Neutrophils in the era of immune checkpoint blockade

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ABSTRACT
The immune checkpoint blockade-based immunotherapies are revolutionizing cancer management. Tumor-associated neutrophils (TANs) were recently highlighted to have a pivotal role in modulating the tumor microenvironment and the antitumor immune response. However, these cells were largely ignored during the development of therapies based on programmed cell death receptor or ligand-1 and cytotoxic T lymphocyte antigen-4 immune checkpoint inhibitors (ICIs). Latest evidences of neutrophil functional diversity in tumor raised many questions and suggest that targeting these cells can offer new treatment opportunities in the context of ICI development. Here, we summarized key information on TAN origin, function, and plasticity that should be considered when developing ICIs and provide a detailed review of the ongoing clinical trials that combine ICIs and a second compound that might affect or be affected by TANs. This review article synthesizes important notions from the literature demonstrating that: (1) Cancer development associates with a profound alteration of neutrophil biogenesis and function that can predict and interfere with the response to ICIs, (2) Neutrophil infiltration in tumor is associated with many features of resistance to ICIs, and (3) TANs play an important role in resistance to antiangiogenic drugs reducing their clinical benefit when used in combination with ICIs. Finally, exploring the clinical/translational aspects of neutrophil impact on the response to ICIs offers the opportunity to propose new translational research avenues to better understand TAN biology and treat patients.

INTRODUCTION
Immune checkpoint blockade is the latest revolution in the care of patients with cancer. Anticytotoxic T lymphocyte antigen-4 (CTLA-4) and antiprogrammed cell death receptor or ligand-1 (PD-1/PD-L1) antagonist antibodies, which work by inducing the receptor or ligand-1 (PD-1/PD-L1) antagonism and cytotoxic T lymphocytes (Th1) or by restoring the antitumor activity of exhausted CD8+ T cells,1 2 are giving promising results (long-term survival increase) in a subset (25%–40%) of patients with melanoma, lung cancer, colorectal cancer (CRC),3 and urothelial carcinoma.4 However, important challenges to improve the clinical benefit of these therapeutics, globally designated as immune checkpoint inhibitors (ICIs), concern the identification of predictive markers and the characterization of resistance mechanisms.5 6 Major efforts are made to decipher the immunosuppressive molecular and cellular mechanisms in the tumor microenvironment (TME), to better stratify patients who will benefit from ICIs, and to identify new targetable pathways to enhance their antitumor immune responses. In this context, myeloid cells and particularly neutrophils have recently emerged as major immune contributors to cancer progression and resistance to ICIs.

Neutrophils are frequently identified as CD66b+, CD15+CD14+, CD33+ cells in humans and as CD11b+Ly6G+Ly6C+ cells in mice.7 Although tumor-associated neutrophils (TANs) seem to be predominantly associated with bad prognosis and poor response to therapy in multiple solid tumor types,8 they also show some plasticity with important consequences on disease progression. Indeed, they can polarize toward tumor-promoting (N2-TAN) or antitumor (N1-TAN)9 cells in mice. Neutrophils in blood can be subdivided into normal-density neutrophils and low-density neutrophils (LDN) from gradient centrifugation. Immunosuppressive neutrophils from blood and tumors are also frequently designated as granulocytic-myeloid derived suppressor cells (G-MDSC) and more recently as polymorphonuclear-MDSCs (PMN-MDSC). In the most publications, PMN-MDSC and G-MDSC refer to a population of cells that express Ly6G in mice, CD66b and/or CD15 in humans able to inhibit T cell proliferation, cytokine secretion and cytotoxic activity in vitro. For a very detailed review on neutrophil markers in the context of cancer, see Jaillon et al.10 In mouse models of breast cancer, TANs can display protumor functions by reactivating dormant lung metastasis growth during inflammation.11 On the other hand, they can also have antitumor activity12 and protect against early stages lung cancer12 and breast cancer.
metastasis in mouse lung. Hence, TANs constitute a pool of cells with a broad range of activities and phenotypes. Their functional diversity is still poorly understood, but might be explained by signals coming from the TME, and probably also by other regulatory pathways involved in neutrophil biogenesis or differentiation.

We will see that in many circumstances, immunosuppressive and tumor-promoting neutrophils can negatively interfere with ICIs. To apprehend neutrophils biology in the context of ICI treatment, we decided to simplify the nomenclature of these cells by designating neutrophils present in the TME as ‘TANs’ and we use the term neutrophil for referring to cells present in the blood or the bone marrow.

Yet, TANs and their impact on the TME are usually ignored during the development and evaluation of ICI-based therapies, although they could help to explain the success or failure of many clinical trials. In this review, we propose that innovant therapies can affect TAN behavior, and consequently might deeply modify the care of patients with cancer by increasing ICI strength and success rate in the near future.

**SPECIFIC TAN EFFECTS AND RELATED TUMOR CHARACTERISTICS LINKING NEUTROPHIL WITH ICI RESPONSE**

Neutrophils and TANs are associated with key features of resistance to ICIs: (1) adaptive immune cell polarization and suppression, (2) tumor neoangiogenesis, (3) immune exclusion, and (4) cancer cell intrinsic characteristics (figure 1).  

