The Role of Neutrophil Extracellular Traps in Cancer

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Neutrophils are vital components of innate and adaptive immunity. It is widely acknowledged that in various pathological conditions, neutrophils are activated and release condensed DNA strands, triggering the formation of neutrophil extracellular traps (NETs). NETs have been shown to be effective in fighting against microbial infections and modulating the pathogenesis and progression of diseases, including malignant tumors. This review describes the current knowledge on the biological characteristics of NETs. Additionally, the mechanisms of NETs in cancer are discussed, including the involvement of signaling pathways and the crosstalk between other cancer-related mechanisms, including inflammasomes and autophagy. Finally, based on previous and current studies, the roles of NET formation and the potential therapeutic targets and strategies related to NETs in several well-studied types of cancers, including breast, lung, colorectal, pancreatic, blood, neurological, and cutaneous cancers, are separately reviewed and discussed.

Keywords: neutrophil extracellular trap, cancer, neutrophil, immunity, inflammation

INTRODUCTION

Neutrophils are recognized as the most abundant leukocytes in the blood, comprising approximately 50%–70% of all circulating leukocytes in humans and 10%–25% in mice (1–3). They are widely acknowledged as vital members of both innate and adaptive immune responses and defendants against exogenous invaders, including various kinds of bacteria, viruses, and fungi (4–6). Under microbial infection or foreign invasion, neutrophils are rapidly activated and accumulated, which contribute to the restriction and clearance by triggering reactive oxygen species (ROS) production, endocytosis, degranulation, etc. (7). In 2004, Brinkmann et al. observed a special form of neutrophil degranulation, consisting of DNA fibers decorated with granule proteins, which were initially termed neutrophil extracellular traps (NETs) (8). They revealed that the formation of extracellular DNA traps contributed to the constraining and killing of invasive bacteria (8). In the last 17 years since their initial discovery and definition, numerous studies have been devoted to uncovering the characteristics of NETs. Studies have also demonstrated the physiological and pathological functions of self-defense mechanisms in various types of disorders via the involvement of inflammatory and immune responses (9–12). Recently, NETs have been demonstrated to be involved in the pathogenesis and progression of malignant tumors. An increasing number of studies have revealed the pro-tumor effects of NETs. These effects are mediated via mechanisms including the establishment of an inflammatory microenvironment and interaction with other pro-tumor mechanisms such as inflammasomes and autophagy (13–17).
this review, knowledge on the biological characteristics of NETs is presented. Furthermore, the roles of NETs and potential therapeutic targets and strategies related to NETs in several types of cancer, including breast, lung, colorectal, pancreatic, blood, neurological, and cutaneous cancers, is discussed in detail through a review of the latest related studies.

**PART I: BIOLOGICAL CHARACTERISTICS OF NETS**

NETs are extracellular strands of decondensed (unwound) DNA fibers in complex with histones and neutrophil granule proteins, including matrix metalloproteinase (MMP), neutrophil elastase (NE), myeloperoxidase (MPO), cathepsin G, complement factors, and other enzymatically active proteases and peptides (18–20). In the first few years since its initial report in 2004 (8), the term “neutrophil extracellular trap–osis (NETosis)” was widely used in related studies instead of NETs. The extensive use of the term NETosis was based on reports demonstrating that most extrusion of DNA strands resulted in their death, which allowed neutrophils to serve in immune reactions after their death (21–24). However, a strong concern was raised, since an increasing number of studies had reported that the occurrence of NETs does not necessarily lead to neutrophil death (9, 25–27). Therefore, it was strongly recommended by the Nomenclature Committee on Cell Death (NCCD) in 2018 that the term “NETosis” should be replaced with “NETs” or “NET formation” to include the DNA extrusion in the absence of cell death (28). Given this consideration, in this review, “NETs” or “NET formation” is used instead of “NETosis” in the following sections.

