stromal microenvironment to promote metastasis [18]. Tumour cell-derived OPN promotes the expression of cancer-associated fibroblast markers—such as αSMA, tenascin-C, SDF1 and FSP1—in human MSCs (Fig 1). Interestingly, the expression of these markers is enhanced in

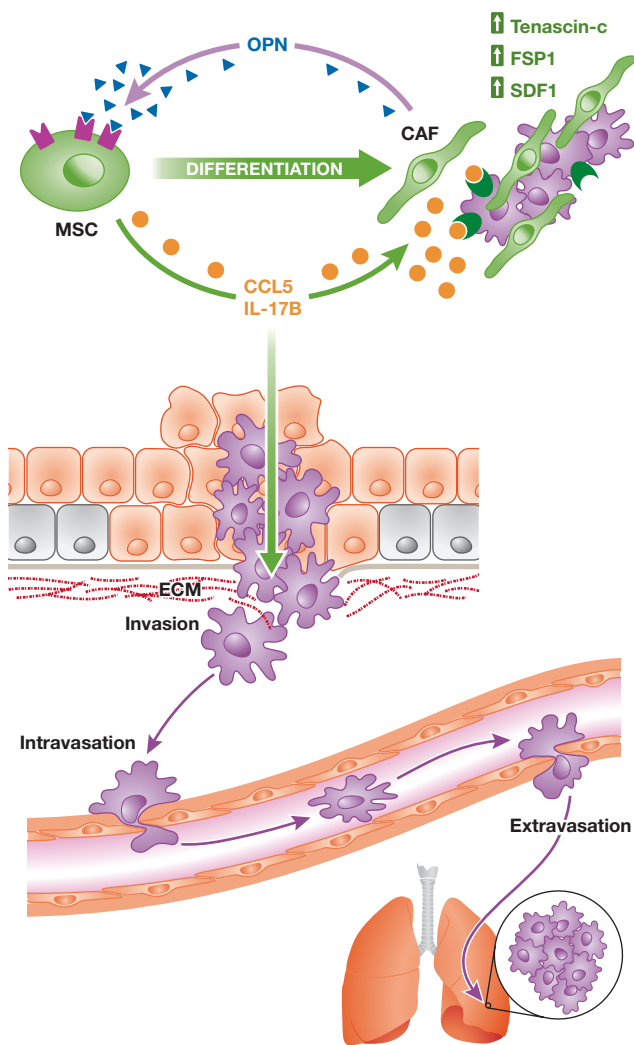


Fig 1 | Mesenchymal stem cells promote cancer metastasis. MSCs can secrete inflammatory cytokines—such as CCL5 and IL-17B—to facilitate cell motility, which is necessary for tumour cell invasion through the surrounding ECM, intravasation into blood vessels and extravasation from the circulation at target sites. In addition, MSCs can differentiate into CAFs, which express tenascin-C, FSP1 and SDF1 to support tumour progression and create a pro-metastatic microenvironment. Feedback mechanisms sustain these pro-metastatic effects of MSCs on tumour cells, as is the case of tumour-secreted OPN promoting both CCL5 transcription and CAF marker expression. Such MSC–tumour cell interactions have been observed both in the primary tumour as well as at the sites of metastases. CAF, cancer-associated fibroblast; CCL, CC chemokine ligand; ECM, extracellular matrix; FSP1, fibroblast-specific protein 1; IL, interleukin; MSC, mesenchymal stem cell; OPN, osteopontin; SDF1, stromal-derived factor 1.

