

Expanding Neutrophil Horizons: New Concepts in Inflammation

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Keywords

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Abstract

Research into neutrophil biology in the last 10 years has uncovered a number of unexpected aspects of this still mysterious innate immune cell. Advances in technology have allowed visualisation of neutrophil trafficking to sites of inflammation, and, remarkably, neutrophils have been observed to depart from the scene in what has been termed reverse migration. There has also been increasing appreciation of the heterogeneity of neutrophils with ongoing categorisation of neutrophil subsets, including myeloid-derived suppressor cells and low-density granulocytes. Newly recognised neutrophil functions include the ability to release novel immune mediators such as extracellular DNA and microvesicles. Finally, studies of neutrophil cell death, both apoptotic and non-apoptotic, have revealed remarkable differences compared to other cell types. This review will highlight important discoveries in these facets of neutrophil biology and how the new findings will inform treatment of diseases where neutrophils are implicated.

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Introduction

The neutrophil has long been recognised as a pivotal player in the innate immune response. Neutrophils employ well-characterised strategies of phagocytosis, reactive oxygen species (ROS) generation and release of pro-inflammatory mediators to provide the critical first-line defence against pathogens. At the end of their lives, neutrophils undergo apoptosis, an orderly programmed dismantling of the cell, are phagocytosed and promote the resolution of inflammation [1]. In the last 10 years, there has been a growing appreciation that neutrophils play roles beyond this relatively simplistic schema (Fig. 1). Investigation of neutrophil trafficking using intravital microscopy has allowed researchers to monitor diapedesis and migration in vivo, leading to novel paradigms in homing and trafficking. Furthermore, within the population of neutrophils, there are subsets that differ in phenotype and functionality, and there has been growing appreciation that neutrophils can modulate the function of other immune cells [2]. This integrated role of neutrophils acting within the immune system is highlighted by their importance in autoimmune diseases [3]. This review will focus on progress in the understanding of several facets of neutrophil biology that reflect these new roles and functions [4].