**TANS AND TUMOR-INFILTRATING T LYMPHOCYTES**

Comparison of different cell lines generated from the KPC pancreatic ductal adenocarcinoma mouse model confirmed that the cell lines forming PD-1⁺ CD8⁺ T cell-rich tumors are sensitive to combination treatments containing gencatimibe, paclitaxel, anti-CD40, anti-CTLA-4, and anti-PD-1 molecules. Conversely, tumor-infiltrating T lymphocyte (TIL)-poor tumors are resistant and contain a higher fraction of TANs. Abolishing CXCL1 or CXCL5/CXCR2 signaling in TIL-poor tumors reduces TAN infiltration, increases PD-1⁺ CD8⁺ T-cell abundance, and sensitizes tumors to the anti-CD40, anti-CTLA-4, and anti-PD-1 combination immunotherapy (table 1). Similarly in breast cancer, TAN-rich tumors display lower macrophage and TIL infiltration and are resistant to anti-PD-1 treatment. Hence, the relationship between TANs and antitumor T cell response suggests the existence of a direct TAN-mediated suppression of Th1 and cytotoxic T lymphocytes in tumors. In this regard, arginase-1 (ARG1)-expressing human granulocytic cells leads to CD3ζ chain downregulation on T cells though L-arginine depletion and ultimately inhibit T-cell proliferation and cytokine secretion. In patients with non-small cell lung cancer (NSCLC), the presence of peripheral ARG1⁺ neutrophils increases with the tumor stage in treatment-naïve patients and negatively correlates with the proportion of CD8⁺ T cells. Similarly, in patients with gastric cancer, a population of CD11b⁺CD33⁻CD66b⁺ cells was identified in peripheral blood that suppress CD8⁺ cell activity. ARG1 expression has also been observed in peripheral CCR5⁺ LDNs from patients with melanoma. The proportion of such cells in peripheral blood mononuclear cells and their ARG1 expression increase with disease progression. It has been proposed that expression of CCL3, 4, and 5 in the tumor mass drives the recruitment of ARG1⁺CCR5⁺ LDNs in tumors. Interestingly, in the KrasG12D mouse model of lung cancer, ARG1 inhibition reduces tumor growth and increases T-cell homing and function in tumors. Besides ARG1 expression, the release of prostaglandin-E2 (PGE2) by neutrophils and TANs represents another important mechanism of T-cell suppression. Very recently, it was reported that FATP2 expression plays a critical role in PMN-MDSC and TAN-mediated T-cell suppression. FATP2 is involved in triglyceride-containing arachidonic acid uptake by PMN-MDSC, and consequently, in the production of immunosuppressive PGE2 through its degradation by the prostaglandin E synthase (COX2). FATP2-deficient LDNs show lower ability to suppress T-cell amplification in vitro, and FATP2 inhibition using lipofenorma reduces tumor growth and increases sensitivity to anti-CTLA-4 immunotherapy in the EL4, CT26, LLC, and TG-1 cancer cell-derived mouse models. TANs can also suppress T-cell proliferation and function, or induce their apoptosis through the production of reactive oxygen species (ROS) and nitric oxide (NO). Fas ligand, and TRAIL.

More information on the real impact of these pathways may be revealed by ongoing clinical trials. Indeed, ARG1 inhibitors (INCB001158 and pegzilarginase) are currently tested in combination with pembrolizumab. Celecoxib (COX2 inhibitor) is evaluated in combination with nivolumab and nivolumab plus ipilimumab. The PGE2-receptor inhibitor grapiprant is also tested in combination with pembrolizumab (table 1). Finally, the NO synthase inhibitor monomethyl-L-arginine (L-NMMA) is assessed in combination with pembrolizumab and durvalumab (table 1). Of note, expression of iNOS and ROS2, which are involved in NO and ROS synthesis respectively, is not a strict marker of tumor-promoting TANs because both proteins can also participate in cancer cell killing by TANs.

In addition to the direct inhibition of effector T cell functions, TANs have also been implicated in regulatory T cell (Treg) recruitment. In KP mice, neutrophil depletion, using an anti-GR1 antibody or the combination of anti-Ly6G antibody and a secondary antirat antibody, has been associated with Treg reduction in lung tumors. Similarly, CCR5⁺ARG1⁺PD1⁺ TAN infiltration is associated with increased Treg proportion in tumors from the human RET transgenic mouse model of melanoma. In this model, CCR5 inhibition reduces neutrophil recruitment in tumors and Treg proportion, and increases mouse survival. Finally, in the ApcΔ/Δ, Cdx2CreER2 mouse

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Model of CRC and in CRC samples from patients, TAN infiltration correlates with increased transforming growth factor-beta (TGFβ) signaling, whereas TAN depletion using the anti-GR1 antibody and CCR2 inhibitor combination results in increased T-cell homing in the tumor mass, characterized by a smaller proportion of Tregs. Interestingly, in this CRC model, TGFβ is mostly expressed by epithelial cells, monocytes and stromal cells, and the matrix metalloproteinase-9 (MMP9), which is abundantly expressed by TANs, converts pro-TGFβ into its active form. Consequently, inhibition of MMP9 and/or TGFβ receptor abolishes the neutrophil immunosuppressive and tumor-promoting functions.26

