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Macrophages in tumor cell migration and metastasis

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Tumor-associated macrophages (TAMs) are a phenotypically diverse, highly plastic population of cells in the tumor microenvironment (TME) that have long been known to promote cancer progression. In this review, we summarize TAM ontogeny and polarization, and then explore how TAMs enhance tumor cell migration through the TME, thus facilitating metastasis. We also discuss how chemotherapy and host factors including diet, obesity, and race, impact TAM phenotype and cancer progression. In brief, TAMs induce epithelialmesenchymal transition (EMT) in tumor cells, giving them a migratory phenotype. They promote extracellular matrix (ECM) remodeling, allowing tumor cells to migrate more easily. TAMs also provide chemotactic signals that promote tumor cell directional migration towards blood vessels, and then participate in the signaling cascade at the blood vessel that allows tumor cells to intravasate and disseminate throughout the body. Furthermore, while chemotherapy can repolarize TAMs to induce an anti-tumor response, these cytotoxic drugs can also lead to macrophage-mediated tumor relapse and metastasis. Patient response to chemotherapy may be dependent on patientspecific factors such as diet, obesity, and race, as these factors have been shown to alter macrophage phenotype and affect cancer-related outcomes. More research on how chemotherapy and patient-specific factors impact TAMs and cancer progression is needed to refine treatment strategies for cancer patients.

KEYWORDS

cancer, macrophages, metastasis, tumor cell migration, chemotherapy, EMT

1 Introduction

Metastasis - the systemic spread of cancer - causes the majority of cancer-related deaths (1). To metastasize, cancer cells must be able to migrate through the tumor microenvironment (TME) and intravasate. Though not all tumor cells are inherently capable of such feats, the migratory and invasive phenotypes needed to accomplish these tasks can be induced through interactions with other types of cells in the TME, including endothelial cells, immune cells, and fibroblasts. Of the cellular components within the TME, macrophages are key players in the induction of pro-metastatic phenotypes in cancer cells. In this review, we provide an introduction to macrophages and their origin, discuss macrophage polarization, and then review the latest understanding of the role of macrophages in tumor cell migration and metastasis, including the promotion of 1) epithelialmesenchymal transition (EMT), 2) pro-tumoral extracellular matrix (ECM) remodeling, 3) tumor cell chemotaxis towards blood vessels, and 4) tumor cell intravasation. We also explore the impact of chemotherapy and host factors including diet, obesity, and race on tumor-associated macrophages (TAMs). Overall, we attempt here to summarize recent studies, discuss these new findings in the context of what is already known about the role of TAMs in tumor cell migration and metastasis, and highlight new potential avenues for refining therapeutic interventions.

1.1 Macrophage ontogeny

Macrophages have two distinct ontogenies. The first of these is monocyte-derived macrophages (MDMs) which originate from progenitors in the bone marrow and other hematopoietic niches (2), progress through several stages of differentiation, and enter the systemic circulation as monocytes. Circulating monocytes are recruited to tissues in response to locally released chemoattractants where they differentiate into macrophages (3). Once inside the tissue, MDMs may be short- or long-lived, and their population is maintained through recruitment of new circulating monocytes as well as proliferation of pre-existing MDMs (4, 5). The second group, known as tissue-resident macrophages, arise early in embryonic development, migrating from the yolk sac or fetal liver into developing organs where they differentiate into tissue-specific macrophages, including Kupffer cells (liver), osteoclasts (bone), and microglia (brain) (5). In adults, these macrophages self-renew largely independently of the bone marrow (6, 7). Macrophages in the TME are referred to as tumor-associated macrophages (TAMs). While most TAMs are monocyte-derived, tissue-resident macrophages make up a considerable percentage of TAMs in some tumor types (8-10).

1.2 Macrophage polarization

Once inside the tumor, macrophages take on various phenotypes and functions in response to stimuli in the microenvironment. These phenotypes are referred to as "polarization states." There is a wide spectrum of macrophage polarization states ranging from pro-inflammatory (M1) to antiinflammatory (M2).

M1 macrophages (historically called "classically activated"), are pro-inflammatory cells that participate in the host immune response against pathogens and can have anti-tumor activity. As such, environmental factors associated with infection and inflammation (including interferon (IFN)-y, bacterial lipopolysaccharide (LPS), and granulocyte-macrophage colony stimulating factor (GM-CSF)) promote M1 polarization (11). These signals cause macrophages to express surface proteins related to antigen presentation and T cell activation (including HLA-DR, CD80, and CD86) (11-14), and secrete inflammatory cytokines such as tumor necrosis factor (TNF)-a and interleukin (IL)-1 β to further enhance the immune response (11). M1 macrophages promote tumor cell killing through strong antigen presentation and effective activation of the innate and adaptive immune responses (15). Indeed, high M1 macrophage infiltration is correlated with positive outcome in cancer patients (16, 17).

M2 (or "alternatively activated") macrophages, are antiinflammatory cells that are involved in tissue repair and immune suppression. While these cells are essential in maintaining homeostasis in healthy tissues, they can also promote tumor growth and metastasis in the TME. M2 macrophages are induced by anti-inflammatory cytokines in the microenvironment, including IL-4 and IL-10 (11). These signals cause macrophages to express surface proteins such as CD163 and CD206, which are involved in tissue "clean-up" and homeostasis, and to secrete additional anti-inflammatory factors, such as IL-10 and transforming growth factor (TGF)- β , which further suppress the immune response (11, 18, 19). M2 macrophages also express high levels of vascular endothelial growth factor (VEGF), which promotes tumor vascularization, enhancing the delivery of oxygen and nutrients to the tumor (11). M2 macrophages are poor antigen presenters and suppress both innate and adaptive anti-tumor immunity (15). Furthermore, M2 macrophages are implicated in chemoresistance and metastasis, and high M2 infiltration is associated with poor prognosis in cancer patients (20-26).