MSCs isolated from the metastatic site, suggesting another mechanism whereby OPN contributes to cancer metastasis. Human and mouse MSCs have been shown to transdifferentiate into CAFs when co-injected with Skov3 ovarian cancer cells [14] or with MKN45 gastric cancer cells [87]. In addition to secreting tumour growth-promoting factors such as EGF and IL-6, these MSC-derived CAFs also express extracellular matrix- and angiogenesis-regulating proteins to create a metastasis-promoting stromal microenvironment

for the primary tumour. Human MSC-derived CAFs are also found at the sites of metastasis, have similar characteristics as in the primary tumour [18,92,93] and probably create a fostering environment for tumour cells—the metastatic niche'. In this regard, it is of particular interest that tenascin-C, which is derived from cancer cells and myofibroblasts, has been recently shown to generate a metastatic niche to facilitate lung metastasis [94]. As noted above, tenascin-C expression is also increased in OPN-stimulated human MSCs.

CAFs are thought to be one of the most important and aggressive supporters of tumour growth and invasion [95–100], but they have also been recently implicated in facilitating metastasis after the cancer cells have entered the circulation [101,102]. Periostin—a crucial extracellular component in bone and heart formation—is upregulated in α SMA⁺ vimentin⁺ myofibroblasts in metastatic lungs [103]. Periostin deficiency in spontaneous mammary tumour-bearing mice significantly reduces pulmonary metastasis and is thought to disrupt the putative cancer stem cell niche at the metastatic site. Despite such similarities in the characteristics between the MSC-derived CAFs that have been recently identified and CAFs of various sources, it remains to be determined whether MSCs actively promote metastasis as a differentiated CAF at the metastatic site (Sidebar A).

The emerging theory of the pre-metastatic niche proposes that target organs can be primed by secreted factors from primary tumours to create a more accommodating microenvironment before the arrival of metastatic tumour cells [25,104,105]. For example, the accumulation of VEGFR1-positive myeloid progenitors in pre-metastatic lungs creates favourable docking sites for lung carcinoma tumour cells [25]. Tumour-secreted factors, such as placental growth factor (PlGF), are transmitted from the primary tumour to the metastatic organ, causing resident fibroblasts to produce fibronectin at future sites of metastasis. The secreted factors also stimulate the recruitment of bone marrow-derived progenitor cells that express VEGFR1 and integrin VLA4, which is a fibronectin receptor [25]. Although this study showed that VEGFR1 activity is crucial for metastasis, another report claims that VEGFR1 deficiency in a genetic model that lacks the tyrosine kinase domain of VEGFR1 does not affect spontaneous metastasis of Lewis lung carcinoma and melanoma [106]. This aspect of CAF targeting clearly requires further study, as does whether the resident fibroblasts of the pre-metastatic niche arise from local MSCs or from the bone marrow. In addition, although MSCs isolated from sites of metastasis seem to have stronger CAF-associated marker expression than those in the primary tumour, whether and how metastatic MSCs differentially contribute to primary and secondary tumorigenesis and whether MSC-derived CAFs precede tumour seeding to facilitate the colonization process remain open questions (Sidebar A).

Another molecule recently implicated in promoting metastasis through an MSC-dependent mechanism is the pro-inflammatory cytokine IL-17B [107], which is secreted by human MSCs that migrate to the primary tumour in a TGF- β -dependent manner [93]. The IL-17BR is a prognostic indicator associated with invasive tumour progression [108,109], and its ectopic expression in MDA-MB-231 and SUM1315 breast cancer cells leads to increased migration *in vitro*, as well as increased frequency of lung and liver metastases *in vivo* (Fig 1). IL-17BR expression is significantly higher in metastatic bone lesions of SUM1315 cells and is thought to promote organ-specific metastasis to the bone, although ectopic IL-17BR alone is not sufficient to promote metastasis [93]. The IL-17 family consists of six structurally related members, which

