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Neutrophils in the tumor microenvironment

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Abstract

Neutrophils are the first responders to sites of acute tissue damage and infection. Recent studies suggest that in addition to neutrophil apoptosis, resolution of neutrophil inflammation at wounds can be mediated by reverse migration from tissues and transmigration back into the vasculature. In settings of chronic inflammation, neutrophils persist in tissues, and this persistence has been associated with cancer progression. However, the role of neutrophils in the tumor microenvironment remains controversial, with evidence for both pro- and anti-tumor roles. Here we review the mechanisms that regulate neutrophil recruitment and resolution at sites of tissue damage, with a specific focus on the tumor microenvironment. We discuss the current understanding as to how neutrophils alter the tumor microenvironment to support or hinder cancer progression, and in this context outline gaps in understanding and important areas of inquiry.

Neutrophils at the crossroads of inflammation and cancer

Neutrophils are the most abundant circulating leukocyte and are the first responders to sites of infection and tissue damage. The primary function of neutrophils is to mediate host defense through multiple mechanisms including phagocytosis and intracellular killing of pathogens, release of granules containing antimicrobial peptides and proteases, and the formation of neutrophil extracellular traps (NETosis). Neutrophils are highly motile and display rapid recruitment to a variety of signals including chemokines, lipid mediators, and pathogen signals to mediate host defense (reviewed in [1-3]). However, neutrophils are not just “killing machines” but play a key role in orchestrating the innate and adaptive immune responses by releasing cytokines and chemokines and through antigen presentation [1, 2]. Indeed, neutrophils are much longer lived than initially thought and can survive for 5 or more days in the circulation [4] and may potentially live even for weeks in tissues. While it is well known that neutrophils are crucial for normal host defense and survival [5], uncontrolled neutrophil activation can contribute to chronic inflammation and tissue damage. Due to the balance needed for proper host protection and tissue homeostasis, understanding how neutrophils are recruited and subsequently resolve inflammation is an

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important question with broad implications including understanding the role of neutrophils in tumor progression.

The tumor microenvironment is characterized by persistent inflammation, and is often referred to as the “wound that does not heal” [6]. It has been known for some time that neutrophils are present in the tumor microenvironment; however, their role in tumor biology including tumor progression and invasion has remained controversial with both detrimental and beneficial effects reported. Several recent studies have highlighted the role of neutrophils in cancer biology using different cancer models in both mice and more recently zebrafish (see Box 1). It is known that the presence of neutrophils in tumors often correlates with poor patient outcome in humans; however whether the presence of tumor-associated neutrophils (TANs) directly contributes to disease progression is unclear. There is substantial evidence for a pro-tumor role for neutrophils in cancer progression. For example, a study by Bekes et al. showed that neutrophils produce MMP9 within the tumor microenvironment and this contributes to angiogenesis, tumor progression, and metastasis in mouse transplantation models [7]. Additionally, inhibition of myeloid cell recruitment into tumors with CXCR2 inhibition increases the efficacy of chemotherapy in breast carcinoma models, suggesting that targeting neutrophil recruitment may be beneficial [8]. By contrast, other studies have suggested that neutrophils can play an anti-tumor role by activating the immune response against tumors and promoting tumor cell clearance [9]. Indeed, neutrophils display plasticity and can be polarized into either an anti-tumoral (N1) or pro-tumoral (N2) phenotype depending on environmental factors [10]. Here we review these and other recent studies that have revealed mechanisms of neutrophil recruitment and resolution at sites of tissue damage, with a specific focus on the tumor microenvironment and how neutrophils may contribute to cancer progression.

Neutrophils and tissue damage

Neutrophil recruitment to tissue damage

There are common mechanisms that mediate neutrophil recruitment to wounds and cancer. In the case of the wound response, many signals mediate neutrophil recruitment to damaged tissues including DAMPs (Damage-Associated Molecular Pattern molecules) and chemokines (Figure 1) [11]. Wound induced recruitment signals have been elucidated using both mouse and zebrafish model systems (Box 1). In the zebrafish model, one of the earliest attractants after wounding is a hydrogen peroxide burst that is released by damaged tissue [12]. This is sensed by neutrophils through the Src family kinase, Lyn, which is required for early neutrophil response to wounds in zebrafish [13]. Other signals include the release of lipid mediators like LTB4. In mouse wounding models, LTB4 has been shown to elicit neutrophil swarming to areas of cell death in damaged tissues [14]. Neutrophils at the site of inflammation also secrete LTB4, further amplifying neutrophil recruitment. Necrotic tissue after sterile injury has also been shown to release ATP, activating the NLRP3 inflammasome via the P2X7 receptor that can mediate a chemokine gradient of CXCL2 (MIP-2) and CXCL1 [15]. A novel pathway in mouse liver injury showed that CXCL2 is regulated by TLR2 and S100A9 and mediates neutrophil recruitment [16]. Moreover, CXCL1 and G-CSF were shown to locally recruit neutrophils to areas of heart injury [17].

Additionally, chemokines like CXCL8/IL-8 recruit neutrophils via the CXCR1 and CXCR2 chemokine receptors to tissue damage [18]. Other G-protein coupled receptors are also involved in neutrophil recruitment. For example, in liver inflammation and sterile injury, the formylpeptide receptors Fpr1 and Fpr2 mediate neutrophil recruitment [15, 19]. While many pathways have been identified that mediate neutrophil recruitment to wounds, the temporal and spatial relationship between these different pathways in the context of tissue damage remains poorly understood.