In conclusion, although TANs can exert a direct pressure toward TIL eradication through NO and ROS cytotoxicity, they can also suppress T-cell activation, cytokine secretion, and proliferation via L-arginine depletion or PGE2 production and drive Treg recruitment and perhaps expansion, establishing a T-antigen specific tolerance.
Table 1  Clinical trials combining immune checkpoint blockade and drugs that may affect resistance linked with neutrophil biology

<table>
<thead>
<tr>
<th>Target</th>
<th>Drug</th>
<th>Immune checkpoint inhibitor</th>
<th>Clinical trial</th>
<th>Cancer type</th>
<th>Phase</th>
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Continued
In the tumor mass, neoangiogenesis leads to the development of an immature microvascular network that shapes the TME architecture and composition and also contributes to disease progression and resistance to ICI treatment. TANs are considered a major source of growth factors involved in tumor angiogenesis, and their infiltration has been associated with resistance to anti-vascular endothelial growth factor (VEGF) therapy. The anti-VEGF and anti-angiopoietin-2 bispecific antibody vencizumab has been evaluated in a phase I clinical trial in combination with atezolizumab (anti-PD-L1 antibody) in patients with advanced cancer with acceptable safety profile and promising effect on tumor angiogenesis and cellular density were observed (NCT01688206). In pancreatic, breast, and brain mouse tumor models, treatment with anti-VEGFR and anti-PD-L1 antibodies together with a lymphotoxin-β receptor agonist led to the...
formation of high endothelial venules in tumors accompanied by cytotoxic lymphocyte infiltration. These findings linking angiogenesis to the TME have been reviewed elsewhere and suggest that the connections between TAM infiltration and tumor angiogenesis play an important role in ICI response.

We identified 111 clinical trials that are testing the combination of bevacizumab (Avastin; anti-VEGF-A antibody) with PD-L1/PD-L1 or CTLA-4 blockade. Therefore, it is important to determine the role of TAM infiltration in the response to this strategy. Mouse neutrophils can release VEGF-A on CXCL1 stimulation through HCK and FGR Src tyrosine kinase signaling, but it is not known whether human neutrophils can also do it. Moreover, TAM negative impact on bevacizumab treatment remains poorly understood. In nude mice xenografted with human cancer cell lines (A-673, Calu-6, HM7, HPAC, Jurkat), Bv8/prokineticin-2 is upregulated in bone marrow neutrophils. The Bv8 soluble factor induces endothelial cell proliferation and angiogenesis through the activation of EG-VEGFR/PRK1 and EG-VEGFR/PRK2 signaling. Importantly, inhibition of Bv8 signaling using a specific antibody reduces CD11b’/GR1’ myeloid cell mobilization in blood, TAM infiltration in tumors, and tumor angiogenesis in all tested models. Furthermore, in these models, Bv8 inhibition and anti-VEGF-A agents have a cumulative effect, demonstrating that both Bv8 and VEGF-A contribute to tumor neoangiogenesis. Considering TAM negative impact on bevacizumab efficacy, TAM infiltration should be specifically analyzed in clinical trials that combine ICIs and antibodies against VEGFR-2 (tanimibumab and ramucirumab), or VEGFR-2 inhibitors, such as apatinib. Multitargeted RTK inhibitors, such as pazopanib and sorafenib that inhibit PDGFR and VEGFR1–3 are currently tested in combination with ICIs. Importantly, sunitinib, which inhibits PDGFR and VEGFR-1–3 are currently tested in combination with ICIs. Hence, identifying the molecular differences between TAM-related and TAN-related tumor angiogenesis may help to determine their specific contribution to tumor growth and to treatment response in different solid tumor types (figure 2).

**TANS AND TUMOR CELL INTRINSIC CHARACTERISTICS**

The three major tumor-related diagnostic factors that are functionally associated with the response to ICIs are: (1) PD-L1 expression, (2) tumor mutational burden, and (3) specific oncogene/tumor-suppressor alterations in cancer cells.

Specifically, the type of oncogenic signaling and tumor-suppressor gene alterations differently affects the response (resistance or sensitivity) to immune checkpoint blockade in a way that might involve neutrophils. For example, deletion of the tumor suppressor LKB1/STK11 is associated with higher tumor-promoting TAM recruitment and immune checkpoint blockade resistance in lung cancer. The PTEN signaling pathway is also a great example of oncogenic pathway that links ICI resistance to TAM recruitment. In cancer cells, loss of PTEN is associated with phosphatidylinositol-3-kinase (PI3K)-AKT signaling overactivation that drives the expression of immunosuppressive cytokines, lowering T-cell recruitment, and enhancing TAM accumulation in the tumor mass. Therefore, PI3Kα/β inhibitors (duvelisib and GSK2636771) that target PTEN mutated cancer cells might also remodel the TME. Furthermore, the MET inhibitor capmatinib increases ICI efficacy in various solid cancer mouse models through alteration of neutrophil recruitment at sites with T cell-induced inflammation. A phase II clinical trial (NCT01439317) is currently testing the efficacy of pembrolizumab in NSCLC combined or not with capmatinib, irrespective of the cancer MET mutation status. Finally, clinical innovations may come from cobanztinib, a MET, RET, AXL, VEGFR2, FLT3, and KIT inhibitor that reprograms TANs toward antitumor activity. There are currently 48 and 47 ongoing clinical trials in which the pan-TKI cobanztinib is combined with PD-1/PD-L1 or CTLA-4 blockade, respectively. Preliminary observations from the clinical trials with cobanzitib are promising, possibly due to