are produced by and act on various immune cells [110,111] to regulate immune function in inflammation, autoimmunity, host defence against bacterial and fungal infections and tumorigenesis [107,112]. IL-17A⁺ immune cells in the tumour microenvironment have been shown to promote tumour growth by increasing the production of pro-angiogenic factors—such as VEGF and prostaglandins [113,114]—and recruiting neutrophils to the primary tumour through the production of IL-8 [115,116]. Interestingly, IL-17A reduces TNF-induced CCL5 expression in mouse lung fibroblasts [117], whereas IL-17E, which binds to IL-17BR, has the opposite effect [118]. Although the roles of IL-17A and E in tumorigenesis have been characterized in numerous studies [119–121], the function of IL-17B in both tumorigenesis and immunity remains largely unknown. Both IL-17B and C cause neutrophil infiltration by the induction of TNF and IL-1 β expression in monocytes [118,119]. In the light of the physiological role of IL-17B, IL-17B-producing MSCs could facilitate the metastatic process by promoting tumour angiogenesis and recruiting neutrophils to the metastatic site. This would support β_2 -integrin-mediated docking of the tumour cells to the endothelial wall [122].

Notably, there are significant variations in how tumour cells use MSCs to enhance their metastatic potential. Although MDA-MB-231 and SUM1315 are both innately metastatic to the bone and can be induced to metastasize to the liver and lungs upon ectopic expression of IL-17BR, only SUM1315 cells express significantly higher levels of IL-17BR in bone metastasis. shRNA-mediated knockdown of CCR5 in MCF7/Ras and HMLER cells does not affect metastatic potential, whereas lung metastasis is seriously compromised in MDA-MB-231 and MDA-MB-435 cell lines under these conditions. In addition, the metastasis-promoting potential of MSCs can be independent of their effect on primary tumour growth, as is the case in the enhancement of MCF7/Ras metastasis by MSC admixture [13]. By contrast, MSCs apparently inhibit hepatocellular carcinoma metastasis by suppressing TGF- β and MMP2, although they promote primary tumour growth *in vivo* [123]. Therefore, further investigation into how various cancer subtypes alter their metastatic potential in response to MSCs and whether these interactions are tumour type-dependent and context-dependent will be crucial for developing new therapeutic strategies to block the metastasis-promoting function of MSCs (Sidebar A).

MSC-mediated immunomodulation in metastasis

The contribution of MSCs to tumour growth and progression is similar to that in wound healing; both processes benefit from physical support of the stroma provided by MSCs with their versatile plasticity. However, one major difference between the endogenous role of MSCs healing a wound and their tumorigenic role at the ‘never-healing’ wound site is their influence on immune cells—the former is immunostimulatory whereas the latter is generally immunosuppressive [124]. Human MSCs suppress the proliferation of T cells through mediators such as TGF- β and HGF [125], and induce apoptosis of activated T cells through indoleamine 2,3-dioxygenase secretion [126]. MSCs also interfere with dendritic cell differentiation [127–129], which is crucial for the production of pro-inflammatory cytokines—including TNF- α , IFN- γ and IL-12—as well as B- and NK-cell proliferation [130,131]. This markedly hampers the normal inflammatory function of immune cells (Fig 2). The MSC-driven promotion of T_{reg} cell expansion and immunosuppressive activity is of particular interest, as many reports suggest that T_{regs} are a prognostic factor in

Sidebar A | In need of answers

- (i) How are tumour-associated MSCs, T_{regs} and other bone marrow-derived cells different from those present in normal tissues?
- (ii) Do bone marrow-derived cells have a role in a particular stage of metastasis, distinct from other stages?
- (iii) What is the nature of tumour metastatic niches and how do they differ from normal adult stem cell niches, such as the haematopoietic stem cell niche? What are the roles of the various bone marrow-derived cells in such niches?
- (iv) How can we interfere with the interaction of bone marrow-derived cells with tumour cells to inhibit tumour metastasis?

a variety of cancers (discussed later). These FoxP3⁺ CD25^{hi} CD4⁺ cells expand after stimulation by MSC-derived HLA-G5 [132] and can maintain their immunosuppressive activity for an extended period of time when cultured with MSCs *in vitro* [133]. Most of the aforementioned studies have led to promising results in the use of MSCs to treat graft-versus-host disease, allogeneic bone marrow transplantations and skin transplantation [124,134].