Under normal conditions, most DNA strands in neutrophils are highly wrapped around histones into heterochromatin within the nucleus (18, 29, 30). The protein–DNA interactions largely constrain the potential energy of DNA to extend, which leads to transcriptional inactivity (31). Under certain stimuli, such as microbial and sterile agents in vivo or phorbol 12-myristate 13-acetate (PMA), lipopolysaccharide (LPS), and intracellular calcium ion flux in vitro, the condensed DNA strands in neutrophils are uncoiled as fibrous polymers. The decondensation of DNA strands leads to the release of such potential energy, thus facilitating the formation of NETs (illustrated in Figure 1) (32, 33). To date, two proteases have been commonly acknowledged to be vital in the process of NET formation (18, 34–36). The first is peptidyl arginine deiminase 4 (PAD4), which catalyzes the conversion of arginine in histones to citrullines. Such citrullination significantly weakens the original positive charge of histones and weakens the strong histone-DNA binding, which leads to the decondensation of nuclear DNA and/or mitochondrial DNA. Besides PAD4, the other vital protease is NE, which is considered to facilitate the destruction of histone-DNA binding by cleaving histones. Deficiency in either PAD4 or NE in mice has been shown to prevent generation of NETs (37–39). After the decondensation of chromatin and disintegration of nuclei, DNA structures decorated with histones and granule proteins are extruded throughout the cellular membrane with the assistance of gasdermin D. Gasdermin D has been shown to function in the formation of pores, which results in the release of NETs (40, 41). However, several studies have demonstrated that PAD4 is not always necessary for NET formation. For instance, some researchers have argued that the role of PAD4 in NADPH-oxidase (NOX)-dependent NETs remains controversial. They observed the formation of NETs in the absence of detectable histone deamination (32, 42, 43). These studies indicate the complexity of the processes and mechanisms of NET formation.

To date, various signaling pathways have been shown to be involved in NET formation. Changes in the levels of signaling pathway-related proteins have been detected, including protein kinase C (PKC), extracellular regulated protein kinase 1/2 (ERK1/2), c-Jun N-terminal kinase (JNK), Akt, and Scr (44–46). In addition, several mechanisms, including those involved in inflammasomes and autophagy, have been shown to interact with NETs in diseases (47–49). Since the initial discovery of bacterial infection as a stimulating factor for NETs, an increasing number of stimuli have been uncovered in the recent decade to trigger NET formation. In addition to bacteria, several other factors, including interleukin (IL)-8 (a major neutrophil chemoattractant) (50, 51), PMA (a PKC activator) (52–54), LPS (a component of gram-negative bacteria) (55, 56), certain kinds of toxins, intracellular calcium ion flux, ionomycin, and A23187 (18, 42) have been shown to stimulate NET formation. It is notable that the mechanisms for the stimulation of NET formation differ between in vitro and in vivo conditions. For instance, LPS has been demonstrated to be effective in inducing NET formation in vitro through direct stimulation of neutrophils. However, it has been shown that LPS can only induce NET generation by binding activated platelets (thrombocytes) in vivo (57). Due to the complexity of in vivo conditions and the relative limitations of research techniques, a limited number of studies have addressed the specific inducers of NET formation in vivo (33). To date, several factors, including exogenous infections such as bacteria and fungi (10, 58, 59), inflammatory cytokine stimulation (60, 61) and interaction with activated platelets (62) have been recognized as stimuli in vivo. However, few specific agents have been proven to be effective in inducing NET formation in vivo.

Based on the knowledge of the biological and morphological characteristics of NETs, several methods have been developed for the detection and monitoring of NET formation, including immunofluorescence, immunohistochemistry, intravital microscopy, live cell imaging, DNA-intercalating dyeing techniques, and immunoblotting. These methods mainly target NET-related proteins, such as PAD4, NE, MPO, and MMP (33, 63–65). By means of these techniques and methods for NET detection, NETs have been demonstrated to be highly involved in the pathogenesis and progression of several kinds of disorders, including inflammatory bowel disease (66–68), multiple sclerosis (69, 70), atherosclerosis (71, 72), ischemic stroke (73), and some other autoimmune diseases (11, 20, 74). Moreover, an increasing number of studies have uncovered the involvement of NETs in the onset and development of malignant tumors, which will be discussed in detail in the following sections.
PART II: NETS IN CANCER