The role of neutrophils at the wound, beyond clearance of pathogens and host defense, still remains unclear. Tissue damage and hypoxia can produce VEGF-A which recruits neutrophils expressing the pro-angiogenic matrix metalloproteinase MMP9, contributing to revascularization of the wounded tissue [20]. MMP9 is also important for remodeling of collagen after tissue injury to promote tissue repair and regeneration [21]. Neutrophils also recruit additional immune cells such as macrophages, which phagocytose apoptotic neutrophils and cellular debris [22]. This is important for resolution of inflammation at wounds, which is a key step in tissue repair. Neutrophils are also important for turning down the hydrogen peroxide burst through the delivery of myeloperoxidase (MPO) [23]. In chronic inflammation, neutrophils impair wound healing. A recent study in a diabetic mouse model showed that formation of neutrophil extracellular traps (NETs) impairs wound healing and that by disrupting NETosis, wound healing was enhanced [24]. On the other hand, neutrophil NETs may also limit inflammation by leading to the degradation of chemokines [25]. Further progress in characterizing the role of neutrophils at wounds will likely increase our understanding of the role of neutrophils in the tumor microenvironment.

Resolution of neutrophilic inflammation by reverse migration

Once infection and tissue debris have been cleared, it is important for neutrophil-mediated inflammation to resolve to prevent further tissue damage. It has long been known that resolution of neutrophil infiltration in tissues involves neutrophil apoptosis and their subsequent clearance by macrophages (reviewed in [22, 26]). However, more recent work has shown that neutrophils can also leave sites of tissue damage and undergo reverse migration back into the blood stream (Figure 1). Neutrophil reverse migration was first suggested in a rat model of glomerular inflammation [27] and was subsequently observed in human neutrophils co-cultured with endothelial cells in vitro [28]. The full process of neutrophil reverse migration was first visualized in vivo in the transparent zebrafish using live imaging of neutrophils after wounding [29]. In more recent studies, neutrophil reverse transmigration has also been live-imaged in mouse models [30]. Interestingly, in zebrafish, macrophages can induce reverse migration of neutrophils from a wound site by ROS-Src family kinase signaling, suggesting that macrophages may modulate resolution of neutrophil-mediated inflammation through both phagocytosis and/or a repellent mechanism [31]. In mice, recent evidence demonstrates neutrophil reverse transendothelial migration is regulated by an LTB₄-neutrophil elastase signaling axis [32], suggesting that similar cues may direct both recruitment and reverse migration. Using microfluidic devices, reverse migration, or retrotaxis, of human neutrophils has also been demonstrated [33, 34]. These studies have shown that as many as 90% of human neutrophils are able to reverse migrate and that this process is enhanced in the presence of the anti-inflammatory lipid mediator,

lipoxin A4, or the antioxidant compound, Tempol. There is some speculation that activated neutrophils undergoing reverse migration may lead to systemic activation of the immune response but this possibility remains to be studied. Work in zebrafish has shown that while neutrophils that have reverse migrated exhibit an activated morphology, they mount a normal response to a secondary insult [35]. This is important as neutrophils survive longer than previously thought in humans and these activated neutrophils may be able to respond to multiple tissue insults in the context of chronic inflammation or cancer.

Importantly, if neutrophils fail to resolve inflammation, acute inflammation can become chronic inflammation, leading to a cycle of tissue damage. Chronic neutrophil-mediated inflammation is observed in many diseases, including COPD, rheumatoid arthritis, diabetes, and others which are reviewed in detail elsewhere [36-40]. Importantly, the role of neutrophils in acute inflammation, as well as the transition from acute to chronic inflammation can potentially yield clues as to the role of neutrophils in cancer. Tumor progression can be considered a type of chronic inflammation where there is persistent neutrophil presence, and understanding neutrophil reverse migration and wound-healing in this context may provide a framework that is informative for cancer biology.

Neutrophils in cancer

Evidence of neutrophils' pro-cancer role

The role of neutrophils in cancer remains controversial, and is likely to be context dependent. It has long been known that neutrophils are present in many different types of cancers. For example, neutrophils are within the tumor or surrounding tissues in many cancers including glioblastoma, renal cell carcinoma (RCC), melanoma, colorectal cancer, hepatocellular carcinoma (HCC), pancreatic ductal carcinoma, and head and neck cancer, among others [41-49]. Many of these studies, including a recent meta-analysis [50], suggest that enhanced levels of neutrophils within the tumor tissue, or tumor-associated neutrophils (TANs), correlate with poor patient prognosis. It is important to note that it is not clear whether neutrophils themselves contribute to poor patient outcomes or if they merely correlate with a more aggressive disease phenotype. Recent studies have begun to address the role of neutrophils within the tumor microenvironment, revealing both beneficial and detrimental effects.