Though macrophage polarization is a spectrum with M1 and M2 on opposing ends, it is common in the literature to oversimplify this state and treat macrophage polarization as a dichotomy (M1 or M2). Given that many "anti-inflammatory" macrophages also participate in inflammatory signaling and vice versa, the terms "M1" and "M2" should merely give a sense of how a macrophage is predominantly functioning in a particular environment. It is also important to note that macrophage phenotype is highly plastic. Similar to other components of the innate immune system, macrophage phenotype can quickly change in response to environmental cues (27). Indeed, in vitro and in vivo studies confirm that macrophages may repolarize in response to particular stimuli (28-30), an effect that has been leveraged in several immunotherapy clinical trials (31). Promising macrophagetargeting therapies and the challenges associated with their development are reviewed elsewhere (32-37). Given their significant, plastic, and diverse roles in cancer progression, understanding the mechanisms behind macrophage-mediated

cancer progression and the effects of chemotherapy and host factors is crucial to refining cancer treatment strategies.

2 TAMs in EMT

Epithelial-mesenchymal transition (EMT) is the process by which epithelial cells lose their characteristic apical-basal polarity and tight cell-cell junctions, and gain features associated with mesenchymal cells, including the ability to migrate and invade surrounding tissue (38, 39). In healthy tissues, EMT is used in critical processes such as embryonic development and wound healing. However, cancer cells hijack this program to gain migratory and invasive phenotypes. Cells that have undergone EMT are characterized by the loss of E-cadherin and the increase in N-cadherin and vimentin. E-cadherin, often used as an epithelial cell marker, is an important cell-cell adhesion protein involved in contact-mediated inhibition of cell growth (40). During EMT, the transcription factor SNAIL directly represses E-cadherin transcription and is thus crucial in EMT regulation (41). As Ecadherin decreases, the mesenchymal cell markers N-cadherin and vimentin increase and support tumor cell survival and migration (42, 43). During this process, tumor cells pass through a series of epithelial/mesenchymal (E/M) hybrid states that reflect varying degrees of plasticity and metastatic potential (44). Tumorassociated macrophages have long been known to play a role in EMT induction (45, 46), and more recent evidence shows that TAMs also promote progression to later E/M hybrid states (44). A number of recent studies have further elucidated the mechanisms behind this relationship, pointing to feedback loops in which tumor cells undergoing EMT attract and polarize macrophages, which then secrete factors that further promote EMT in tumor cells (45–47) (Figure 1).

Macrophages can induce EMT in cancer cells by secreting various factors, including TGF- β (48), CCL2 (49), and IL-6 (50), all of which ultimately lead to SNAIL upregulation and subsequent EMT in tumor cells. For instance, IL-6 activates the JAK2/STAT3 pathway upon binding the IL-6 receptor (47, 50, 51) (Figure 1A). The JAK2/ STAT3 axis is a critical signal transduction pathway that participates in many cellular functions including proliferation, differentiation, and survival, and components of this pathway are hyperactivated in many cancers (52, 53). After IL-6 receptor activation, STAT3 inhibits the transcription of tumor suppressor microRNAs including miR-34a (50, 51). MiR-34a suppresses SNAIL, and loss of miR-34a leads to SNAIL upregulation and subsequent EMT (54, 55), as well as tumor cell proliferation and migration (50, 56) (Figure 1B). Macrophageinduced mesenchymal-like tumor cells then secrete increased amounts of CCL2, which recruits macrophages (47), and IL-6, which leads to M2 polarization (51, 57) (Figure 1C), further propagating EMT in a positive feedback loop.



FIGURE 1

(A) Macrophages secrete IL-6, which binds to the IL-6 receptor on tumor cells, activating the JAK2/STAT3 pathway. After IL-6 receptor activation, STAT3 translocates to the nucleus and suppresses transcription of miR-34a, which leads to SNAIL upregulation. (B) The increase in SNAIL leads to loss of E-cadherin, and EMT programs become active and increase the expression of N-cadherin and vimentin. The tumor cell takes on a mesenchymal-like phenotype, which affords enhanced migration capacity. (C) Mesenchymal-like tumor cells secrete factors that recruit macrophages to the TME (e.g. CCL2, CCL5, and CXCL2) and that promote M2 polarization (e.g. IL-6). M2 polarization is characterized by the expression of surface markers such as CD163 and CD206. Figure created with BioRender.com.

These studies reveal several therapeutic targets with the potential to reduce the co-induction of EMT and M2 polarization. Inhibiting IL-6 signaling by targeting IL-6 itself, or its receptor (with anti-IL-6R monoclonal antibodies like tocilizumab), reduces EMT, decreases M2 polarization and increases M1 polarization (51). The tumor suppressor miR-34a is also a potential target. MiR-34a suppresses SNAIL and reinstates an epithelial phenotype in mesenchymal-like cancer cells (55). MiR-34a expression in tumor cells also promotes macrophage M1 polarization, demonstrating that the microRNA can favorably modify both the tumor cells and the immune microenvironment (51). Indeed, nanoparticle-delivered miR-34a has shown promise in treating several types of cancers (58, 59). In addition to promoting EMT in cancer cells, SNAIL is also involved in macrophage recruitment. SNAIL expression in tumor cells increases their secretion of CCL2, CCL5, and CXCL2, all of which attract macrophages to the TME (60-62) (Figure 1C). Indeed, SNAILoverexpressing tumors show a significant increase in macrophage infiltration, M2 polarization, and metastasis (60, 62).