General Ideas on the Relations Between NETs and Cancer

Notably, a search on the database of PubMed (www.ncbi.nlm.nih.gov/pubmed/) using the keywords “neutrophil extracellular traps” and (“cancer” or “tumor”) yielded 672 results by June 2021, with studies published in the last 4 years (2018–2021) accounting for 65.9% (443 in 672). Demers et al. (75) first reported the role of NETs in cancer. Since then, the prevailing view is that NETs produce a pro-tumor effect through the promotion of cancer cell proliferation, differentiation, metastasis, and other pathological characteristics in various types of malignant tumors (13, 76–81). For instance, it was demonstrated that NE, an important granule protein in NET microvesicles, could degrade the extracellular matrix and induce the phosphatidylinositol-4,5-bisphosphate 3-kinase (PI3K) pathway in cancer cells. The induction of the PI3K signaling pathway promotes cancer cell proliferation and migration (76, 77). Another member of the granule proteins, MMP, was also reported to promote tumor growth and metastasis through proteolysis of the extracellular matrix (13, 78). In addition, it has been revealed that the proteases in NETs can induce the remodeling of laminin, which triggers the integrin signaling pathway in cancer cells and the awakening of dormant tumor growth (13). Moreover, Teijeira et al. demonstrated that CXCR1 and CXCR2 agonist-induced NETs could wrap and coat cancer cells, which shielded them from clearance and the cytotoxic effects of cytolytic cytotoxic T lymphocytes (CTLs) and natural killer (NK) cells (79). These data indicate that NETs may function as a physical barrier preventing the interaction between cancer cells and surrounding inflammatory and immune populations (79, 80). It was further revealed that the levels of intratumoral NETs and supranormal preoperative serum MPO-DNA, regarded as a NET marker, were significantly increased in metastatic cancer tissues (81). They
showed that NETs could facilitate the growth of stressed cancer cells by altering their bioenergetics, while the inhibition of NETs leads to cancer cell death (81).

Cancer-associated platelet activation has been shown to facilitate tumor progression and metastasis through protecting cancer cells from shear forces and assault of immune cells, opening the capillary endothelium to induce the epithelial–mesenchymal transition (EMT) program and secreting pro-tumor growth factors, etc. (82–84). Patients with malignant tumors demonstrate an increase in platelet activation (82). NETs have been shown to contribute to the formation of arterial, venous, and cancer-associated thrombosis (15, 85). These findings suggest that targeting NETs may serve as a potential and promising approach to reduce thrombosis and limit tumor progression and metastasis.

In addition, NETs have also been regarded as an important member of the dynamic tumor immune microenvironment (TIME), which may contribute greatly to impeding metastatic dissemination (51, 86). Several factors have been revealed to contribute to the formation of the TIME. Among them, cancer-associated fibroblasts (CAFs) are regarded as one of the most vital pro-tumor factors. Regarding the influence of NETs on CAFs, it was reported that the formation of NETs originated from CAFs in pancreatic ductal adenocarcinoma, thus producing a pro-tumor microenvironment (87). For the mechanism by which NETs influence the conversion of normal fibroblasts into CAFs in tumors, a nuclear factor (NF)-κB-dependent manner in stomal cells induced by activated neutrophils has been reported (88). Furthermore, the overwhelming production of fibroblast growth factors (FGFs) triggered by NETs in cancer was also shown to be a potential mechanism for the promotion of CAFs (89). However, due to the complexity of the conversion of normal fibroblasts to CAF, more research is required to explore the specific mechanisms mediated by NET regulation.

Although abundant experimental data suggest the pro-tumor effect of NETs on cancer, it is not in the absence of controversy (90, 91). Some researchers have argued that baseline NETs could produce an anti-tumor effect by directly killing cancer cells or limiting tumor growth and metastasis via stimulation of the immune system (90). In addition, NETs have been shown to limit the growth of cancer-related intestinal microbiota populations, thus inhibiting the proliferation and metastasis of colorectal cancer cells (91). Therefore, to ultimately take advantage of NETs for cancer treatment, more knowledge should be gained regarding the precise regulation of NETs.