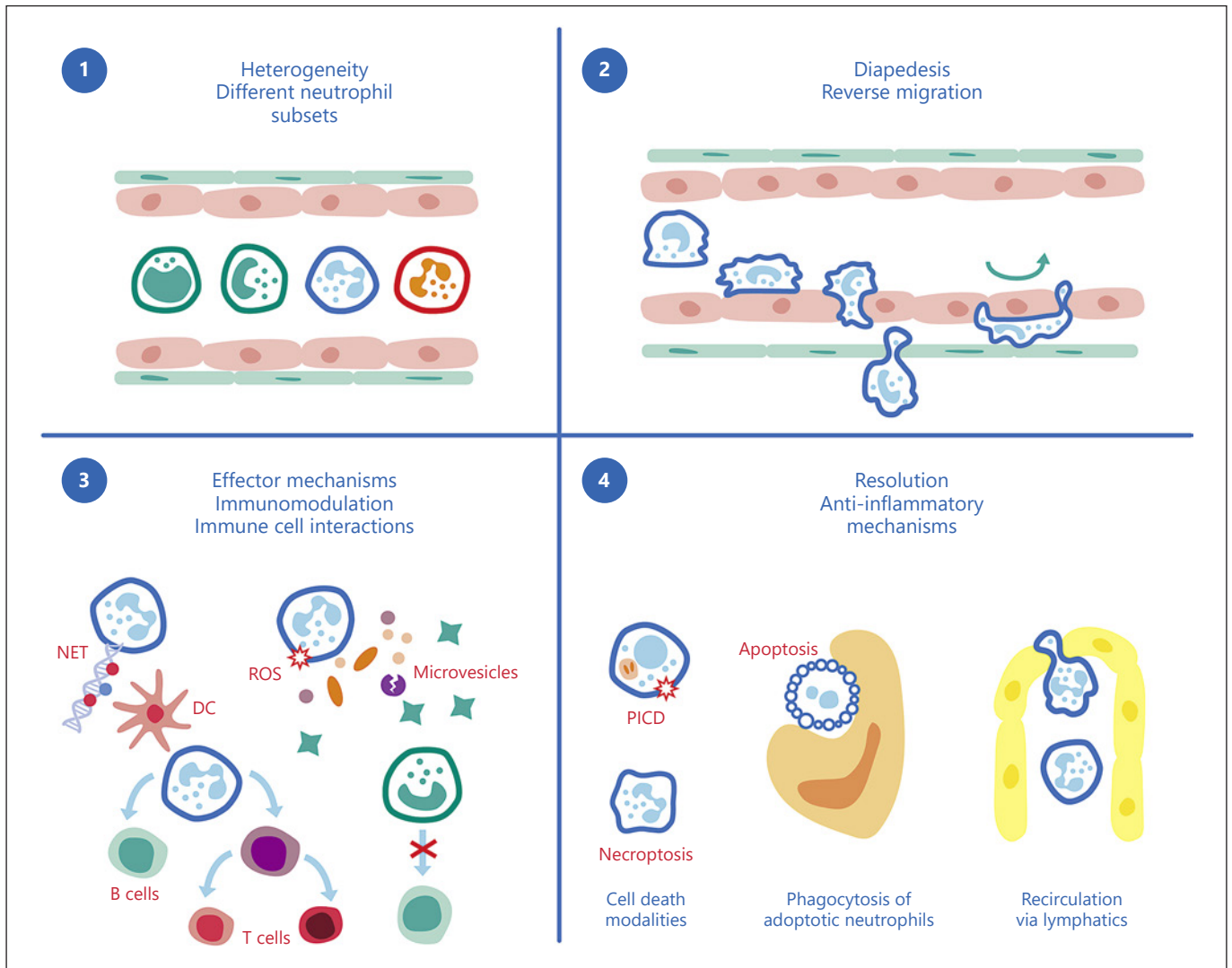


Fig. 1. Schematic overview of neutrophils. (1) Neutrophil heterogeneity: in inflammatory or pathological conditions, neutrophil subsets have been characterised as including immature neutrophils, hypersegmented neutrophils, low-density granulocytes and granulocyte myeloid-derived suppressor cells. (2) Neutrophil diapedesis: transendothelial migration (rolling, adhesion, crawling and migration to inflamed sites) and reverse migration (returning to the circulation after having migrated into tissues). (3) Effector mechanisms of neutrophils: oxidative burst with production of superoxides, hydrogen peroxide (reactive oxygen species [ROS]) and hypochlorous acid; release of anti-pathogen proteases, defensins, cytokines and chemokines; release of specialised pro-resolving molecules, neutrophil extracellular traps (NETs) and neutrophil

microvesicles; and phagocytic clearance of bacterial and cell debris. Neutrophils interact with dendritic cells (DCs), B lymphocytes, natural killer cells, T lymphocytes and endothelial cells and thus influence both the innate and adaptive immune responses. (4) Resolution of inflammation: the modality of cell death (e.g., phagocytosis-induced cell death [PICD], necroptosis and apoptosis) may influence the response of surrounding tissue cells and either perpetuate inflammation or promote resolution. Macrophage efferocytosis of apoptotic neutrophils favours the resolution of inflammation. Recirculation of neutrophils shuttling antigens in lymphatics shapes the adaptive immune response. Interference with or defects in these pro-resolving mechanisms may promote inflammation and autoimmunity.

Neutrophil Trafficking and Homing: No More a One-Way Trip

Neutrophil migration to sites of inflammation has until recently been characterised as a linear progression of

intravascular rolling, tethering and adhesion before irreversible diapedesis into tissues [5]. This schema has recently been challenged, particularly in respect to tissue migration. In a number of model systems taking advantage of significant progress in in vivo imaging, reverse

migration of neutrophils from tissues back into the circulation has been observed and characterised [6, 7]. The ability to migrate back across the endothelial cell layer is dependent on neutrophil-derived neutrophil elastase-degrading JAM-C, a key endothelial adhesion molecule [8, 9]. The fate of neutrophils after migrating to the site of sterile inflammation was further characterised using similar *in vivo* two-photon strategies [10]. Once again, moving away from the dogma that neutrophils reach the end of their lifespan at the site of inflammation and either die or are phagocytosed, neutrophils were observed to re-enter the vasculature, migrate to the lungs initially and then home to the bone marrow, where they underwent apoptosis. With this new technology, it remains to be studied whether there is a dominant pattern of neutrophil behaviour in all forms of inflammation or whether the context, in particular sterile and non-sterile, drives different migration patterns. Species-specific effects also need to be considered, although at this time, in the absence of the ability to deeply study neutrophil dynamics in humans, paradigms will continue to be guided by animal models.