Finally, in addition to cytokines and chemokines, more recent evidence has revealed that exosomes can also mediate macrophagetumor cell feedback loops related to EMT and M2 polarization. Tumor cells that have undergone EMT secrete exosomes containing microRNAs that are taken up by macrophages and induce M2 polarization (63). For example, it was shown that tumor cell derived exosomes contain miR-106b-5p, which upon uptake by macrophages, activates the PI3K γ /AKT/mTOR signaling pathway to induce M2 polarization by downregulating the pathway inhibitor PDCD4 (63). Similarly, SNAIL expression directly upregulates the transcription of miR-21 in tumor cells. This microRNA is then transferred to macrophages through exosomes and also targets macrophage PDCD4, leading to M2 polarization (64).

Exosomal microRNAs can be transmitted from macrophages to tumor cells as well (65). Tumor cell uptake of M2 macrophagederived exosomes leads to downregulation of E-cadherin and upregulation of N-cadherin and vimentin (65). Lu et al. found that these exosomes contain miR-23a-3p, which downregulates the tumor suppressor PTEN (65) – a known regulator of EMT (66). In a positive feedback loop, tumor cells treated with M2 macrophagederived exosomes express higher levels of CCL2, leading to increased macrophage recruitment and M2 polarization (65).

In summary, tumor cell EMT and macrophage recruitment and polarization are intimately connected and co-regulated by several molecular mechanisms.

3 TAMs in extracellular matrix remodeling

3.1 TAMs in matrix stiffness

Macrophages also promote tumor cell migration and invasion through ECM remodeling. The ECM of non-cancerous soft tissue is characterized by "curly," non-dense collagen fibers that lay parallel to the epithelium (67). This soft ECM is involved in maintaining an epithelial phenotype, and matrix stiffening has been shown to play a direct role in promoting EMT (68, 69). Indeed, clinical conditions characterized by a stiff ECM – including cirrhosis of the liver (70), pulmonary fibrosis (71), and mammographically dense breast tissue (72) – are associated with a higher incidence of cancer in the respective tissues. In tumors, collagen deposition increases, and fibers become stiff, cross-linked and linearized – a process known as desmoplasia that has been associated with immune evasion and metastasis (67, 73–76). Indeed, tumors have been shown to be stiffer than healthy tissue in breast (77), pancreas (78), bladder (79), and ovarian (80) cancers. The stiffened matrix of tumors promotes malignant transformation, proliferation, and invasion of tumor cells, and acts as a "highway," guiding tumor cells towards the vasculature, where they further invade and intravasate (67, 69, 76, 77, 81–85). Recent work sheds light on the mechanistic role of TAMs in pro-tumoral matrix stiffening.

Macrophages promote matrix deposition and stiffening in both cancerous and healthy tissue (86, 87). In pancreatic cancer, macrophages foster desmoplasia indirectly by activating pancreatic stellate cells. Mechanistically, macrophages internalize and degrade surrounding collagen, which leads to an increase in inducible nitric oxide synthase (iNOS) and the production of reactive nitrogen species (RNS). RNS then activate pancreatic stellate cells leading to increased collagen deposition and desmoplasia (87).

In the desmoplastic reaction, excessive ECM deposition is followed by cross-linking, which confers increased stiffness to the TME. ECM crosslinking is mediated primarily by lysyl oxidase (LOX) and lysyl oxidase-like (LOXL) proteins, which are expressed by a variety of cells in the TME (88). In pancreatic cancer, LOXL2 expression is positively associated with tumor burden and metastasis (89). Macrophages both express their own LOXL2 and promote its expression in tumor cells (89, 90). Alonso-Nocelo et al. recently demonstrated that macrophage depletion leads to a significant decrease in LOXL2, collagen fibril orientation, and metastasis in mice, indicating that macrophages promote matrix stiffness (89). In a positive feedback loop, the stiffened matrix then promotes macrophage infiltration and M2 polarization (89). Mechanistically, macrophages increase matrix stiffness by secreting oncostatin M (OSM), which upregulates LOXL2 in tumor cells (89). In turn, the stiffened matrix activates integrin $\beta 5$ in macrophages, leading to FAK/MEK/ERK activation and LOXL2 upregulation, further supporting ECM crosslinking in the TME (90). In addition to promoting tumor cell migration, stiffened matrices cause macrophages to take on a more immunosuppressive phenotype (89, 91). Indeed, macrophages cultured on stiff matrices recruit cytotoxic T cells less efficiently than those cultured on softer matrices (91).

Together, these studies indicate that macrophages support the development of a stiff ECM through direct and indirect mechanisms. In turn, the stiff ECM promotes macrophage recruitment, M2 polarization, tumor cell migration, and metastasis.

3.2 TAMs in matrix degradation

Equally as important as matrix stiffness for cancer progression is matrix degradation, which is mediated by matrix metalloproteinases (MMPs). MMPs are a group of zinccontaining proteolytic enzymes responsible for degrading the extracellular matrix (92). MMPs are upregulated in nearly every type of cancer, and their activity has been shown to facilitate angiogenesis, tumor cell immune evasion, migration, and metastasis (92, 93). MMPs are expressed by a variety of stromal, immune, and tumor cells, and a growing body of evidence reveals the role of MMPs in the dynamic pro-metastatic interplay between macrophages and tumor cells.