Currently, there is still little knowledge on the specific mechanisms of NET formation in cancer. Although some findings have been obtained, there is no consensus regarding this issue. A previous study revealed that NETs were produced by neutrophils in a toll-like receptor 4 (TLR4) and high-mobility group box 1 (HMGGB1)-dependent manner in lung cancer cells (92). In addition, it has also been reported that the chronically inflammatory microenvironment contributes to the formation of NETs via an MPO–NETs–antineutrophil antibody (ANCA) axis in cancer (93, 94). Researchers have shown that NETs might act as a central factor in the neutrophil–NET–cancer cycle, thus aggravating the pathogenesis and progression of cancer. Other researchers have revealed that NETs might be associated with other cellular processes, including inflammasomes and autophagy, which will be discussed in the following sections.

**INFLAMMASOMES, NETS, AND CANCER**

Inflammasomes are multi-protein complexes that are widely regarded as important factors of inflammation and innate immunity responsible for the activation of inflammatory and immune responses through the recognition of pathogen-associated molecular patterns (PAMPS) or danger-associated molecular patterns (DAMPs) (95). To date, several types of inflammasomes have been reported, including the NOD-like receptor family, pyrin domain-containing 1 (NLRP1), NLRP2, NLRP3, NLR family caspase recruitment domain-containing protein 4 (NLRC4), and double-stranded DNA sensors absent in melanoma 2 (AIM2) (96–98). As previously reviewed by us, the activation of inflammasomes involves the cleavage of procaspase-1 into caspase-1, which subsequently catalyzes the production and secretion of inflammatory cytokines, including IL-1β and IL-18 (99, 100). The involvement of NF-κB signaling in the initiation of inflammasome activation leads to increased production of inflammasome components (99, 100). Recently, numerous studies conducted by our laboratory and others have demonstrated the roles of inflammasomes in various diseases, including inflammatory bowel disease, multiple sclerosis, atherosclerosis, stroke, and malignant tumors (96, 101–104). In cancer, it has been revealed that the over-induction of inflammasomes largely influences cancer cell death, proliferation, and even tumor growth microenvironment, such as intestinal microbiota populations and fibroblasts, which indicates the wealth of putative inflammasome-based targeted therapies for cancer (104–110).

As important components in immune reactions, NETs have been shown to be closely linked to inflammasomes in some autoimmune diseases and cardiovascular disorders (19, 111, 112). For instance, it has been previously described that the formation of NETs by neutrophils could trigger the synthesis of inflammasome-related IL-1β and IL-18 in macrophages through cathelicidin LL-37-mediated potassium efflux from the cells. The production of IL-1β and IL-18 subsequently promotes NET formation during the occurrence of cardiovascular diseases (19, 113). Similar crosstalk between inflammasomes and NETs has been reported in cancer. It has been demonstrated that NET-associated serine proteases such as NE could act as alternative enzymes for processing inflammasome-related IL-1β and IL-18, which subsequently leads to the modulation of PGRN inactivation and MMP-9 activation in cancer (114). In addition, Albrengues et al. (13) reported in lung cancer that NETs and NETs-mediated extracellular matrix remodeling acted as critical mediators of awakening of dormant cancer cells by LPS-mediated formation of inflammasomes in mice. This process is mediated by the binding between NET-DNA and extracellular matrix protein laminin and bringing NE and MMP-
9 to their substrates (13). To date, little evidence has been available regarding the crosstalk between inflammasomes and NETs in cancer (13, 114, 115). However, future studies should further elucidate such interactions as they may unravel potential novel therapeutic strategies for the treatment of cancer.

**Autophagy, NETs, and Cancer**

Autophagy, commonly recognized as a vital metabolic mechanism relying on lysosomes, functions in degrading and recycling long-lived, misfolded proteins and damaged organelles to maintain cellular homeostasis (116–118). As previously described, under organic stress, such as nutrient deprivation or inflammatory loading, cytoplasmic materials are targeted and sequestered into autophagosomes, which are regarded as the functional units of autophagy (119–121). The autophagosomes fuse with lysosomes to form autolysosomes (119–121). Notably, Yoshimori Ohsumi was awarded the 2016 Nobel Prize in Medicine or Physiology for exploring the cellular autophagy processes (122). Since its discovery by Christian de Duve in the 1960s (123), numerous studies from our laboratory and others have uncovered the involvement of autophagy in various diseases, including cardiovascular disorders, autoimmune diseases, metabolic abnormalities, malignant tumors, neurodegenerative diseases, and gastrointestinal diseases (124–129).