The immunomodulatory properties of MSCs have also been proposed to have an influential role in tumour growth, although evidence for this is limited. Injection of MSCs with B16 melanoma cells into syngeneic recipients allows tumour growth, whereas the tumour cells are rejected in the absence of MSCs or if their numbers are reduced [135]. More recently, MSC-secreted TGF- β 1 has been shown to promote T_{reg} cell expansion, and blocking TGF- β 1 or depleting T_{regs} has been shown to increase CTL- and NK-mediated lysis of T47D breast cancer cells *in vitro* [136]. Finally, CAFs polarize the tumour microenvironment to an immunosuppressive T_H2 cytokine profile, which can stimulate T_{reg} expansion and promote 4T1 tumour growth and metastasis to the lungs [137]. This suggests that differentiated progenies of MSCs might also contribute to immunosuppression. Although these studies confirm that tumour-associated MSCs can exert immunosuppression, they do not provide unequivocal evidence that MSC-mediated immunosuppression can promote metastatic tumour growth independently of other MSC-dependent mechanisms discussed above.

The prospect of targeting MSCs to render the tumour microenvironment more immunoreactive has been discussed but must be considered in the light of the controversy surrounding tumour immunosurveillance. Burnet and Thomas proposed in 1957 that tumour cells with their ‘new antigenic potentialities’ can induce an immune response against the primary tumour; successful tumorigenesis occurs if the tumour cells are able to escape recognition by the innate and adaptive immune systems [138–140]. Deficiency of Rag [141], the depletion of CD4⁺ and CD8⁺ T cells, and neutralization of IFN- γ in MCA-treated mice [142] have all been shown to significantly increase the incidence of tumorigenesis, supporting the immunosurveillance hypothesis. However, other studies have shown that immunoeediting—a process by which tumour cells can lose their immunogenicity through loss of the major histocompatibility complex [138]—is not observed in other strains of spontaneous cancer-prone mice. For example, tumours spontaneously arising from simian virus 40 T antigen (SV40 Tag) can elicit a Tag-specific B-cell and T-cell response, but this response cannot suppress tumorigenesis [23]. In fact, some key studies have demonstrated a pro-tumorigenic role for adaptive immune cells [143,144]. CD4⁺ effector T cells, which secrete cytokines to support CTL proliferation, also secrete cytokines such as

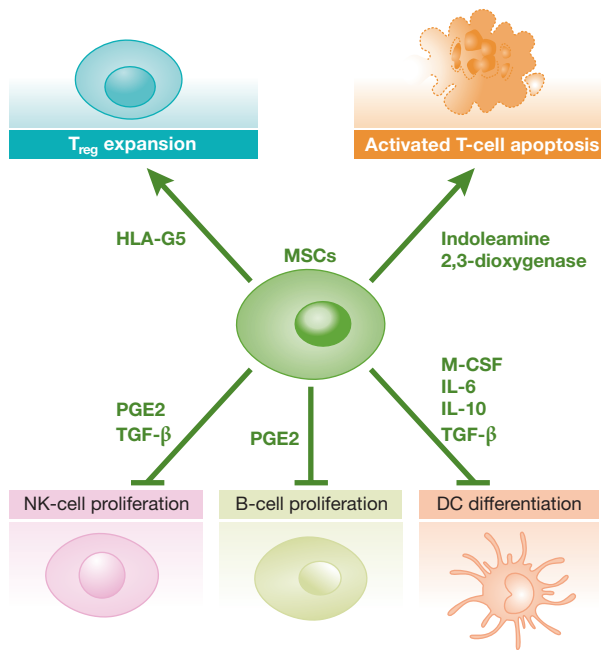


Fig 2 | Immunomodulatory properties of mesenchymal stem cells. MSCs can secrete a variety of immunomodulatory factors to promote an immunosuppressive environment. They induce apoptosis of activated T cells and the expansion of immunosuppressive T_{regs} through the secretion of indoleamine 2,3-dioxygenase and HLA-G5, respectively. MSCs can also produce PGE2 and TGF- β to inhibit NK-cell proliferation, as well as secrete PGE2 to inhibit B-cell proliferation. MSC secretion of M-CSF, IL-6, IL-10 and TGF- β also suppresses DC differentiation. DC, dendritic cell; HLA-G5, human leukocyte antigen G5; IL, interleukin; M-CSF, macrophage colony-stimulating factor; MSC, mesenchymal stem cell; NK, natural killer; PGE2, prostaglandin E2; T_{reg} cell, regulatory T cell; TGF- β , transforming growth factor- β .