Study in the KP mouse model of lung cancer showed that neutrophil depletion (using an anti-GR1 antibody) increases tumor vasculature coverage by alpha-SMA+ pericytes, reduces hypoxia, reverts tumor immune exclusion and sensitizes tumor to anti-PD-L1 therapy. In the CCR2-deficient mouse model of cervical cancer, tumor-associated macrophages (TAM) proportion is reduced compared with control tumors, and TANs support tumor angiogenesis in the absence of macrophages. Thus, when considering tumor angiogenesis, TAMS and TANS sometimes display overlapping functions through shared signaling mechanisms. Depending on the tumor models, infiltration can be predominantly by TAMs or TANS, and this will differentially affect the response to ICIs.

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the simultaneous targeting of cancer cell signal pathways (RET, MET, AXL), neutrophil function (MET, potentially KIT, and FLT3), and tumor angiogenesis (VEGFR2).52

TAN DIVERSITY AND RECRUITMENT AS THE CORNERSTONE TO UNDERSTAND ICI FAILURE IN SOLID TUMORS

While preexisting adaptive immune response in tumor is required for effective ICI response,44 53 TAN infiltration is predominantly associated with an adverse outcome across all cancer types53 and these cells present specific features linked with immunosuppression in tumors.10 21 Of note, approaches designed to inhibit neutrophil survival might trigger a compensatory neutrophil production in bone marrow, leading to the release of young neutrophils in the circulation, as our recent experiments in C56BL/6J mice treated with an anti-Ly6G antibody demonstrated.25 54 These newly generated neutrophils might develop opposite functions compared with those of mature neutrophils present in non-treated mice.55 Therefore, although neutrophil involvement seems obvious, it is often difficult to determine whether the observed biological effects are due to neutrophil number reduction (depletion) or increased renewal in tissues (turnover).25 54 Furthermore, association between neutrophils and response to ICIs is not restricted to the TME. Indeed, an alteration of neutrophil biogenesis and survival as reflected by an elevated absolute neutrophil count (ANC) in peripheral blood and an increased in the neutrophil-to-lymphocyte ratio (NLR) predicts the response to ICIs across multiple cancer types.56 57

Figure 2  TAN, tumor angiogenesis, and therapeutics used in combination with ICIs TANs and macrophages play a key role in tumor neoangiogenesis. Tumors are highly infiltrated by macrophages or neutrophils, frequently in a mutually exclusive manner. However, in mouse models, these two cell types can substitute each other during neo-angiogenesis. Neutrophils secrete PIGF, PDGF, VEGFA, and BV8 that directly contribute to neoangiogenesis. These cells also express MMP9 that plays an important role in neoangiogenesis. Neutrophil infiltration drives bevacizumab resistance, while the impact of this cell type on the response to anti-VEGFR2 antibodies (tanibirumab and ramucirumab) and small molecule inhibitors (sorafenib and pazopanib) remains to be characterized. Cabozantinib and lenvatinib, which block VEGFR1–3 and PDGFR and other tyrosine kinase receptors, inhibit neutrophil tumor-promoting function and neoangiogenesis. TANs promote the formation of an immature blood vessel network that lack pericyte coverage and that is not permissive to immune cell extravasation and recruitment in tumors. Ultimately, this contributes to immune exclusion. All the indicated drugs are currently tested in combination with ICIs. ICIs, immune checkpoint inhibitors; MMP9, matrix-metalloproteinase-9; PDGFR, platelet-derived growth factor receptor; PIGF, placental growth factor; TANs, tumor-associated neutrophils; VEGFA, vascular endothelial growth factor-alpha; VEGFR, VEGF receptor.
NEUTROPHIL DIVERSITY IN PATIENTS WITH CANCER

TAN phenotypic diversity in cancer was formally demonstrated by single-cell transcriptomic analysis in samples from patients with lung cancer and from KrasLSLG12D/WT; p53fl/fl (KP) mice with lung cancer.58 This study identified five populations of neutrophils in human tumors and six in mouse lung adenocarcinoma, and also six neutrophil clusters in blood samples from patients.58 Particularly, this analysis showed the existence of a subpopulation of human TANs characterized by the expression of peptidase inhibitor 3 (PPI3).58 This cell subpopulation displays a similar transcriptional profile as the SiglecF+ tumor-promoting TANs identified earlier in KP mice.59 Interestingly, this tumor-specific neutrophil cluster identified in mouse lung tumor is also characterized by elevated expression of LOX1/OLR1, a potential marker of human LDN in peripheral blood of patients with cancer.60 Hence, similarity and differences between mouse and human tumor-promoting TANs remains to be clarified however the observation listed above suggest that both mice and human tumor promoting TANs might share, at least partially, similar transcriptomic identity and origin.