The influence of autophagy on the pathogenesis and progression of cancer tends to be regarded as a double-edged sword (130–134). In contrast, autophagy has been considered to maintain stemness, induce recurrence, and develop resistance of cancer cells to anticancer agents. The administration of rapamycin, an autophagy inducer, has been shown to be effective in alleviating cancer (131–133). Additionally, autophagy has also been reported to inhibit tumor initiation through the induction of autophagic cell death, and chloroquine, an autophagy inhibitor, has been used in anti-tumor therapy (131, 134). These studies indicate the complexity of autophagy mechanisms in cancer.

The dichotomous effects and complex mechanisms of interactions between autophagy and NETs have also been demonstrated in cancer. It has been reported that the induction of autophagy (formation of autolysosomes) in leukemia cells leads to the release of NETs, which results in the deterioration of acute promyelocytic leukemia (135). In addition, Boone et al. (136) revealed that NETs were upregulated in pancreatic cancer via receptor for advanced glycation end products (RAGE)-dependent/autophagy-mediated pathways. Regarding the effect of autophagy and NET crosstalk in cancer, it was previously revealed that the administration of chloroquine significantly reduced the hypercoagulability in pancreatic cancer by inhibiting NETs, suggesting a positive effect of suppressing autophagy–NET interaction in the alleviation of cancer (137). However, autophagy has been reported to be involved in the effects of interferon (IFN)-γ on cell growth inhibition and cytotoxicity in lung epithelial malignancies via the induction of PAD4-mediated NETs (138). Consequently, to ultimately take advantage of interactions between autophagy and NETs in cancer treatment, further studies are required to explore this issue.

**PART III: NETS IN DIFFERENT TYPES OF CANCER**

During the 16 years since the initial discovery of NETs in 2004, an increasing number of researchers have focused on the study of NETs in cancer. Fortunately, to date, the effects of NETs in several popular types of cancer have been widely revealed to illustrate the whole picture. In this section, the roles of NETs and potential therapeutic targets and strategies related to NETs in several types of cancers, including breast, lung, colorectal, pancreatic, blood, neurological, and cutaneous cancers are described and discussed in detail based on the latest studies available in the current database (listed in Table 1).

**Breast Cancer**

Breast cancer is regarded as one of the three most commonly diagnosed cancers worldwide, especially in women (160). Breast cancer is one of the most studied type of cancer-related to NETs. It was first reported by Demers et al. (75) that, in a murine late-stage breast cancer model, the formation of NETs corresponded with cancer-associated thrombosis in the lung. The formation of thrombosis contributed to a poor prognosis and cancer-caused death. In addition, the expression levels of PAD4 genes were shown to be high in murine breast cancer 4T1 cells and PAD4-mediated NETs, which contributed to the release of cancer extracellular chromatin networks (CECN) both in vitro and in vivo (139). PAD4-mediated NETs were demonstrated to promote breast tumor growth and cancer metastasis into the lung, since the deletion of PAD4 genes in mouse models largely attenuated breast cancer cell proliferation and migration (139). In addition, Martins-Cardoso et al. (140) revealed that NETs promoted a pro-metastatic phenotype in human breast cancer cells by inducing the EMT program.