IL-4 that stimulate EGF production from tumour-associated macrophages (TAMs) in a spontaneous MMTV-PyMT murine mammary adenocarcinoma model. This, in turn, promotes EGF receptor-mediated invasion and subsequent metastasis to the lungs [144]. In line with these reports, CCL5 and IL-17B—which have been discussed above as mediators of cancer metastasis—are inflammatory cytokines known to recruit and activate adaptive immune cells. Therefore, further investigation is needed regarding the immunomodulatory properties of MSCs in the context of tumorigenesis and metastasis.

T_{regs} and metastasis

There is rapidly growing evidence that T_{regs} infiltrate tumours and positively correlate with poor prognosis in cancer patients [15,145–147], with the exception of renal cell carcinoma and certain haematological malignancies [148]. T_{regs} maintain immune tolerance and prevent inflammatory disease by suppressing cytotoxic T-cell activity and the proliferation of effector T cells [149–151]. Various subsets of T_{regs} of different immunophenotypes are found in both lymphoid and non-lymphoid organs [152–154]. As in the case of MSCs, there are many markers—such as CD25, FOXP3 and CTLA4—used to identify T_{regs}, the immunosuppressive role of which has been well-characterized [15]. Impairment of T_{reg} function induces a range of autoimmune and inflammatory diseases, such

as type I diabetes [155], rheumatoid arthritis [156] and systemic lupus erythematosus [157]. Hypomorphic alleles of the *FoxP3* gene and the loss of CTLA4 expression—which are both required for T_{reg} maintenance and function [158–160]—lead to severe systemic autoimmunity and lymphoproliferative disease in humans and animal models [15,161,162]. This suggests that T_{reg} infiltration into the tumour mass could render the tumour microenvironment immunosuppressive and thus prevent the inflammatory response to tumours.

Indeed, depletion of T_{regs} suppresses tumour progression in models of breast cancer [163], leukaemia, myeloma, fibrosarcoma [164], colon adenocarcinoma [165] and lung cancer [166]. Furthermore, the infiltration of T_{regs} into the primary tumour has been recently proposed to promote metastatic potential (Fig 3; [19,163,167]), although some studies disagree with these findings due to the differences in animal models and cell lines [168,169]. Administration of autologous CD25 antibody—which depletes the T_{reg} pool—alone or with concomitant stimulation of NKT cell activity can suppress pulmonary metastasis of 4T1 murine mammary tumour cells [163]. This suggests additive effects on the suppression of tumour metastasis when T_{regs} are inactivated and NKT cells are relieved from T_{reg}-mediated suppression (Fig 3).