While immunosuppressive TAN and circulating neutrophils (frequently named PMN-MDSC or G-MDSC) are suspected to originate from immature neutrophils,7 in KP mice, tumor-promoting SiglecF+ TANs are mature, non-proliferating and long-lived cells that sediment in both low-density and high-density gradient layers.61 Furthermore, we have recently shown that maintenance of SiglecF+ tumor-promoting TAN infiltration requires metabolic adaptation of these cells that associates with Glut1 induction and Glut3 loss in the KP mouse model of lung cancer. Conditional knockout of Glut1 in neutrophils increases TAN turnover and reduces accumulation of SiglecF+ TANs in the TME delaying tumor growth and sensitizing them to radiotherapy.62 These observations emanating mostly from mouse models suggest that tumor-promoting neutrophils, as exemplified by SiglecF+ TAN and immunosuppressive neutrophils are referring to different types of cells, each of them contributing to disease progression and refractoriness to treatment.

Although TANs seem to be predominantly associated with bad prognosis and poor response to therapy in multiple solid tumor types, they also show some plasticity with important consequences on disease progression. Mirroring a concept initially developed to describe tumor-associated macrophages (TAMs), neutrophils were shown to polarize toward an antitumor phenotype (N1) in presence of interferon (IFN), by opposition of the acquisition of immunosuppressive and tumor-promoting functions (N2) in presence of TGFB.9 However, usage of this denomination (N1 vs N2) became rare in latest publications and might not recapitulate the functional diversity of these cells in cancer patients. Nevertheless, TAN function was shown to be versatile in tumors ranging from antitumor to tumor-promoting activity along disease progression. TANs display antitumor activity at early stages of disease development through various mechanisms. In early-stages lung cancer TANs were shown to stimulate T-cell responses through the expression of the co-stimulatory molecules, prime tumor-antigen (T-Ag)-specific T-cell responses, and cross-present T-Ag to CD8+ T cells.63 In early-stages mouse model of uterine cancer neutrophils induce tumor cell detachment from the basement membrane reducing tumor growth,64 while in early-stage mouse model of CRC, neutrophils were proposed to restrict tumor-associated microbiota ultimately avoiding uncontrolled IL17-mediated inflammatory response and disease progression.65 Finally, tumor-suppressing neutrophils play a major role in trained immunity models. Indeed, mice treatment with β-glucan induces epigenetic modification of neutrophil progenitors and precursors driving the emergence of tumor suppressing neutrophils capable of controlling B16 melanoma tumor growth.66

These observations revealed that TAN diversity has to be taken in consideration while evaluating ICI treatments. Together with signals present in the TME, TAN diversity might also be orchestrated by a profound alteration of neutrophil biogenesis during cancer progression. Indeed, a study in the KP mouse model of lung cancer revealed that tumor cells remotely induce osteoblasts that regulate neutrophil biogenesis, leading to the accumulation of the SiglecF+ subpopulation of tumor-promoting TANs.69 Interestingly, the appearance of SiglecF+ neutrophils in a model of heart infarction was shown to be initiated in the bone marrow directly from neutrophil precursors,67 arguing that neutrophil plasticity during heart injury, as in tumor models, might originate from distal alteration of the neutropoiesis. Supporting a profound alteration of circulating neutrophil diversity in cancer patients, single-cell RNA sequencing comparing blood neutrophils from healthy donors and patients with NSCLC identified a cluster representing 8% and 40% of neutrophils in healthy donors and patients, respectively.68

Hence, TAN accumulation and diversity in tumors are the result of neutrophil biogenesis alterations (for review, see Bergenfetz and Leandersson7), intratumor polarization, and prosurvival signals present in the TME as well as of active recruitment mechanisms, each of them offering actionable targets to reduce TAN negative impact on the response to ICIs (figure 3).