Regarding the exploration of therapeutic targets related to NETs in the treatment of breast cancer, a brilliant study by Yang et al. (141) uncovered a potential specific mechanism for the influence of NETs in breast cancer metastasis. In this study, the researchers revealed that the DNA components of NETs (NET–DNA) could act as a chemotactic factor to attract breast cancer cells rather than merely “trap” them, thus leading to the occurrence of liver metastases in patients with early-stage breast cancer (141). They also suggested that the transmembrane protein CCDC25 might act as a potential NET–DNA receptor in breast cancer cells by sensing extracellular DNA. The activation of CCDC25 consequently enhanced cell motility through the subsequent activation of the ILK-β-parvin pathway. These data indicate an appealing therapeutic strategy that takes advantage of targeting CCDC25 for cancer metastasis prevention (141). It has been commonly revealed that patients with breast cancer are at a relatively higher risk of developing thrombosis. Gomes et al. (142) demonstrated that blockade of inflammasome-related IL-1β production and
secretion attenuated cancer-associated thrombosis in a NETs-dependent breast cancer model, indicating the vital role of the crosstalk between NETs and inflammasomes in the treatment of breast cancer.

**Lung Cancer**

Lung cancer is the most diagnosed and frequent cause of cancer-related deaths worldwide, with approximately 1.8 million newly diagnosed patients and 1.6 million deaths each year (161). Thus far, NETs have been demonstrated to modulate the biological characteristics of lung cancer cells, thus influencing the pathogenesis and progression of lung carcinoma. For instance, a brilliant study published in Science revealed that sustained lung inflammation mediated by smoke exposure or nasal instillation of LPS could awaken dormant cancer cells and facilitate metastasis through the induction of NET formation (13). Mechanistic analysis revealed that this process was potentially via NE/MMP-9-induced cleavage of laminin, which subsequently led to the activation of integrin α3β1 signaling in dormant cancer cells (13). However, it is worth mentioning the difference between lung cancer patients and tumor-bearing mice models pointed out by Arpinati et al. (162). They argued in a recent study that no predisposition of neutrophils to release NETs in patients with lung cancer was found to be similar to that in mice compared with healthy controls. These results indicate that attention should be paid to the translation of experimental results obtained from animal studies into clinical applications (162).

Regarding therapeutic targets related to NETs in lung cancer, it was previously revealed that the DAMP protein HMGB1 released by lung cancer cells contributes to the induction of NETs that was dependent on the activation of TLR4 (92). In addition, extracellular RNAs (exRNAs) produced by lung cancer cells have also been demonstrated to induce NET formation, which promotes cancer cell proliferation and migration (143). In addition to exploring therapeutic targets, some researchers have focused on the development of therapeutic strategies for the treatment of lung cancer-related to NETs, using certain natural compounds. Recently, Li et al. (144) revealed that emodin, the main bioactive component of Rheum palmatum, could significantly prevent hypercoagulation and lung carcinogenesis by suppressing NET formation in lung cancer animal models. However, most related research is still in the stage of animal models. Therefore, more clinical studies are required for the development of cancer therapies based on NETs.

**Colorectal Cancer**

According to the analyzed data in 2019, colorectal cancer is considered to be the fourth most deadly cancer worldwide, with 900,000 deaths annually (163). Aging, western lifestyle, and bowel inflammatory conditions (e.g., inflammatory bowel diseases) serve as risk factors for the oncogenesis of colorectal
cancer (163, 164). Richardson et al. (165) demonstrated that patients with colorectal cancer have significantly increased NET formation compared with healthy volunteers, and high levels of NETs were shown to be associated with adverse patient outcomes. Furthermore, NETs have been reported to be involved in venous thrombogenesis in patients with colorectal cancer via the induction of procoagulant activity (PCA), which led to a close interaction with platelets and endothelial cells (145). These data indicate that targeting NETs may be a potential and promising way to tackle thrombosis in colorectal cancer.

Exploring potential therapeutic targets related to NETs in colorectal cancer, Rayes et al. (146) revealed that NETs-associated carcinoembryonic antigen cell adhesion molecule 1 (CEACAM1) acts as an essential element for the interaction between NETs and colorectal cancer cells, and deficiency in CEACAM1 significantly decreases cancer cell adhesion, migration, and metastasis. Regarding the induction of NET formation, KRAS mutation has been shown to contribute to neutrophil recruitment and NET formation through exosomes in colorectal cancer, which uncovered a novel mechanism of regulation of NET formation in colorectal cancer (147). In addition, IL-8, an inflammatory cytokine, has been reported to be part of a positive loop connecting NETs and colorectal cancer liver metastasis (148). However, further studies are required to develop effective therapeutic strategies for colorectal cancer, taking advantage of these targets.