MMP production can be induced through several major signal transduction pathways including STAT3, ERK, and NF-KB (94-102). Evidence shows that macrophages provide multiple ligands for these pathways that cooperate to promote MMP expression. For example, macrophages secrete AEG-1, TGF-B, and IL-6, which all increase MMP-9 expression in tumor cells by activating STAT3 (94, 95, 97). Indeed, inhibiting STAT3 in tumor cells, or its activators in macrophages, causes a significant decrease in MMP expression and migration in tumor cells (95, 97). Macrophages also secrete TNF- α and IL-1 β , which activate the NF- κ B pathway. Yamanaka et al. found that IL-1 β activates NF- κ B in gastric cancer cells, and this leads to increased MMP-9 expression and tumor cell invasion (103). Furthermore, tumor cells cultured in M2 macrophage-conditioned media express significantly increased levels of MMP-9 (98). This effect can be seen to a lesser (but still significant) extent when tumor cells are stimulated with TNF- α alone, suggesting that macrophages provide multiple ligands that stimulate MMP production (98).

MMPs expressed by macrophages also play a significant role in tumor cell invasion and metastasis. Macrophage - but not tumor cell - expression of MMP-11 is a negative prognostic marker in breast cancer (104). MMP-11-overexpressing macrophages secrete increased amounts of CCL2. CCL2 then activates MAPK signaling in tumor cells and increases tumor cell migration and MMP-9 expression (104). In Wilms' tumor and gastric cancer, MMP-9 is upregulated in M2 macrophages, and MMP-9 initiates EMT and increases tumor cell invasion (105, 106). Mechanistically, macrophage-derived MMP-9 activates the PI3K/AKT pathway in tumor cells leading to the upregulation of SNAIL and subsequent EMT (105, 106). These studies identify MMP-9 as a promising therapeutic target. Indeed, MMP-9 inhibition increased the efficacy of chemotherapy and decreased metastasis to the lungs in a mouse model of gastric cancer (106). Together, these studies identify macrophages as important regulators of tumor cell MMP production.

4 TAMs in tumor cell chemotaxis

Beyond promoting a mesenchymal phenotype in cancer cells, TAMs also supply ligands and chemotactic factors that support tumor cell migration and invasion in the tumor microenvironment (Figure 2).

In vitro and *in vivo* migration assays and intravital imaging show that tumor cells and macrophages migrate through the TME together using a CSF1/EGF paracrine loop that leads to invasion and metastasis (107–110) (Figure 2A). High levels of CSF1 (111–113) and the EGF receptor (114–117) are correlated with metastasis and poor prognosis in a number of solid tumors. CSF1 secreted by

tumor cells both recruits macrophages to the tumor microenvironment and promotes macrophage expression of EGF (108). TAM-derived EGF then binds to the EGF receptor on tumor cells, leading to increased CSF1 production and activation of pathways associated with migration (107, 108, 118) (Figure 2A). Using this paracrine loop, macrophages and tumor cells migrate together along fibronectin-collagen I ECM fibers towards chemotactic gradients. Leung et al. found that in the TME, the primary chemo-attractant for the macrophage-tumor cell pair is hepatocyte growth factor (HGF), which is secreted by endothelial cells (119) (Figure 2). Within 500 µm of a blood vessel, tumor cells may perform sustained directional migration towards HGF gradients with or without macrophages. However, tumor cells at greater distances can only move towards blood vessels by comigrating with a macrophage (119). Thus, while tumor cells may chemotax along these HGF gradients alone, co-migrating with macrophages greatly supports their ability for sustained directional migration and extends the chemoattractive influence of the blood vessels.

In addition to participating in tumor cell chemotaxis, macrophages support tumor cell migration by both promoting the formation of tumor cell invadopodia and prolonging their activity (Figure 2). Invadopodia are F-actin-rich protrusions with MMP activity used to degrade the ECM and create new physical pathways through the tumor (120, 121). TAMs promote the formation of invadopodia by secreting EGF, which activates the EGF receptor in tumor cells. EGF receptor activation initiates the assembly of invadopodial precursors through the recruitment of actin regulatory proteins such as cortactin, Arp2/3, and cofilin (118, 120, 122, 123) (Figure 2A). Phosphorylation of cortactin activates actin polymerization and leads to maturation of a precursor. The actin regulatory protein Mena (encoded by the ENAH gene) supports this polymerization by localizing to the barbed ends of polymerizing actin filaments and temporarily interfering with the capping proteins that block polymerization (124, 125). In non-invasive tumor cells, invadopodia can form, but do not mature, as cortactin is rapidly dephosphorylated by the tyrosine phosphatase PTP1B that is constitutively bound to Mena. This lack of maturation dramatically limits the invasive capacity of invadopodia by limiting the amount of matrix they can degrade (126).