NLR AS A MARKER OF THE RESPONSE TO ICIS

The NLR, which is based on the determination of the ANC and absolute lymphocyte count (ALC) in peripheral blood of patients with cancer, is a strong non-invasive predictive marker of the response to ICIs. At baseline, high NLR (>2.5) has been associated with lower progression-free survival and lower disease-free survival in patients with NSCLC,69-72 small cell lung cancer,73 and melanoma74 treated with ICIs. Furthermore, ANC increase during anti-PD-1 immunotherapy is an independent predictive factor of rapid disease progression in patients with advanced gastric cancer.75 Conversely, in patients with RCC77 and NSCLC78 a decrease of the NLR value at week six post-treatment with anti-PD-1/
PD-L1 antibodies is associated with better objective response rate and overall survival (OS). As a strong evidence of the important predictive value of NLR in patients treated with ICIs, a study performed in 509 patients with advanced melanoma, NSCLC, HNC, bladder cancer, or sarcoma treated with anti-CTLA-4 and/or anti-PD-1/PD-L1 antibodies found that pre-treatment NLR value <5 and a 10% to 28% decrease of the NLR value during treatment are associated with longer OS. Conversely, NLR increase during treatment correlates with shorter OS. Of note, NLR predictive value is not restricted to immune checkpoint blockade, but concerns also other antibody-based therapies, such as cetuximab (anti-EGFR antibody) and the anti-EGFR (cetuximab or panitumumab) and anti-VEGF (bevacizumab) antibody combination in patients with CRC. This is also true for conventional chemotherapy in patients with large-cell neuroendocrine lung cancer, hepatic cell carcinoma, and prostate cancer. While diminution of the ANC during treatment is associated with a better patient response to ICI in lung cancer, high baseline ALC associates with better response to ICI in HNC and patients with NSCLC. Hence, the NLR prognostic value comes from the cointegration of the ANC and ALC, which it is determined by biogenesis as well as survival of both neutrophils and lymphocytes (figure 3).

**Combining Immune Checkpoint Blockade and Inhibition of Neutrophil Recruitment in Tumors**

The expression of ELR⁺ CX-C chemokines (CXCL1, 2, 5, 6, and 8) and their specific receptors CXCR1 and CXCR2 represent the main axis regulating neutrophil recruitment in inflamed tissues and tumors. In agreement, CXCR1/2 inhibitors (SX-682 and navarixin) significantly inhibit TAN accumulation in tumor tissues and increase the efficacy of anti-PD-1 immunotherapy in mice. These results might be translated into the clinic thanks to a newly developed...
anti-IL8/CXCL8 antibody (BMS-986253) currently tested in a phase II clinical trial (NCT04050462) in combination with nivolumab (anti-PD-1 antibody) in patients with advanced hepatocellular carcinoma (HCC).

More recently, the expression of the CCR5 ligands CCL3/ MIP1α, CCL4/ MIP1β, and CCL5/RANTES also has been implicated in neutrophil recruitment to tumor tissues in patients with melanoma and mouse models as well as in patients with NSCLC. Therefore, several trials are assessing whether the combination of the CCR5 inhibitor vicriviroc with pembrolizumab (NCT03631407) or of the CCR2/5 inhibitor BMS-813160 with nivolumab (NCT03184870) could be more effective than immune checkpoint blockade alone by inhibiting TAN-mediated cancer promotion and immunosuppression.

TAN accumulation might be reduced also through inhibition of CXCR4 because it has been suggested that it directly contributes to neutrophil recruitment in the premetastatic niches in mice. Furthermore, CXCR4 inhibition could impair hematopoietic niches and neutrophil biogenesis by blocking CXCL12-mediated hematopoietic stem cell homing. Interestingly, the phase II clinical trial NCT02826486 recently demonstrated that the combination of BL-8040/BKT140 (a CXCR4 inhibitor) and pembrolizumab in patients with pancreatic cancer reduces TAN infiltration and increases cytotoxic T-cell function.

Besides the classical chemotaxis pathways, members A and C of the IL-17 cytokine family contribute to neutrophil recruitment in various solid tumors mouse models. Hence, IL-17A secretion has been associated with granulocyte/macrophage-colony stimulating factor (GM-CSF)-dependent cancer neutropoiesis in mouse models. Although inhibiting IL-6 or GM-CSF signaling might interfere with IL-17A-related neutropoiesis and TAN infiltration, there is currently no therapeutic approach specifically targeting IL-17 signaling in combination with immune checkpoint blockade. Similarly to IL-17A secretion that links antibacterial immunity to the anti-cancer immune response in mice and human, release of the complement component 5a (C5a) in the TME is another potent attractant for neutrophils and macrophages in mice and on purified human neutrophils in vitro. Although the impact of C5a receptor (C5aR) signaling on TAN plasticity and function remains unclear in cancer, the anti-C5aR antagonist antibody IPH5401 is currently tested in the combination with durvalumab anti-PD-L1 antibody (NCT03665129).

Finally, based on observations made in mice and humans, artificial activation of apolipoprotein E signaling using the LXR agonist RGX-104 is considered as one of the most promising approaches to inhibit neutrophil accumulation in tumors and in the periphery. This approach reduces TAN survival in mice and sensitizes tumors to anti-PD-1 blocking antibodies. Furthermore, a phase I clinical trial (NCT02922764) showed that RGX-104 reduces the fraction of circulating neutrophils in CRC, RCC, sarcoma, uterine cancer, and melanoma. In addition to the direct effect on neutrophil survival, mouse models suggest that RGX-104 also alters neutrophil biogenesis through modulation of the hematopoietic niche function (figure 3, table 1).

TARGETING TUMOR-PROMOTING TAN POLARIZATION TOGETHER WITH ICI TREATMENT

TAN functional diversity is partly the result of polarizing and survival signals that neutrophils receive while infiltrating the TME such as the balance between IFN and TGFβ signals, that drive N1 and N2 functions, respectively. However, whether neutrophil polarization towards the N1 phenotype may have a favorable impact in cancer therapy remains to be demonstrated. We identified 14 clinical trials in which ICIs are combined with type-I IFN (Sylatron, PegIFNαb, HDI IFNa-2b, VSV-IFNβ, IFNβ), 3 trials that are testing ICIs with TGFβ signaling inhibitors (LY3200882, TEW-7197), and 11 clinical trials that use M7824, a bispecific antibody against TGFβ and PD-L1. These combined treatments might have a pleotropic activity on a broad range of immune populations, including TAN polarization towards the N1 phenotype.