Neutrophil recruitment to tumors

Similar to acute wound responses, tumor cells and the surrounding microenvironment produce cues that actively recruit neutrophils. These signals include chemokines and cytokines, many of which are also important for responses to acute wounding (Table 1 and Figure 2). Neutrophils express the chemokine receptors CXCR1 and CXCR2 and these receptors are important for chemotaxis [51]. Cancer cells express various ligands for these receptors that facilitate recruitment of TANs. Mouse T cell lymphoma and Lewis lung carcinoma (LLC) cells express the Liver X Receptor (LXR) ligand oxysterols which have been shown to recruit neutrophils through CXCR2 [52]. Tumors also express a host of chemokines and cytokines, including CXCL8, CXCL5 and CXCL6 among others, which are involved in neutrophil recruitment [53]. CXCL5 was found to recruit neutrophils in a mouse

model of HCC, and high levels of CXCL5 correlate with poor prognosis in human HCC patients [54]. Interestingly, the CXCL5/CXCR2 axis was also found to be involved in epithelial-to-mesenchymal transition of HCC cells and contributes to tumor invasion [55], although it is not known if this is directly linked to increased neutrophil infiltration. Cytokines like TNF α have also been implicated in the recruitment or persistence of neutrophils in the tumor microenvironment, such as in a mouse skin carcinogenesis model [56]. The cytokine IL17 produced by gamma delta T cells enhances neutrophil recruitment and promotes tumor growth and metastasis in two separate mouse models of metastatic breast cancer [57, 58], suggesting that tumor-infiltrating lymphocytes can also modulate neutrophils in the tumor microenvironment. One very interesting connection between inflammation and cancer is what happens when there is chronic wounding or when wounding occurs near neoplastic cells. Using a zebrafish model of chronic wounding it was recently demonstrated that wound-induced inflammation leads to transformation of nearby melanocytes. Moreover, inflammatory cells that are recruited to a wound can influence transformed cells by increasing their proliferation [59]. This has important clinical implications as tumor biopsies are routinely used for diagnosis, and this inflammatory wound environment could have a detrimental impact on disease progression. Other work using zebrafish has demonstrated that, like in wounds, hydrogen peroxide is one of the initial signals that recruit leukocytes to transformed cells [60]. While it is known that these and many other chemotactic signals are upregulated in cancer, the direct role of these signals in recruiting neutrophils to the tumor microenvironment requires further investigation.

Role of neutrophils in tumor progression, angiogenesis, and invasion

There has been recent progress to identify potential mechanisms for how neutrophils contribute to cancer progression. It is likely that TANs play a role in cancer progression by releasing factors that modulate the extracellular matrix (ECM) and inflammation in the tumor microenvironment. More recent evidence also suggests that neutrophils may have more direct effects on cancer cell proliferation and invasion (Figure 2). Neutrophils release granules containing neutrophil elastase (NE), neutrophil collagenase (MMP8), and gelatinase B (MMP9), factors that remodel the ECM and affect cancer progression [61]. For example, MMP9 is produced by neutrophils and affects keratinocyte proliferation and invasion in skin cancer models [62]. While it is important to note that tumor-associated macrophages are also likely a source of MMP9 within the tumor microenvironment, it has been demonstrated that inhibition of neutrophil infiltration in a chick spontaneous metastasis model decreases tumor angiogenesis and dissemination, and the effect was rescued by neutrophil production of MMP9 [7]. Neutrophils can also enhance tumorigenesis through the release of reactive oxygen species (ROS) and reactive nitrogen species (RNS) which likely contribute to DNA damage and genetic instability in chemically induced carcinogenesis models [63-65]. Importantly, neutrophils produce growth factors like hepatocyte growth factor (HGF), which influences the invasion of human pulmonary adenocarcinoma cells [66]. Interestingly, HGF signaling through the Met receptor also can recruit neutrophils that have anti-tumor effects [67] suggesting a potential feedback loop mediated by HGF signaling. Neutrophils have also been shown to release cytokines like Oncostatin M, a member of the IL-6 family, which induces VEGF production and increases angiogenesis and tumor cell invasion in breast cancer [68], as well as affecting infiltrating

neutrophils in lung cancer [69]. Moreover, it was reported that neutrophil release of MMP9 increases VEGF, while inhibiting neutrophils impairs the VEGF-induced angiogenic switch in a mouse pancreatic cancer model [70]. In a zebrafish xenograft model, VEGFR inhibition reduces tumor vascularization but also enhances neutrophil infiltration and promotes tumor invasion, potentially through effects on collagen remodeling [71]. In another zebrafish model of oncogene-transformed cells, the presence of neutrophils was observed to enhance epithelial-to-mesenchymal transition of keratinocytes [72] at least in part through CXCR2 signaling, suggesting that neutrophils may directly affect the invasive behavior of transformed cells. Neutrophils also produce prostaglandin E2 (PGE2) which supports proliferation of early transformed cells [73], supporting the beneficial effects of targeting COX2-PGE2 in cancer prevention. In an in vitro co-culture model of RCC cells, neutrophils enhance the migration and invasion of cancer cells by upregulating estrogen receptor β , VEGF α , and HIF-2 pathways, providing further insight into how neutrophils affect cancer progression [74]. Although there has been recent interest and progress, there remain significant gaps in understanding how neutrophils affect cancer cell invasion and progression, in part because these effects are likely dependent on factors such as tumor type and stage of progression.