Another study supporting the role of T_{regs} in promoting metastasis showed suppression of 4T1 lung metastasis after administration of CD25 antibody, and that the secretion of CCR4 ligand, CCL17 and CCL22 in the lungs can recruit CCR4⁺ 4T1 and CCR4⁺ T_{regs} [167]. In addition, the loss of lung metastasis in NOD/SCID mice can be restored by transferring CD25⁺ CD4⁺ T_{regs} from BALB/c mice [166]. Most importantly, this study identified a molecule responsible for T_{reg}-mediated apoptosis of NK cells, β -GBP, which is secreted by T_{regs} and specifically affects NK cells [167]. The same research group also identified a population of immunosuppressive B cells that express one of the T_{reg} markers, CD25, as well as B-cell markers B220 and CD19 [170]. This immunosuppressive population can also promote 4T1 lung metastasis, as shown by its reduction after B220 antibody treatment. Culturing CD25⁺ B220⁺ CD19⁺ regulatory B cells (B_{reg}) with non-regulatory CD4⁺ T cells leads to increased expression of FOXP3 in the latter population and the acquisition of immunosuppressive properties *in vitro*, suggesting that B_{reg} cells might promote T_{reg} conversion (Fig 3). Furthermore, only transfer of both regulatory B cells and non-T_{regs}—and not of B_{reg} cells alone—can restore lung metastasis in NOD/SCID mice, demonstrating that B_{reg} have an indirect, supporting role in T_{reg}-mediated lung metastasis. These studies provide fairly convincing evidence that T_{regs} have a crucial role in lung metastasis, but they also strongly suggest that there could be many other subpopulations involved in this process. For example, a specific subset of CD4⁺ CD25⁺ T_{regs} that express CCR6 is more frequently found in breast and colorectal tumours than its CCR6⁻ counterpart, and relies on TGF- β signalling for propagation *in situ* [171]. CD11c⁺ dendritic cells, which secrete TGF- β and stabilize FOXP3 and CCR6 expression, and TAMs, which recruit CCR6⁺ T_{regs} through CCL20 secretion, mediate this subtype-specific immunosuppressive response [171]. Other immune cells, such as myeloid-derived suppressor cells (MDSCs), secrete various cytokines—such as TGF- β —to recruit and activate T_{regs} in tumorigenic settings (Fig 3; [172,173]).

It has also been shown, however, that T_{regs} might promote metastasis independently of their immunosuppressive activity. RANK activity can repress the expression of maspin, which regulates cell adhesion and has been characterized as a metastasis inhibitor in

breast cancer [174–176]. T_{regs} in murine mammary carcinoma, which are induced by the overexpression of ErbB2, are a major source of RANKL and promote metastasis by suppressing maspin expression (Fig 3; [19]). In this case, T_{reg} cell-mediated metastasis does not occur through immunomodulation, as pulmonary metastasis can be enhanced with exogenous RANKL in the complete absence of T cells.

Therapeutic potential

Effective treatment for metastatic disease remains elusive, as the dissemination of tumour cells severely hinders the ability to deliver effectively therapeutic agents to numerous metastatic colonies [2]. Furthermore, various components of the metastatic microenvironment offer tumour cells protection and resilience against therapy [177]. Therefore, ideal candidates for the treatment of cancer metastasis should be able to efficiently home to metastatic sites and disrupt the protective environment of the metastatic lesions.

Application of MSCs. The tumour-targeting affinity of MSCs has been characterized, although some controversy remains, and proposed to be a potentially effective mode of drug delivery for cancer treatment [178–180]. Furthermore, MSCs might incorporate into the tumour stroma as CAFs [11], thereby effectively infiltrating the tumour microenvironment. Many therapeutic agents, ranging from oncolytic adenoviruses in ovarian cancer [181] to cytokines in melanoma [182] and various adenocarcinomas [183], have been delivered by tumour-tropic MSCs and shown to suppress tumour growth in experimental models. MSCs engineered to produce and deliver TRAIL in metastatic breast cancer [184] and pancreatic carcinoma [185] have been recently shown to suppress lung metastasis. The administration of TRAIL holds high therapeutic potential, as it induces apoptosis in transformed cells with minimal toxicity towards normal cells [186,187]. The application of a delivery platform is crucial, however, as the unstable pharmacokinetics of TRAIL in the circulation hinder efficient delivery to the tumour [188]. Studies demonstrating the targeting of MSCs to metastatic organs [18,92,93] suggest that TRAIL could be delivered to these metastatic sites effectively by using this strategy. Indeed, TRAIL-expressing MSCs have been shown to target metastatic lungs in an MDA-MB-231 experimental metastasis model [184].