Macrophages also play a role in promoting invadopodium maturation (thus increasing degradative activity) by stimulating the expression of a splice variant of Mena called MenaINV (Figure 2B). MenaINV prolongs the degradative activity of invadopodia by sequestering PTP1B and preventing the dephosphorylation/deactivation of cortactin and the subsequent disassembly of actin filaments (127, 128). The MenaINVstabilized invadopodium then degrades the ECM in its path, facilitating tumor cell migration towards blood vessels (Figure 2C). Macrophages promote the alternative splicing of Mena through cooperative Notch1/NF-KB signaling (129, 130) (Figure 2B). Mechanistically, macrophages secrete $TNF\alpha$, which activates the NF-KB pathway in tumor cells, leading to p65 nuclear translocation. Inside the nucleus, p65 binds to the kB binding sites on the ENAH promoter, initiating ENAH transcription. Macrophages also express the Notch1 ligand Jagged1, which



Macrophages and tumor cells co-migrate through the TME along fibronectin-collagen I ECM fibers towards HGF gradients secreted by endothelial cells using an EGF/CSF1 paracrine loop. (A) Macrophage-derived EGF activates the EGF receptor on the tumor cell, leading to the upregulation of genes associated with cell migration and invadopodium formation. (B) Cooperative Notch1/NF-xB signaling between the macrophage and tumor cell leads to an increase in MenaINV expression, which enhances invadopodium stability and degradative activity. (C) The invadopodium degrades ECM in its path, facilitating tumor cell migration towards the blood vessel. Figure created with BioRender.com.

engages the Notch1 receptor on tumor cells, causing nuclear translocation of the Notch1 intracellular domain (NICD). Nuclear NICD enhances the nuclear retention of p65, leading to sustained *ENAH* transcription and to alternative splicing (130). Prior work has shown that this alternative splicing is the switch that turns non-invasive tumor cells into invasive tumor cells (131).

In summary, macrophages partner with tumor cells to enhance directional migration and metastasis by guiding tumor cells towards blood vessels and promoting the assembly and invasion capacity of tumor cell invadopodia.

5 TAMs in tumor cell intravasation

Intravasation – the process by which tumor cells enter the vasculature – represents a key step in the metastatic cascade. Macrophages not only assist with tumor cell intravasation but are crucial for the process.

Breast cancer cells disseminate from the primary tumor through tumor microenvironment of metastasis (TMEM) doorways (132,

133). TMEM doorways are stable, tri-cellular structures (occurring primarily at vascular branch points) composed of a Menaexpressing tumor cell, a perivascular Tie2^{High} macrophage, and an endothelial cell in direct physical contact (132, 134-137). TMEM doorway density (hereafter referred to as TMEM doorway score) in the primary breast tumor is a clinically validated prognostic marker of distant metastasis (136, 138). Arwert et al. investigated the process of TMEM doorway assembly by systemically depleting macrophages and then tracking the fate of newly-recruited monocytes in the TME (139). They found that upon extravasation into the TME, monocytes become motile TAMs that begin to express CXCR4 and are then recruited back to the perivascular space by CXCL12-expressing perivascular fibroblasts. Once at the blood vessel, these motile TAMs become sessile, forming TMEM doorways with adjacent tumor and endothelial cells (139). Signaling between the three TMEM doorway cells results in the release of vascular endothelial growth factor-A (VEGFA) (132). The secreted VEGFA leads to the dissociation of local vascular endothelial cell-cell junctions, which causes a transient, localized vascular permeability event. Harney et al. used real-time

multiphoton intravital imaging of a murine mouse model of breast cancer to show that these transient vascular permeability events are regulated and occur concurrent with tumor cell intravasation (132). Neither transient vascular permeability nor tumor cell intravasation occurs away from TMEM doorways (132). Furthermore, TMEM doorway score increases concomitantly with circulating tumor cells, and macrophage depletion leads to a significant reduction in TMEM doorways, vascular permeability, and circulating tumor cells, highlighting the essential role of TMEM doorway-associated macrophages in tumor cell intravasation (132, 133).

In vitro and in vivo studies confirm that invadopodia formation is necessary for tumor cell intravasation (140–143). In addition to initiating invadopodium formation through paracrine EGF signaling, macrophages can also initiate this process through direct contact (129, 140). In TMEM doorways, contact between the TMEM doorway-associated macrophage and tumor cell induces invadopodium formation in the tumor cell (129, 140). The invadopodium then degrades the basement membrane surrounding the vascular endothelium and functionally "holds the door open" for other migratory tumor cells to enter the blood stream (129, 140). Mechanistically, macrophage-tumor cell contact activates RhoA signaling in tumor cells, which initiates invadopodium formation in the tumor cell (140, 144). Indeed, RhoA knockdown reduces tumor cell invadopodium formation, matrix degradation, and intravasation (140, 144).

Importantly, targeting TMEM doorway-associated macrophages with the Tie2 inhibitor rebastinib has shown therapeutic promise by decreasing TMEM doorway function and metastasis in preclinical studies of breast cancer and pancreatic neuroendocrine tumors (133, 145). Mice treated with rebastinib have significantly reduced TMEM doorway activity, circulating tumor cells, and metastases compared to mice treated with vehicle control (133, 145).

Together, these studies identify macrophages as key mediators of tumor cell intravasation and demonstrate that blocking crucial macrophage signaling pathways may be a strategy to block tumor cell dissemination in patients.

6 TAMs in response to chemotherapy

Cytotoxic chemotherapies are characterized by their ability to directly prevent proliferation and promote apoptosis of dividing cells. Chemotherapeutic agents, including anthracyclines, platinum-based drugs, and other alkylating agents induce apoptosis by damaging DNA and preventing DNA replication and repair (146). Taxanes and vinca alkaloids prevent cell division by interfering with the mitotic spindle, and antimetabolites – structural analogs of nitrogenous bases – prevent DNA synthesis by getting fraudulently inserted into replicating DNA, as well as by preventing the synthesis of proper bases (146). Increasing evidence suggests that many of these drugs also exert indirect effects by modulating the immune microenvironment. While many of these indirect effects support tumor cell killing, some promote drug resistance and metastasis. In this section, we review the current understanding of how common chemotherapies affect macrophages in anti- and pro-tumoral ways.