Neutrophils are also involved in tumor immunity by orchestrating the activity of other components of the immune response. In addition to proinflammatory factors, neutrophils produce factors that suppress anti-tumor immunity, similar to the effects described for granulocytic myeloid-derived suppressor cells (G-MDSCs, see Box 2 for further discussion). One such factor is arginase 1, an inhibitor of T cell function, which is produced by neutrophils in response to CXCL8 signaling [75]. Accordingly, depletion of neutrophils in mouse lung tumors results in increased activation of CD8⁺ T cells and decreased tumor growth that can be mediated by TGF- β [10]. However, in some contexts, such as early-stage human lung cancer, TANs are not necessarily inhibitory to T cell function but can instead promote T cell mediated immunity through the production of OX-40L and 4-1BBL costimulatory molecules which enhance proliferation of CD4⁺ and CD8⁺ T cells and increase their cytotoxic abilities in early stage lung cancer [9], further demonstrating the complexity of TAN crosstalk within the tumor microenvironment. It is an intriguing idea that TANs may differentially affect anti-tumor immunity depending on the stage or type of cancer. An additional element is the possibility that neutrophils may act as antigen presenting cells to promote the targeting of cancer cells by cytotoxic T cells, similar to what has been reported in response to microbial environments [76] or in the presence of chemokines (recently reviewed here [77]).

One new avenue of research focuses on the role of neutrophil extracellular traps (NETs) in cancer progression (recently reviewed in [78]). There is evidence that the presence of NETs within tumors correlates with poor prognosis [79]. In various mouse models of cancer, it has been seen that tumor cells can increase NET production by TANs, and potentially circulating neutrophils as well [80]. NETs appear to have a pro-tumor effect both by enhancing proliferation of cancer cells and by inhibiting apoptosis [79-82], however the precise mechanism by which NETs enhance tumor progression is unclear. One potential mechanism is that the release of NETs is associated with the release of pro-tumor factors including MMP9, NE, and cathepsin G [78]. As NETs are traditionally involved in the

capture of pathogens, it has also been suggested that NETs may enhance adhesion and metastasis of escaped circulating tumor cells. Importantly, inhibition of NETs has been shown to decrease adhesion of lung carcinoma cells and formation of metastases [83]. By contrast, a recent study of chronic inflammation in mice suggests that the formation of NETs by high density neutrophils can degrade pro-inflammatory cytokines and chemokines and alleviate inflammation [25]. While this role has not yet been addressed in the context of cancer, it suggests once again that neutrophils can play a dual role in inflammation and cancer. A caveat to the study of NETs' role in tumor progression is that it can be difficult to distinguish NETs, which are frequently identified by DNA stains or histone labeling, from the DNA of other dead cells including neutrophils themselves. However, co-labeling with proteins such as NE can help to clarify the presence of NETs. As live-imaging techniques advance and other models, including zebrafish (Box 1) are developed, it may be possible to better identify NET producing neutrophils within the tumor microenvironment through observation of their behavior and NET dynamics.

Targeting neutrophil-mediated inflammation in the tumor microenvironment

There is increasing interest in the idea that targeting neutrophils may be a useful therapeutic approach to affect cancer progression. For example, inhibition of CXCL8-CXCR1/2 signaling by CXCL8 antibodies [7] or small-molecules targeting CXCR1 and/or CXCR2 [84-86] has been shown to decrease tumor growth and progression in mouse models of cancer. Importantly, CXCR2 inhibition in a mouse metastatic breast cancer model enhanced response of both tumor and micrometastases to chemotherapy treatment, which was attributed at least in part to the dampening of the chemoresistant, pro-myeloid and granulocyte recruitment effects mediated by CXCL1/2 signaling [8]. Additionally, anti-CXCL6 antibodies impaired neutrophil recruitment to melanoma in mouse models and affected tumor growth [87]. Inhibitors of NE are also being tested and have shown some promise in mouse models of lung cancer [88], however in some cases such as the K14-HPV16 mouse model of squamous cell carcinoma, knockdown of NE had no effect on tumor development [89]. However, targeting neutrophils can be associated with side effects as neutrophils are critical for host defense against infection. An intriguing target for future studies is the idea that neutrophil resolution of inflammation could be induced within the tumor microenvironment. In particular, the idea that neutrophilic inflammation resolves by neutrophil reverse migration provides an avenue to promote resolution rather than blocking recruitment to sites of infection or tissue damage. Moreover, reverse migrated TANs may provide a mechanism for systemic activation of the immune response to cancer, either by direct stimulation of peripheral cytotoxic T cells or antigen presentation. The identification of drugs that promote neutrophil reverse migration from wounds [90] raises the idea that drugs that promote neutrophil reverse migration and resolution in the tumor microenvironment may be developed. Another intriguing possibility is the use of nanoparticles to target TANs. This has recently been described in a mouse model of inflammation to block neutrophil adhesion [91]. In addition, since wound associated macrophages can have neutrophil repelling functions [31], it is an interesting idea that

macrophages could be engineered to promote resolution of neutrophil inflammation in the tumor microenvironment.