Despite the potential advantage of using MSCs for drug delivery, their pro-tumorigenic and pro-metastatic effects described above are a cause for concern and could possibly outweigh the benefits of efficient delivery to tumour metastasis. Nevertheless, MSC drug delivery studies so far show that this is not the case [179,185,189,190] and it remains unknown whether such engineered MSCs lose their innate potential to promote tumour progression from genetic manipulation, or the pro-tumorigenic effect of such engineered MSCs has minimal impact in the context of such experimental settings.

However, a major caveat of the aforementioned MSC-based therapeutics is that the preclinical studies began MSC-mediated drug delivery relatively early in tumour progression—10 days after subcutaneous injection of pancreatic carcinoma cells [185] and 7 days after intravenous injection of breast adenocarcinoma cells [184]—whereas patients with metastatic disease would require treatment at much later stages of progression. Therefore, whether MSC drug delivery would elicit anti-metastatic effects in a clinical setting still needs to be explored.

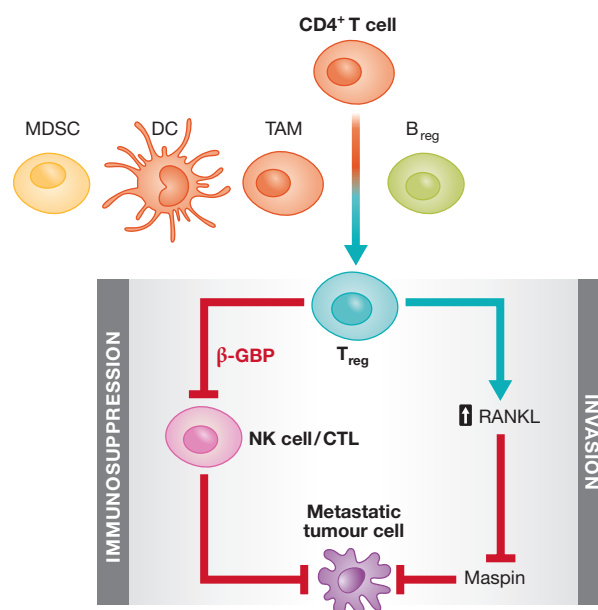


Fig 3 | T_{regs} promote cancer metastasis through immunosuppression and by stimulating invasive behaviour. Tumour-associated B_{regs} enhance the conversion of CD4⁺ T cells into immunosuppressive T_{regs}, which further promote cancer metastasis through immunosuppression and stimulating cell motility. Other immune cells, including MDSCs, DCs and TAMs, secrete various cytokines to recruit and activate T_{regs}. T_{regs}, in turn, inhibit the proliferation of tumour-reactive NK cells and CTLs to create a metastasis-permissive immune environment. T_{regs} can also produce abundant RANKL, which has been shown to promote tumour cell invasion through maspin suppression. B/T_{reg}: regulatory B/T cell; CTL, cytotoxic T lymphocyte; DC, dendritic cell; MDSC, myeloid-derived suppressor cell; NK, natural killer; RANKL, receptor activator of nuclear factor-κB (ligand); TAM, tumour-associated macrophage.

The greatest challenge in searching for therapeutic targets to disrupt tumour–MSC interactions lies in the fact that certain molecular mechanisms defined under pathological conditions are also crucial components of physiologically normal conditions. For example, CCL5 is secreted by tumour-associated MSCs to promote primary tumour growth and metastasis in a malignant setting, but its endogenous function is to recruit and activate inflammation-inducing immune cells [191]. Targeting of CCL5 as a therapeutic option, therefore, might be problematic and result in severely adverse effects unrelated to the intended therapeutic impact on cancer. Alternatively, extensive characterization of cancer-associated MSCs as compared with normal MSCs might yield unique cancer-associated MSC-specific properties as potential therapeutic targets. This is especially important when devising a therapeutic strategy to inhibit MSC activity, as normal MSCs are crucial in wound healing and tissue regeneration [54]. In fact, normal MSCs and their derivatives are being considered in numerous clinical trials for the treatment of bone fractures, stroke, multiple sclerosis and leukaemia [192–194]. For example, osteoprogenitors expanded *in vitro* on hydroxyapatite scaffolds have been used to successfully treat 4 patients with diaphyseal segmental defects in long bones [195,196]. Therefore, a detailed molecular characterization of cancer-associated MSCs in various metastasis

organ sites might provide new therapeutic approaches to target organ-specific metastasis.