The IL6/STAT3 pathway, which plays a pivotal role in cancer-induced inflammation, might also be involved in cancer-induced neutropoiesis and TAN recruitment. Indeed, IL6 production by tumor-associated fibroblasts in HCC induces PD-L1, CXCL8/IL8, TNFα, and CCL2 expression in TANs, and increases their survival and suppressive activity. Interestingly, ongoing trials are evaluating the potential synergistic effect of the fibroblast activation protein inhibitor talabostat, the anti-IL6 antibody tocilizumab, and STAT3 small molecule inhibitors, such as BBI-608/napabucasin, with ICIs, based on multiple mechanisms, including their expected effect on neutrophil biogenesis, homing, and survival (table 1).

The proinflammatory cytokine IL1β might contribute to TAN accumulation, polarization, and survival in the TME. Here again, combining the anti-IL1β antibody canakinumab or the IL1R antagonist anakinra with ICIs in clinical trials might affect both IL1β-mediated cancer cell proliferation and TANs (table 1).

Similarly, tumor-necrosis factor (TNF) signaling inhibition could synergize with immune checkpoint blockade by directly influencing T cells and also by reducing IL-17 secretion and TAN recruitment. Indeed, RIP1 and/or RIP3 inhibition, which abolishes necroptosis and TNF signaling, increases the antitumor activity of macrophages and reduces TAN recruitment in mouse models of pancreatic cancer. GSK3145095, a RIP1/ RIP3 kinase inhibitor that might damp TAN-mediated tumor promotion, is currently evaluated in combination with pembrolizumab (anti-PD-1 antibody) in patients with pancreatic cancer (NCT03681951) (table 1).

However, RIP1 and NF-kB can contribute to the induction of immunogenic cell death, supporting CD8+ T cell cross-priming. In a model of soft tissue sarcoma, inhibition of IAP family antiapoptotic genes (cIAP1 and 2) that influence TNF and RIP1 signaling in cancer cells increases cancer cell death and favor antitumor immunity.
in a RIP1-dependent manner (for a review, see Annibaldi et al.\(^{113}\)). The IAP (cIAP, XIAP) inhibitor birinapan is now evaluated in combination with pembrolizumab. However, due to its effect on neutrophil recruitment and also on immunogenic tumor cell death in the presence of TNF signaling, more studies are required to determine whether RIP1 activation or inhibition should be privileged to enhance ICI efficacy.

Induction of the GM-CSF/CSF2R pathway activates STAT5 signaling, increasing survival and expression of PD-L1 and fatty acid transport protein 2 (FATP2) in neutrophils, ultimately contributing to TAN immunosuppressive functions in mice.\(^{36}\) However, the combination of sagramostim (recombinant GM-CSF) and ipilimumab (anti-CTLA-4 antibody), currently tested in phase I/II trials, increases OS in patients with melanoma from 12.7 to 17.5 months, compared with ipilimumab alone.\(^{114}\) This raised many questions on how GM-CSF influences TANs and other tumor-infiltrating immune cells when administered as an anticancer drug (figure 5).

The expression of ‘don’t eat me’ signals by cancer cells is a potent mechanism to inhibit tumor cell destruction by neutrophils. In cancer cells, CD47 expression and its interaction with its specific receptor, called signal-regulatory protein alpha (SIRP\(\alpha\)), on the neutrophil membrane inhibits their destruction through trogoptosis.\(^{115,116}\) These data provided the rational for the current evaluation of the CD47 inhibitor ALX148 and anti-SIRP\(\alpha\) (TTI-62) or anti-CD47 (TJC4) blocking antibodies in combination with ICIs. In the same way, immunoglobulin-like transcript-4 (ILT4, gene name \(\text{LILRB2}\)) is an inhibitory receptor that impairs neutrophil phagocytosis and respiratory burst,\(^{117}\) and the anti-ILT4 antibody MK-4830 is currently tested in combination with pembrolizumab (table 1).

Inhibition of CD47/SIRP\(\alpha\) and HLA-G/ILT4 signaling should increase TAN antitumor activity, phagocytosis, and cancer cell killing. However, clear mechanistic investigations in patients and mouse models are lacking to fully evaluate the impact of these treatments on neutrophil functions relative to the immune checkpoint blockade response.