A key caveat in blocking neutrophils in the tumor microenvironment is that not all neutrophils have pro-tumor effects. Recent studies suggest that neutrophils exhibit substantial plasticity and can be polarized to an N1 anti-tumoral or N2 pro-tumoral phenotype in response to the microenvironment, reminiscent of the M1/M2 polarization of macrophages [10, 92]. Tumor-associated N2 neutrophils are characterized by high expression of CXCR4, VEGF, and gelatinase B/MMP9 and can be induced upon exposure to high TGF- β levels. By contrast, N1 neutrophils are induced upon TGF- β blockade and express immunoactivating cytokines and chemokines, low levels of arginase, and are able to kill cancer cells [10]. Additionally, there is recent evidence suggesting that neutrophils from certain healthy donors are able to kill cancer cells and oncogene-transformed cells in a cell-specific manner [93], and that neutrophil killing of cancer cells may be enhanced by β -glucan treatment [94] making neutrophils a compelling candidate for cancer immunotherapy. Indeed, several studies that induce neutrophilia via prolonged G-CSF treatment in tumors demonstrate a shift from a chronic to acute inflammatory environment and an anti-cancer effect [95]. Understanding how neutrophils are polarized to a pro- or anti-tumor phenotype, and if and how they can be reprogrammed to switch this fate, will be crucial to developing successful cancer therapies.

Additionally, neutrophils are now being used to aid in the diagnosis of cancer and to help guide therapeutic decisions. Blood counts such as the neutrophil-to-lymphocyte ratio (NLR) predict patient outcome and can be used to measure response to treatment, where high NLRs correlate with poor prognosis and failure to respond to treatment [96]. Recently, NLRs have also been used to help differentiate primary breast carcinoma from benign proliferative breast disease [97]. NLR has also recently been applied for early detection and as a prognostic marker in gastric cancer [98], early diagnosis of ovarian cancer [99], and prognosis and survival prediction in colon cancer [100] and HCC [101], among others. The use of NLRs and other systemic inflammation markers could help not only aid in treatment choices and prognostic predictions, but also potentially enhance early detection of cancer. Additionally, some of the factors released by neutrophils in the tumor microenvironment can be used as biomarkers during early cancer detection. For example, neutrophil gelatinase-associated lipocalin (NGAL) is a strong biomarker for early pancreatic cancer [102] and ovarian cancer detection [103]. Identification of further TAN proteins could identify other biomarkers for the early diagnosis and characterization of cancer.

Concluding Remarks

There has been recent progress in defining the role of neutrophils in cancer progression, however gaps remain in understanding neutrophil plasticity and the switch between pro- and anti-tumor effects (Outstanding Questions). Recent studies have provided clues as to how neutrophils are recruited to the tumor microenvironment and what roles they play there, including remodeling of the ECM, promoting angiogenesis, and mediating interactions with other cell types including epithelia, stroma, and immune cells. It is clear that neutrophils often play a pro-inflammatory, pro-tumoral role within the tumor microenvironment. As the

presence of neutrophils is a well-characterized phenotype of aggressive disease, it will be important to understand how neutrophil inflammation is resolved following tissue damage and how failure to resolve in conditions of chronic inflammation contributes to cancer initiation and progression. In addition, identification of factors that influence neutrophil recruitment to target tissues and drive neutrophil reverse migration or resolution could lead to novel therapies to reduce neutrophil-mediated inflammation in tumors. However, methods to engineer neutrophils to an anti-tumor role in cancer are also an interesting avenue for future research. In particular, as cancer immunotherapy comes to the forefront of cancer treatments, a better understanding of how neutrophils regulate other immune cells within the tumor microenvironment and how this can influence prognosis will be needed. It is evident from recent work that neutrophils play a far more complex role than previously appreciated, and that further study of the role of neutrophils in wounding and inflammation will provide clues and guide research on the role of neutrophils in tumor progression with the hope of aiding and advancing cancer diagnosis and treatment.

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Box 1: Zebrafish as an emerging model to study neutrophils and cancer

With the emergence of new evidence that neutrophils play a role in cancer progression, researchers are identifying novel tools to gain more insight into the interaction between neutrophils and tumor cells. One of these tools is the larval zebrafish. While the zebrafish model system has traditionally been used to study developmental processes, the optical clarity of zebrafish embryos and larvae allow for unparalleled imaging of cellular processes and cell-cell interactions in an *in vivo* model. Many studies have taken advantage of easy and rapid genetic manipulations to drive oncogenic transformation of various cell types or to create mutants in tumor suppressor genes and generate cancer models. In addition, xenografts of human tumor cells into an immunocompetent zebrafish larval host may also be performed due to the delayed development of the zebrafish adaptive immune system (the use of zebrafish as a cancer model is further reviewed here [104, 105]). Xenografted cells can be labeled and tracked within a zebrafish larva over long periods of time allowing for visualization of tumor cell invasion [71, 106]. Additionally, primary human tumor cells can be readily transplanted in zebrafish tissues, allowing for robust, patient-specific studies of cancer cells within a host microenvironment [107]. With the generation of fluorescently labeled transgenic lines, high resolution live-imaging of neutrophil recruitment and interactions with cancer or transformed cells has allowed researchers to begin to uncover the molecular mechanisms underlying neutrophils' regulation of tumor progression. Zebrafish are an especially useful tool for analyzing neutrophil and other immune cell responses to areas of initial cell transformation that are not easily accessible in a mouse. Several key studies have demonstrated that neutrophils are specifically recruited to early transformed cells within a matter of days following expression of an oncogene and that neutrophils dynamically interact with these cells and promote EMT and invasion [60, 72, 73]. Importantly, the zebrafish system is ideal for large-scale, high-throughput drug screening which may help identify new therapies for cancer [108] and new anti-inflammatory drugs that may inhibit neutrophil recruitment to tumors and/or promote their resolution [90]. Using xenografts of patient-derived tumor cells in zebrafish larvae, it is becoming increasingly feasible to perform patient-specific drug screens and analyze tumor growth and proliferation, invasion, metastasis, and immune cell infiltration to help guide patient treatment in personalized medicine.