6.1 Anti-tumor TAM response to chemotherapy

Paclitaxel - a microtubule stabilizing agent in the taxane group - has been shown to increase the immune response and tumoricidal activity of murine macrophages. In mice, paclitaxel treatment leads to a robust increase in macrophage expression of TNF α and IL-1 β , pro-inflammatory cytokines associated with the M1 phenotype (147-149). Paclitaxel also increases macrophage expression of IL-12, a Th1-type cytokine involved in activating the innate and adaptive immune response (148, 150). As a LPS mimetic, paclitaxel activates toll-like receptor (TLR) 4 on murine macrophages leading to NF-KB activation and increased production of pro-inflammatory signals (148). Though some studies show that paclitaxel also activates TLR4 in human cells (151-153), others show that species-specific differences in the TLR4 accessory protein, myeloid differentiation factor 2 (MD-2), do not allow this activation (154-159). Interestingly, some studies indicate that docetaxel - another taxane - has more potent effects on human macrophages than paclitaxel. Millrud et al. found that docetaxel, but not paclitaxel, promoted an M1 phenotype in human monocytes (160). Furthermore, a clinical study assessing immune responses to taxanes in breast cancer patients showed that, while both docetaxel and paclitaxel lead to an increase in serum M1-associated markers (including IL-6, GM-CSF, and IFN- γ), the effects were significantly more pronounced in patients who received docetaxel (161). The effects of taxanes on macrophages are also highly context dependent. For instance, IFN-y has been shown to "prime" macrophages for tumoricidal activity, and paclitaxel affords macrophages increased cytotoxicity after macrophage exposure to IFN- γ (162). While the exact mechanisms have yet to be elucidated, these studies indicate that taxanes can promote anti-tumor M1 polarization in a context-dependent manner.

In addition to taxanes, platinum-based drugs, antimetabolites, and alkylating agents have also been shown to promote M1 macrophage polarization. The combination of platinum-based agents with antimetabolites is a common first-line treatment for gastric cancer (163). In studies analyzing the TME of matched preand post-treatment biopsies from gastric cancer patients, posttreatment samples harbored significantly more M1-polarized macrophages, and this increase was associated with a favorable response to treatment (164, 165). Furthermore, when given at high doses, the alkylating agent cyclophosphamide is highly immunosuppressive. However, lower doses of the drug strikingly improve anti-tumor immunity (166). This has led to the development and use of metronomic schedules of administration, in which low doses of the drug are administered more frequently (166). In line with observations that low-dose cyclophosphamide improves anti-tumor immunity, several studies show that low-dose, metronomic cyclophosphamide increases macrophage M1 polarization and decreases tumor burden (167-169).

Leukemia and lymphoma are often treated with monoclonal antibodies. While these treatments are largely effective at targeting cancer cells in many niches, cancer cells often become resistant to such antibodies in the bone marrow (170–172). Several studies found that combining antibody therapy with cyclophosphamide prevented antibody therapy resistance in the bone marrow in part by promoting macrophage phagocytosis of antibody-targeted cancer cells (170–172).

In summary, these studies show that in some circumstances, chemotherapy reprograms macrophages to increase antitumor activity.

6.2 Pro-tumor TAM response to chemotherapy

The macrophage response to chemotherapy is a double-edged sword. While some studies show that chemotherapy promotes the anti-tumor activity of macrophages, others show that chemotherapy causes a macrophage-mediated pro-tumoral response.

Chemotherapy causes tumor cell death and tissue damage followed by a cytokine storm that promotes the release of endothelial and immune progenitor cells from the bone marrow (173–175). In response to this tissue damage, cells in the TME initiate a wound healing response by increasing their secretion of CSF1, CXCL12, and other chemokines that recruit these circulating progenitor cells to the tumor (176, 177). One result of this response is that perivascular TAMs increase following chemotherapy (133, 177). These newly recruited perivascular TAMs express high levels of VEGFA and the angiopoietin receptor Tie2, which have been shown to promote relapse and metastasis following chemotherapy (132, 133, 177–180). Several studies show how chemotherapy induces a macrophage-mediated pro-tumoral effect that can be abrogated by targeting macrophages.

Hughes et al. used mouse models of breast cancer to show that treatment with cyclophosphamide causes an increase in CXCR4expressing perivascular macrophages, which promote tumor revascularization and regrowth via VEGFA signaling (177). Blocking CXCR4 signaling prevents the accumulation of perivascular macrophages and delays tumor regrowth (177).

In neuroblastoma, chemotherapy leads to the selective expansion of CCL2-expressing mesenchymal-like tumor cells and macrophage infiltration in patients, which promotes relapse and chemo-resistance (181). In mouse models, combining chemotherapy with CSF1R inhibition prevents macrophage infiltration and tumor regrowth (181).

Furthermore, while chemotherapy increases the infiltration of Tie2+ macrophages, Tie2 inhibitors have been shown to work synergistically with chemotherapy to delay tumor growth (145) and relapse (182).