Application of regulatory T cells. There is a strong clinical correlation between the presence of T_{regs} and poor prognosis of cancer patients [16,17,197]. The amount of T_{regs} in the primary tumours and sentinel lymph nodes has been proposed as a strong prognostic indicator for metastasis-free survival in breast cancer [198,199] and papillary thyroid cancer [200]. For example, the probability of 10-year survival of lymph node FOXP3-negative and FOXP3-positive subgroups of breast cancer patients is 41% and 18%, respectively [198]. Therefore, a great effort to suppress T_{reg} cell expansion and the immunosuppressive activity of this population has been made but has resulted in limited success.

Selective T_{reg} depletion by expressing FOXP3-driven diphtheria toxin receptor improves therapeutic vaccination against ovalbumin-expressing B16 melanoma cells in mice by allowing the accumulation of activated CD8⁺ CTLs [201]. However, T_{reg} cell depletion in the peripheral blood of melanoma patients with the CD25 neutralizing antibody Daclizumab failed to enhance the efficacy of dendritic cell vaccination [202]. By contrast, administration of the recombinant IL-2 diphtheria toxin conjugate denileukin diftitox to renal cell carcinoma patients, followed by RNA-transfected dendritic cell vaccination, significantly increased tumour cell-reactive T-cell responses [203]. This suggests that different cancer types might have different sensitivity to immunosurveillance, as well as different evasion mechanisms [204].

The suppression of cancer metastasis in patients receiving T_{reg}-cell-depleting agents has not yet been reported, although their benefits in the context of tumour progression and metastasis are being assessed in clinical trials (NCT01307618).

Conclusion and future directions

It has become increasingly evident that the distinct stages of tumour progression cannot be viewed as a linear cascade, but rather must be placed in the context of an intricate network of tumour–stroma interactions with multiple signalling and cellular feedback loops. Bone marrow-derived cells are crucial mediators in fulfilling the potential of tumour cells for distant metastasis, by using their innate versatility to perform a wide range of supportive functions. MSCs can put on many faces as accomplices in tumour progression and metastasis. Similarly, T_{regs} contribute to these complex processes not only with their immunosuppressive properties but also by inducing cell motility. Both MSCs and T_{regs} demonstrate how a beneficial endogenous function can be rendered harmful in the context of cancer progression. However, non-discriminatory targeting of MSCs and T_{regs} could lead to dysfunctional wound healing and autoimmune diseases, respectively. Therefore, how various bone marrow-derived populations in the metastatic context can be distinguished from those with normal endogenous function is a crucial goal that requires further research. Not only the MSCs, T_{regs} and other bone marrow-derived cells present at the primary tumour should receive attention, but also the bone marrow-derived cells recruited to the metastatic niche to promote organ-specific metastasis should become a major topic for further investigation. Given the challenges of targeting tumour-specific pathways, investigating the specificity of metastasis-promoting bone marrow-derived cells could provide a new targeted strategy for cancer treatment.

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CONFLICT OF INTEREST

The authors declare that they have no conflict of interest.

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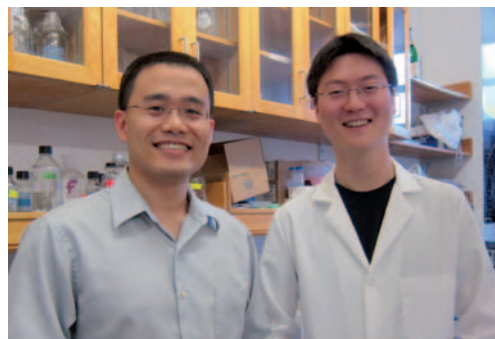
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