**Pancreatic Cancer**

Pancreatic cancer is a highly lethal cancer in the absence of a standard screening program since most patients remain asymptomatic until they reach an advanced stage (166). A clinical study demonstrated that tumor-infiltrating neutrophils (TINs) and their generated NETs could be treated as prognostic factors for pancreatic cancer independent of the TNM staging system (167). NETs have also been shown to promote liver micrometastasis in pancreatic cancer by activating CAFs (87). Notably, it has been commonly recognized that pancreatic cancer induces a hypercoagulable state, which could result in clinically apparent thrombosis (168). An increasing number of studies have demonstrated the influence of NETs on hypercoagulability and thrombogenesis (137, 149, 168). For instance, Yu et al. (149) showed an association between the upregulation of NETs and PCA in patients with pancreatic cancer through the modulation of cell–cell junctions. Furthermore, Hisada et al. (169) revealed that neutrophils and NETs contribute to venous thrombosis in mice bearing human pancreatic tumors and patients with pancreatic cancer, indicating a thrombogenic effect of NETs in pancreatic cancer.

Regarding therapeutic targets related to NETs in pancreatic cancer, Jin et al. (150) revealed that NETs could promote migration and invasion of pancreatic cancer cells via EMT and the IL-1β/epidermal growth factor receptor (EGFR)/ERK pathway. These results indicate that blockade of the IL-1β/EGFR/ERK signaling pathway might serve as a potential strategy for the alleviation of pancreatic cancer. In addition, it has been reported that targeting the interaction between NETs and autophagy reduces hypercoagulability in pancreatic cancer (137). They showed that the administration of chloroquine for the blockade of autophagy significantly lowered the rates of venous thromboembolism in patients with pancreatic cancer by suppressing autophagy-related NET formation. However, further studies are needed for successful clinical application.

**Blood Cancers**

Blood cancers are a group of “invisible” cancers, as tumors are seldomly observed in blood cancers (170). In a broad sense, blood cancers include leukemia, multiple myeloma, and malignant lymphoma. Regarding the effects of NETs on blood cancers, a statistically significant increase in NET levels was found in patients with multiple myeloma (MM) compared to healthy volunteers both in the serum and plasma (171). Furthermore, MM cells have been shown to stimulate citrullination of histone H3 and subsequently lead to the formation of NETs through the activation of PAD4 (151). In addition, NETs have been reported to contribute to thrombogenesis, a major cause of mortality in patients with myeloproliferative neoplasms (MPNs) via platelet activation (152).

Regarding the therapeutic targets related to NETs in blood cancers, it was reported by Wolach et al. (153) that NET formation was increased in mice MPN models via the Janus kinase (JAK)-activator of transcription (STAT) signaling. The administration of ruxolitinib, a JAK2 inhibitor, effectively suppressed NET formation and reduced thrombosis. In addition, NET formation has been previously shown to contribute to the pathogenesis of chronic lymphocytic leukemia (CLL), and ibritinib, a widely used chemotherapeutic drug, alleviated CLL by slightly impairing NET production in patients with CLL (154). Based on these findings, the modulation of NET formation may be a potential and promising therapeutic strategy for the treatment of blood cancers.

**Neurological Cancers**

Neurological cancers are defined as a group of primary or metastatic cancers of the central nervous system (CNS) (172). According to the World Health Organization (WTO) in 2016 (173), neurological cancers comprise diffuse astrocytic and oligodendroglial tumors, other astrocytic tumors, ependymal tumors, other gliomas, lymphomas of the CNS, and metastatic cancers. Regarding the role of NETs in neurological cancers, it was previously reported that NETs could significantly promote cancer-associated arterial microthrombosis in cancer patients presenting with ischemic stroke and large elevations of highly sensitive troponin T (hs-TnT) (155). In addition, in patients with high-grade gliomas, a high level of thrombogenic NETs produced by neutrophils was detected, which resulted in an increase in venous thromboembolism (VTE) (174).