In addition to promoting tumor relapse, macrophages can also increase tumor cell dissemination following chemotherapy. We have previously shown that treatment with paclitaxel causes a robust, macrophage-dependent increase in MenaINV expression, which promotes tumor cell migration, intravasation, and metastasis (130, 133). This indicates that paclitaxel causes a macrophagemediated increase in metastasis-competent tumor cells, though the exact mechanism behind this effect remains unknown.

Furthermore, chemotherapy significantly increases the assembly and function of TMEM doorways, which are portals for tumor cell intravasation. Indeed, circulating tumor cells and lung metastases are more prevalent in mice treated with paclitaxel compared to vehicle control (133). Concerningly, TMEM doorway assembly is also increased in patients with ER+/HER2-breast cancer following neoadjuvant chemotherapy (133), thus increasing their risk of distant metastasis (136). As mentioned in Section 5, targeting TMEM doorway-associated macrophages with the Tie2 inhibitor rebastinib dramatically decreases the prometastatic effects of chemotherapy in pre-clinical studies, indicating that Tie2 inhibition in combination with a cytotoxic agent may improve patient outcomes (133, 145).

Another mechanism by which TAMs promote metastasis in response to chemotherapy is by upregulating the enzyme heparanase. Heparanase cleaves heparan sulfate, which is an important structural component of the ECM (183). Similarly to MMPs, this matrix-degrading enzyme is upregulated in many cancers and correlates with increased metastasis and poor prognosis (184). Unfortunately, heparanase has been shown to increase in some patients following chemotherapy (185). Mechanistically, treatment with chemotherapy leads to an increase in VEGFR3-expressing TAMs which secrete cathepsins that activate heparanase and promote ECM remodeling, lymphangiogenesis, and metastasis (186). Notably, blocking VEGFC/VEGFR3 signaling inhibits chemotherapy-induced lymphangiogenesis and metastasis (186).

In summary, chemotherapy can act on macrophages to promote relapse and metastasis in a variety of ways. Recent preclinical studies show that targeting macrophage recruitment or function is a promising approach to optimize cancer treatment. Indeed, there has been a 3-fold increase in clinical trials on macrophage-targeted therapies in the past 10 years (31). However, due to the diversity of patients, chemotherapies, and macrophage phenotypes, more research is needed to clarify the exact mechanisms of chemotherapy-induced cancer progression to refine treatment strategies and determine biomarkers that can identify good – and bad – candidates for different treatments.

7 Host factors governing TAMs

7.1 Diet and natural compounds

Evidence supporting the role of a healthy diet in cancer prevention and treatment is ever-increasing (187, 188). While natural compounds are known to have a profound role in regulating EMT in cancer cells (189, 190), numerous recent studies have also shed light on how dietary agents and natural compounds target tumor-associated macrophages.

7.1.1 Antioxidants

Perhaps the most well-known dietary anti-cancer agents are antioxidants – compounds that neutralize free radicals that would

otherwise damage DNA and other cellular structures and lead to carcinogenesis. Foods high in antioxidants include berries, fruits, vegetables, walnuts, and pecans (191). Recently, Latronico et al. demonstrated that dietary antioxidants act on macrophages and inhibit the expression and activity of macrophage-derived MMP-2 and MMP-9, which have pro-tumor ECM remodeling activity (see Section 3.2) (192). Macrophages mediate the production of reactive oxygen species (ROS) in the TME (193), and there is an established relationship between ROS and MMP production (194, 195). The authors posit that dietary antioxidants prevent MMP production by removing ROS in the microenvironment (192).

Propolis, a natural resin produced by honeybees, also has antioxidative properties (196–198). It is widely used as a natural additive in both ingestible (i.e. capsules, throat lozenges, food) and topical (i.e. lotions, cosmetics) products. Propolis induces the depolarization and repolarization of M2 macrophages to M0- and M1-like states, respectively. M2-polarized macrophages treated with propolis also express decreased levels of IL-8, IL-10, CCL2, VEGF, and MMP-9 (199). Consistent with this shift of M2 to M1 macrophages, propolis significantly decreases EMT and tumor cell migration and invasion (199).

For centuries, cloves have been used not only as a spice, but also as an herbal remedy due to their antimicrobial and antioxidative properties (200). Kumatakenin, a flavonoid isolated from cloves, has recently shown significant anti-cancer effects by acting on both tumor cells and tumor-associated macrophages (201). In addition to inducing apoptosis in human ovarian cancer cells, kumatakenin reduces tumor cell expression of CCL2 and CCL5 – both implicated in macrophage recruitment, cancer progression and metastasis (201–203). Kumatakenin also prevents M2 polarization and macrophage expression of IL-10, VEGF, MMP-2, and MMP-9 (201).

Together, these studies implicate antioxidants in reducing macrophage-mediated pro-tumoral effects including immunosuppression, angiogenesis, and pro-tumoral ECM remodeling.

7.1.2 Vitamin D and omega-3 poly-unsaturated fatty acids

Vitamin D is a steroid hormone precursor that can come from the diet or be endogenously synthesized in the skin upon exposure to UV radiation (204). Vitamin D exerts its effects by binding to the vitamin D receptor and is primarily responsible for regulating calcium and phosphate levels in the body (205, 206). Though there is no clear consensus on the impact of vitamin D on cancer, a recent study showed that vitamin D may exert anti-cancer effects through macrophages. *In vitro* studies showed that vitamin D reverses M2 polarization, decreases macrophage secretion of TGF- β 1 and MMP-9, and reduces the macrophage-induced proliferation and migration of ovarian cancer cells (207).