Box 2: Neutrophils and G-MDSCs

Myeloid derived suppressor cells (MDSCs) were initially described in 2007 and comprise a heterogeneous population of immature myeloid cells with immunosuppressive phenotypes [109]. There are two subpopulations of MDSC, the monocytic M-MDSCs, and the granulocytic G-MDSCs. Due to similarities in cell surface markers used to distinguish neutrophils and G-MDSCs, there is some confusion and controversy in the field as to whether these are in fact two separate populations of cells or are two different phenotypes or polarities of the same cell type. While this subject is discussed extensively in several recent reviews [110-112], it is important to note some of the key studies focused on tumor-associated granulocytes. Neutrophils and G-MDSC populations are routinely identified in mice using the same markers, for example Gr-1, CD11b, and Ly-6G, making direct comparison of the two populations challenging. In addition, there are several similarities in behavior and function for neutrophils and G-MDSCs, including production of ROS, antigen presentation, and immune suppression. A study directly comparing neutrophils and G-MDSCs in mouse models with and without tumors described neutrophils as mature, highly phagocytic cells that express high levels of proteases and TNF- α , and G-MDSCs as immunosuppressive cells that express high levels of ROS, arginase, and MPO [113]. Additionally, comparison of gene expression has suggested that neutrophils from tumor-free mice and G-MDSCs from tumor-bearing mice are more closely related to each other than to TANs [114], suggesting that the tumor microenvironment may influence TANs to adopt a unique phenotype. Indeed, recent analysis by Sagiv et al. using circulating blood neutrophils from both mouse mammary tumor models and human lung and breast cancer patients demonstrated that there are phenotypically diverse subpopulations of neutrophils that can be separated based on density [115]. The authors describe a population of mature, high density neutrophils which appear to switch to a higher proportion of low density neutrophils, characterized by immune suppression and pro-tumor properties, as cancer progresses. This switch is TGF- β dependent and is reminiscent of N1/N2 polarization, which is also regulated by TGF- β expression in mouse lung cancer models [10]. Sagiv et al. describe G-MDSCs as immature neutrophils; however, it is also likely that G-MDSCs may simply be an intermediary for N1/N2 phenotypic switching. Further defining G-MDSCs, especially in the context of N1/N2 polarity, is needed to provide consensus and clarity moving forward.

Trends Box

- Neutrophils are recruited to wounds and sites of tissue damage by signals including hydrogen peroxide, chemokines, and cytokines, many of which also recruit neutrophils to the tumor microenvironment.
- Neutrophils reverse migrate from target tissues through interactions with macrophages and other cell types, thus contributing to resolution of inflammation and promoting wound healing.
- Tumor-associated neutrophils contribute to tumor progression, invasion, and angiogenesis; however, there is evidence that neutrophils can play both pro- and anti-tumor roles and that they may be polarized to either phenotype based on external cues.
- Targeting neutrophils through blockade of pro-recruitment signals or driving reverse migration of neutrophils from tumors to promote an anti-inflammatory environment could provide therapeutic avenues for the treatment of cancer.

Outstanding Questions

- What are the signals driving neutrophil reverse migration and reverse transendothelial migration? Can these signals be used to exogenously drive reverse migration of neutrophils away from tumor tissues and/or areas of chronic inflammation?
- What mechanisms drive polarization of neutrophils to an N1 or N2 phenotype? Are these mechanisms cell or non-cell autonomous, i.e. are they purely extracellular signals or do neutrophils regulate their own polarity?
- Is TAN recruitment and polarity dependent on tumor type? Are certain tissues more prone to pro-inflammatory neutrophil infiltration?
- Does treatment with currently available anti-inflammatory drugs reduce tumor progression in cancer?
- Does neutrophil infiltration affect tumors differently at different stages of tumor development? Does neutrophil polarity change throughout the course of tumor progression?
- What are the mechanisms of neutrophil communication with other immune cells including tumor-associated macrophages and cytotoxic T cells? As macrophages have a demonstrated role in driving reverse migration of neutrophils during wound response, can tumor-associated macrophages similarly drive reverse migration of TANs?