Regarding therapeutic targets related to NETs in neurological cancers, Zha et al. (156) revealed that NETs generated by TINs mediated the crosstalk between glioma and tumor microenvironment in patients with malignant glioma. These effects were demonstrated to be regulated by the HMGB1/RAGE/IL-8 axis. However, little is known about the specific mechanisms of NET formation in neurological cancers. Consequently, further studies are required to ultimately develop therapies targeting NETs.
Cutaneous Cancers

Cutaneous cancers, also known as “skin cancers”, are a group of cancers that occur in the skin tissue. Cutaneous cancers are divided into melanoma and non-melanoma, with the latter including basal cell carcinoma and squamous cell carcinoma (175). The mechanisms underlying the pathogenesis and progression of cutaneous cancers are complex and remain unclear. To date, modern research has demonstrated that NETs are closely related to cutaneous cancers. NETs have been reported to promote inflammation-mediated skin tumor cell growth in mouse models (176). NETs have also been shown to contribute to spontaneous and immunotherapy-induced adverse reactions of melanoma murine models (157). In addition, similar to other tissues, NET-mediated thrombosis was also found in the skin tissue, indicating the positive effects of NETs on thrombosis in cutaneous disorders (177).

IL-8 has been reported to be a potential therapeutic target related to NETs, since the association of IL-8 and NETs was revealed in patients with metastatic melanoma (158). In addition, type I IFNs, widely used anti-inflammatory agents, have been shown to induce anti-tumor polarization of tumor-associated neutrophils in murine models and melanoma patients. The transfer of neutrophils into an anti-tumor disturbs the formation of NETs and thus produces a tumor-suppressive effect (159). However, additional studies are needed to develop therapies for cutaneous cancers that target NETs.

CONCLUSION

Overall, recent studies have demonstrated the important roles of NETs in cancer through modulation of the biological characteristics of cancer cells including proliferation, differentiation, and metastasis and induction of cancer-related thrombogenesis (illustrated in Figure 2). So far, we have gained significant knowledge on the biological features and regulation of NET formation, as well as the mechanisms of the effects of NETs on different types of cancers. However, because of the limitations of current studies, the specific mechanisms of NETs and the crosstalk between NETs and other cancer-related processes, including inflammasomes and autophagy, remain unclear. The specific mechanisms underlying NET formation also remain unclear. Therefore, additional studies are required for the successful application of knowledge regarding NET formation and function in clinical practice through the development of novel and promising therapeutic strategies against cancer.

**FIGURE 2** | Schematic illustration of mechanism of NETs in cancer. Under cancer-related stimulation, the level and function of PAD4 are enhanced, leading to the citrullination of histones and subsequent decondensation of chromatin. NET microvesicles with DNA-structural fibers decorated by histones, MMP, NE, MPO, cathepsin G, and other granule proteins are extruded from neutrophils. NET formation modulates the biological characteristics of cancer cells in proliferation, differentiation, migration, and metastasis through the involvement and crosstalk with some specific pathways and mechanisms including HMGB1/RAGE/IL-8 axis, IL-1β/EGFR/ERK signaling pathway, JAK/STAT signaling pathway, integrin α3β1 signaling pathway, CEACAM1 signaling pathway, KRAS mutation, Extracellular RNA induction and crosstalk with inflammasomes and autophagy, and cancer-associated fibroblast activation. NETs, neutrophil extracellular traps; PAD4, peptidyl arginine deiminase 4; MMP, matrix metalloproteinase; NE, neutrophil elastase; MPO, myeloperoxidase; HMGB1, high-mobility group box 1; RAGE, advanced glycation end products; EGFR, epidermal growth factor receptor; ERK, extracellular regulated protein kinase; JAK, Janus kinase; STAT, activator of transcription; CEACAM1, carcinoembryonic Ag cell adhesion molecule 1.
AUTHOR CONTRIBUTIONS

B-ZS and YY retrieved concerned literatures and wrote the manuscript. J-PL designed the table and figures. E-QL and N-LC revised the manuscript. All authors contributed to the article and approved the submitted version.

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