Finally, the activation of the PI3K gamma/delta (PI3K\(\gamma/\delta\)) signaling in TANs might also play a major role during the acquisition of tumor promoting functions. In a mouse model of prostate cancer, treatment with the TKI cabozantinib or with the dual PI3K/mTOR inhibitor BEZ235 reduces TAN infiltration and increases the response to the anti-CTLA-4 and anti-PD-1 antibody combination.\(^{50}\) Accordingly, anti-GR1 antibody-mediated neutrophil depletion enhances the response to immunotherapy in this model.\(^{50}\) Similarly, PI3K\(\delta/\gamma\) inhibition or anti-Ly6G antibody-mediated neutrophil depletion sensitizes mice with HNC to ICIs, while their combination (anti-Ly6G plus PI3K\(\delta/\gamma\) inhibition) does not further improve ICI efficacy demonstrating that PI3K\(\delta/\gamma\) acts through neutrophil functional modulation.\(^{48,118,119}\) Indeed, in the CT26 mouse model of CRC, which is characterized by massive TAN infiltration, inhibition of PI3K\(\delta/\gamma\) increases anti-PD-1 immunotherapy efficacy through impairment of TAN immunosuppressive functions.\(^{118}\) Copanlisib, which inhibits PI3K\(\alpha/\beta/\gamma/\delta\), and PI3k\(\gamma/\delta\)-selective inhibitors (idelalisib, tenalisib, itacitinib) are promising drugs that can synergize with ICIs.

All together, these different approaches currently under evaluation in combination with ICIs should help decreasing tumor-promoting TAN polarization and survival and could simultaneously favor the emergence of antitumor neutrophils.

**CONCLUDING REMARKS AND PERSPECTIVES**

Among the 401 clinical trials based on ICI combination with a second drug that might influence or be influenced by TAN and neutrophil biology, 329 (82.45%) want to inhibit tumor vasculature function, and among them, 111 (27.7% of total) are based on VEGF inhibition with bevacizumab. As TANs contribute to bevacizumab resistance, their study becomes highly relevant for such combination settings. Furthermore, 23 (5.7%) of these clinical trials are testing drugs that could revert TAN immunosuppressive functions, and 48 (11.9%) are assessing strategies that might change neutrophil homing and polarization in tumors. This highlights that despite their major impact on the immune response in many solid cancer types, much remains to be investigated to increase the immune checkpoint blockade success through simultaneous TAN targeting. Studying TAN-mediated immune exclusion and immunosuppression implies important efforts to characterize tumor angiogenesis and blood vessel maturation in TAN-infiltrated cancer models.

An important emerging aspect of TAN biology is connecting neutrophil extracellular traps (NETs) with ICI response. Indeed, in patients with CRC\(^{120}\) and in mammary and lung cancer-bearing mice, peripheral blood neutrophils display increased susceptibility to NET formation, which might contribute to cancer-related lung thrombosis.\(^{121}\) Furthermore, NETs have been implicated in metastatic outgrowth in patients with liver cancer after surgical stress through induction of HMG1/TLR9 signaling in cancer cells.\(^{122}\) Similarly, neutrophil elastase and MMP9 released in NETs on lipopolysaccharide-lung metastatic colonization.\(^{11}\) Studying ICI impact on NET formation will most probably help to identify new mechanisms linked to treatment refractoriness, hyperprogression, and might explain some deleterious side effects of immune checkpoint blockade. Indeed, PAD4 inhibitors that abolishes NET formation sensitized the 4T1 breast cancer cell line transplantable model of mammary tumor to anti-PD-1 plus anti-CTLA-4 treatment. The authors suggested that NETs might prevent interactions between tumor cells and cytotoxic effector immune cells.\(^{123}\) Similarly, in a pancreatic cancer model, IL17A signaling inhibition abolished neutrophil recruitment and NET formation in tumors. Inhibition of PAD4 phenocopied IL17A blockade by supressing NET formation, reverting CD8 T cell exclusion form the tumor...
mass and sensitizing to anti-PD-1 plus anti-CTLA-4 treatment\(^{121}\) (for review, see Reference 125).

To conclude, this review of the literature and of ongoing clinical trials raises awareness about the potentially strong impact of neutrophils on the response to ICIs, which should be considered with great attention. Inhibiting TAN tumor-promoting or tumor-protecting functions may substantially increase the success rate of immune checkpoint blockade and the spectrum of patients that can respond to it.

From that, we identified four key questions that should be addressed in the coming years to better understand and exploit neutrophil/TAN biology in the context of ICI development:

► What are the specificity of TAN-related tumor angiogenesis? Understanding how TANS affect tumor vasculature could be remarkable considering immune cell extravasation, drug diffusion and response to antiangiogenic therapy.

► In patients with cancer and in mouse tumor models, TAN or TAM recruitment seems to be mutually exclusive and have a differential impact on the response to ICIs. Are there any specific characteristics of cancer cells that could explain why TAN infiltration is favored in some tumor types rending them resistant to ICIs?

► Can neutrophils and TANS reinforce ICI-induced anti-tumor immune response though intrinsic properties (exacerbation of IFN signaling, cancer cell opsonization by anti-PD-L1 antibody, trogoptosis, tumor antigen presentation and TIL priming) and conversely could they directly be involved in immune-related adverse events and hyperprogression phenomena?

► What are the mechanisms governing TAN and neutrophil diversity in tumors? Can these mechanisms be therapeutically exploited to abolish TAN tumor-promoting functions and favor antitumor TAN activity?

Contributors

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