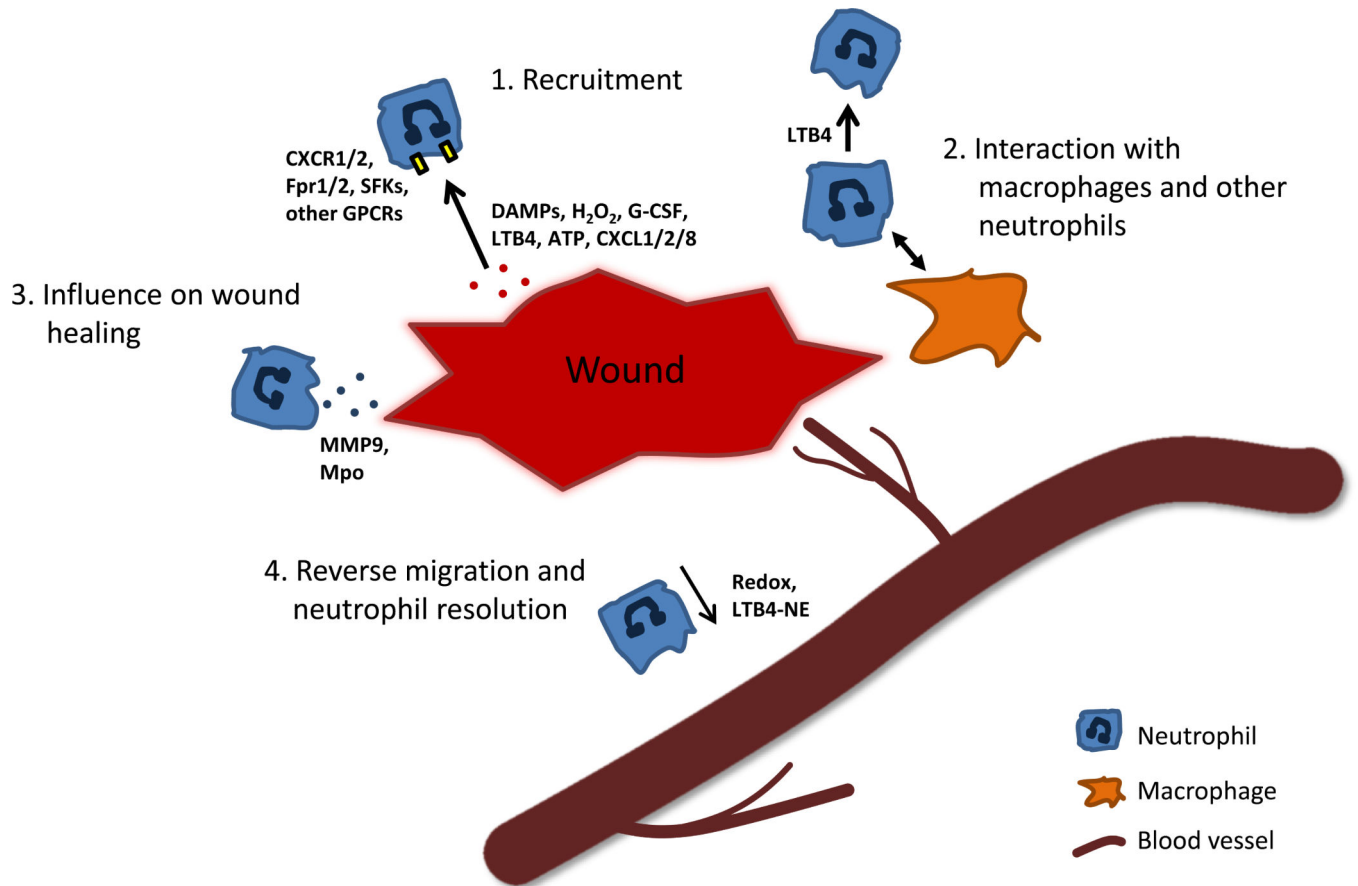


Figure 1. Neutrophils in the wound microenvironment

Neutrophils are recruited to areas of wounding and tissue damage via pro-inflammatory signals including DAMPs, hydrogen peroxide, lipid mediators, and chemokines (1). Neutrophils can further influence the innate immune response to wounding by recruiting additional neutrophils, as well as macrophages which in turn can phagocytose neutrophils or drive reverse migration (2). In addition, neutrophils influence the wound healing by releasing factors including MMP9 and myeloperoxidase (3). Reverse migration of neutrophils is mediated by Redox signaling and LTB4-NE signaling which promotes reverse transendothelial migration back into the blood stream (4).