Omega-3 poly-unsaturated fatty acids (ω -3 PUFAs) – long lauded for their anti-inflammatory properties – also exhibit anticancer effects on tumor-associated macrophages (208). Studies on mouse models of castrate resistant prostate cancer show that a diet rich in ω -3 (vs. ω -6) PUFAs significantly delays tumor progression, decreases M2 polarization, and increases M1 polarization and infiltration of CD4+ T cells. M2 macrophages from tumors in mice fed a high ω -3 diet also show a significant decrease in MMP-9 and VEGF expression (208).

Together, these studies have begun to reveal the mechanisms by which a healthy diet can induce anti-cancer changes in macrophages and the TME more broadly.

7.2 Obesity

Obesity is a fast-growing global health crisis that is associated with an increased risk of cancer, as well as general morbidity and mortality (209, 210). In fact, women with obesity who are diagnosed with breast cancer have an increased risk of distant metastasis and are less likely to respond to some cancer treatments (211-214). Unsurprisingly, adipose tissue macrophages are thought to play a key role in creating a pro-tumorigenic microenvironment (86). CCL2 expression is significantly increased in the adipose tissue of obese compared to lean mice (215) which leads to the accumulation of macrophages. Indeed, it is estimated that macrophages make up <10% of cells in the adipose tissue of lean individuals and nearly 40% of cells in the adipose tissue of obese individuals (216). These recruited macrophages surround dead and dying adipocytes, forming crown-like structures (CLS) that are characteristic of adipose tissue inflammation (217). Overweight and obese patients $(BMI \ge 25 \text{ kg/m}^2)$ with breast cancer are more likely to have CLS compared to patients at a healthy weight (BMI $< 25 \text{ kg/m}^2$), and BMI \geq 25 kg/m² is associated with a shift in CLS macrophage phenotype that may be indicative of metabolic dysfunction and poor treatment outcomes under some conditions (218). These macrophages also interact with pre-adipocytes and prevent their differentiation, instead causing them to take on a fibroblastic phenotype and enhance the synthesis and deposition of ECM components (219, 220). This indicates that macrophages cause increased ECM density in obese adipose tissue, including in the breast where ECM density is a significant risk factor for cancer (72, 221). Further implicating adipose tissue macrophages in breast tumor development, a study on transgenic mice that overexpress CCL2 in the mammary epithelium showed that CCL2 overexpression causes increased macrophage density, stromal density, and ECM crosslinking enzyme LOX compared to non-transgenic controls (222). CCL2overexpressing mice also had an increased susceptibility to DMBAinduced mammary tumors, demonstrating a relationship between macrophages, ECM density, and cancer risk (222).

Though much is still unknown about how obesity shapes cancer development, these studies suggest that obesity promotes macrophage-mediated ECM stiffening that is known to support tumorigenesis (see section 3.1).

7.3 Race

There are widespread racial disparities in the diagnosis, treatment, and outcome of cancer patients (223–226). While some of these disparities are due to systemic racism in the medical field, some studies identify biological differences that could be contributing to this effect (227). Indeed, Black men and

women with prostate and breast cancer, respectively, have significantly worse outcomes than their white counterparts even after socioeconomic and other mediating factors are accounted for (228, 229). We previously evaluated distant recurrence-free survival (DRFS) in breast cancer patients following neoadjuvant (NAC) versus adjuvant (AC) chemotherapy and found that treatment type had no impact on DRFS for white women (230). However, Black women had significantly worse DRFS when treated with NAC (230). Differences in TAMs may contribute to this. Black breast cancer patients have significantly increased macrophage infiltration and M2 polarization compared to white patients, and this is prognostic of progression-free survival (231, 232). Black, compared to white women treated with neoadjuvant chemotherapy for ER+/HER2- breast cancer also have a higher TMEM doorway score and macrophage density in the residual tumor tissue, which may contribute to poorer outcomes (223). In summary, racial disparities in cancer development and outcome may be mediated in part by macrophage infiltration and activity.

8 Conclusion

The diversity and complexity of tumor-associated macrophages leaves their functions highly context-dependent and variable. Despite this complexity, a consensus is emerging in the literature that tumor-associated macrophages support tumor cell migration and metastasis in many ways (Figure 3). TAMs confer migratory abilities in tumor cells by activating EMT. They remodel the ECM to facilitate tumor cell migration and provide ligands to promote invadopodium formation and chemotaxis. Finally, TAMs participate in the signaling cascade that opens TMEM doorways and allows tumor cells to intravasate and disseminate.

The plastic nature of TAMs means that their phenotypes and functions can dramatically change in response to environmental factors, including controllable factors such as chemotherapy, diet, and obesity and immutable factors such as race. Future research elucidating just how these factors play a role in macrophage function and cancer progression are crucial for refining treatment strategies.



FIGURE 3

Macrophages support tumor cell migration and metastasis in many ways. (A) Macrophages promote EMT in tumor cells, which confers a migratory phenotype. They regulate ECM remodeling by enhancing both ECM stiffness (desmoplasia) (B) and degradation (C), which supports tumor cell migration and metastasis. Macrophages also provide signals that promote tumor cell invadopodium formation and directional migration towards blood vessels (D). Finally, macrophages participate in TMEM doorway-mediated tumor cell intravasation (E), which allows tumor cells to disseminate throughout the body. Figure created with BioRender.com.

Author contributions

MF: Writing – original draft, Writing – review & editing. GK: Writing – review & editing. JC: Writing – review & editing. MO: Writing – review & editing. DE: Writing – review & editing.

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Conflict of interest

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