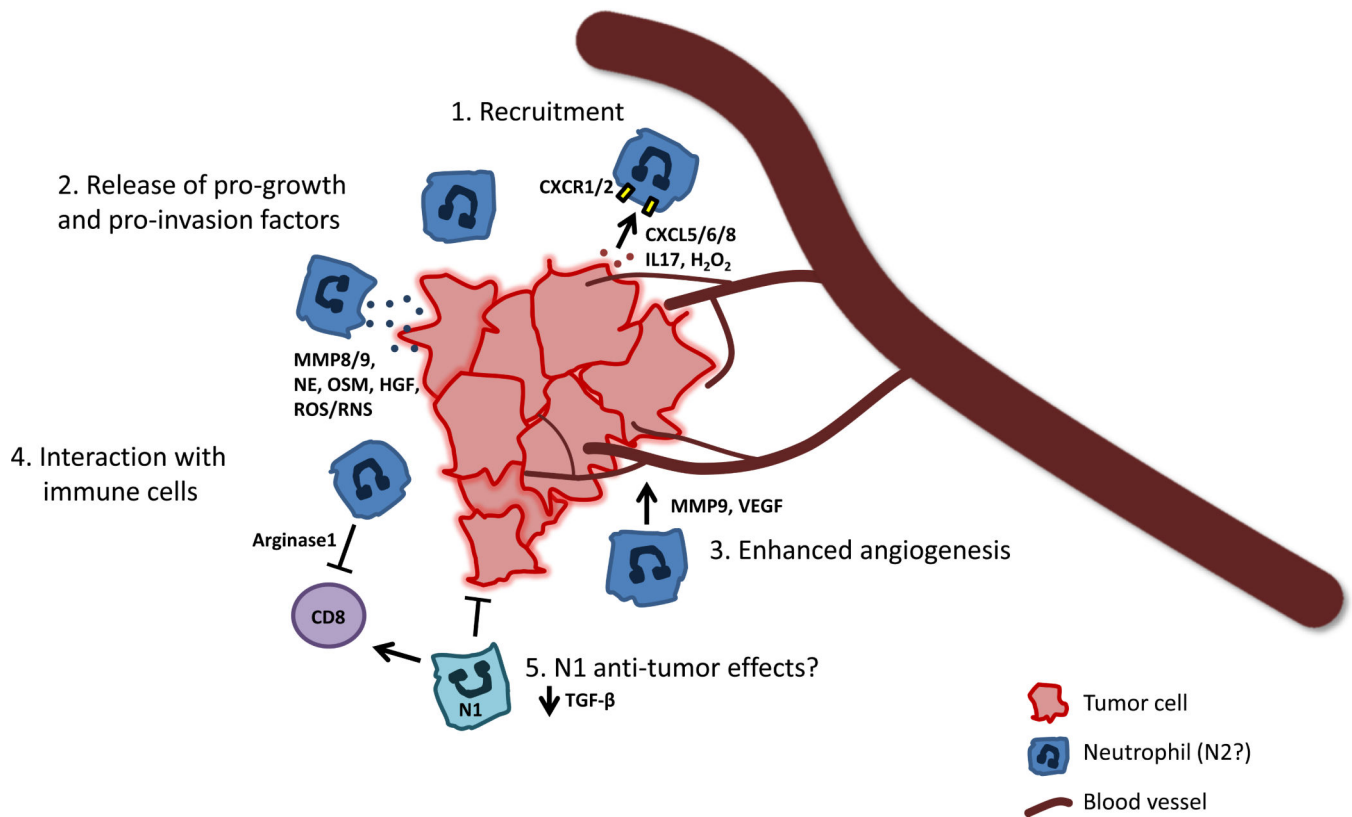


Figure 2. TAN interaction with the tumor microenvironment

Neutrophils are recruited to tumor sites via signals produced by cells of the tumor and microenvironment, including chemokines, cytokines, and hydrogen peroxide (1). These and other signals induce TANs to release factors which can remodel the ECM in the tumor microenvironment or act directly on tumor cells themselves to enhance tumor proliferation and invasion (2). In addition, some of these TAN-produced factors stimulate angiogenesis to support tumor growth and metastasis (3). Further evidence suggests that TANs interact with other immune cells such as CD8⁺ T cells. Depending on their polarity, TANs can have an immunosuppressive or immunostimulatory effects (4). While the TANs represented here are likely of the N2 pro-tumoral phenotype, the role of N1 neutrophils in targeting cancer cells and the N1/N2 polarization of neutrophils in response to cancer is an area of significant interest (5). Understanding how neutrophils are recruited and the complex ways they interact with cells within the tumor microenvironment will be important for the development of new therapies to treat cancer.

Table 1

Factors that mediate neutrophil recruitment and influence on the tumor microenvironment.

Function	Tumor process	Protein or Factor	Cancer/tissue type	Species	Reference
TAN recruitment, pro-tumor	Tumor growth, angiogenesis	Oxysterols	T cell lymphoma, LLC	mouse	52
TAN recruitment, pro-tumor	EMT/invasion, poor prognosis	CXCL5	HCC	mouse, human	54, 55
TAN recruitment, pro-tumor	Tumor initiation	TNF α	skin	mouse	56
TAN recruitment, pro-tumor	Tumor growth, metastasis	IL17	breast cancer	mouse	57, 58
TAN recruitment, pro-tumor	Tumor growth	H ₂ O ₂	skin	zebrafish	60
TAN recruitment, pro-tumor	Tumor invasion and migration	HGF	pulmonary adenocarcinoma, LLC	human, mouse	66, 67
TAN recruitment, pro-tumor	Tumor invasion, angiogenesis	VEGF/VEGFR	pancreatic cancer, transformed endothelial cells	mouse, zebrafish xenograft	70, 71
Pro-tumor	Angiogenesis	MMP9	spontaneous mets model	chick	7, 70
Pro-tumor	Tumor invasion, angiogenesis	Oncostatin M	breast cancer, lung	human, mouse	68
Pro-tumor	EMT/invasion	CXCR2	skin	zebrafish	72
Pro-tumor	Tumor cell proliferation	Prostaglandin E2	skin	zebrafish	73
Pro-tumor	T cell inhibition	Arginase 1	NSCLC	human, mouse	75
Pro-tumor	Metastasis, chemoresistance	CXCL1/2, CXCR2	breast cancer	mouse	8
TAN recruitment, pro-tumor	Tumor growth, metastasis	CXCL6	melanoma	mouse	87
Pro-tumor	Tumor growth	NE	lung	mouse	88
Pro-tumor/Anti-tumor	T cell inhibition/activation, cytotoxicity	TGF-beta	lung	mouse	10
Anti-tumor	T cell activation (CD8+, CD4+)	OX-40L, 4-1BBL	lung	human	9