At this stage, the importance of this mechanism in the process of inflammation remains to be firmly established. Reverse migration has the potential to be deleterious, allowing localised inflammation to disseminate [9]. Alternatively, encouraging neutrophil egress from sites of inflammation could serve as a pro-resolving mechanism, as demonstrated in a zebra fish model [6]. Answering these key questions in reference to human disease will determine how best to harness reverse migration for therapeutic purposes.

Neutrophil Heterogeneity – Not All Neutrophils Are Equal

Recent research suggests that the neutrophil population is not made up of homogeneous innate effector cells. On the contrary, analogous to subsets identified amongst lymphoid cells and myeloid cells [11], there may be subsets of neutrophils with potentially important differences in function.

Neutrophil phenotype analysis has been investigated in various conditions including inflammation, autoimmune disease and malignancy. Agreement remains to be reached on the exact nature of these neutrophil subtypes, particularly in regard to cell surface phenotype and whether these subtypes are conserved across species. However, the data are clear that neutrophils do not com-

prise one uniform population and that this heterogeneity is reflected not only in differences in cell surface marker phenotype but more importantly in function.

One neutrophil subtype of interest is the granulocyte myeloid-derived suppressor cell (G-MDSC). MDSCs comprise a heterogeneous population of cells possessing suppressive capacity and may be derived from both granulocytes and monocytes [12]. Suppression of T cells has been most extensively studied, and potential mechanisms include the release of ROS, nitric oxide and arginase and contact-mediated suppression. The suppressive function of G-MDSCs has been most extensively studied in the context of cancer. In different studies, tumour-infiltrating neutrophils have been found to both promote and inhibit malignancy through mechanisms targeting other immune cells, stromal cells and the tumour itself [13]. In particular tumours, a high circulating neutrophil-to-lymphocyte ratio and the number of neutrophils within the tumour have been identified as poor prognostic indicators suggesting that granulocytes may play a role in cancer progression [14]. The relative role in cancer pathogenesis of subpopulations of granulocytes, and MDSCs in particular, is the subject of ongoing studies.

Low-Density Granulocytes

Over many years, it has been noted that under certain inflammatory conditions, neutrophils can be found in the mononuclear cell fraction of Ficoll gradient separation [15]. These cells, so-called low-density granulocytes (LDGs) or low-density neutrophils, have been better characterised over the last decade, but further research is needed to fully understand their role in various diseases. One study investigated the nature of these cells in humans treated with granulocyte colony-stimulating factor (G-CSF), conducted to mobilise haemopoietic stem cells [16]. Under these conditions, a large percentage of granulocytes are found in the polymorphonuclear cell fraction of Ficoll-separated blood. Intriguingly, both the low-density and normal-density neutrophils demonstrated heterogeneity, primarily in relation to the presence of immature neutrophils characterised in this study by the absence of the surface marker CD10. The mature neutrophils released by G-CSF treatment demonstrated T-cell suppressive activity, whereas the immature cells promoted T-cell survival and proliferation. The pathologic relevance of these cells remains to be established, but G-CSF appears responsible for the mobilisation of neutrophils in certain tumours to promote metastases [17].

In sepsis, LDGs are expanded and have been found to exhibit MDSC characteristics [18]. High levels of G-

MDSCs in sepsis predicted subsequent secondary infections, providing one hypothesis for the relative immune suppression that is seen in sepsis. The other scenario in which LDGs are present in significant numbers is in inflammatory diseases, best characterised in systemic lupus erythematosus (SLE) [19]. The LDGs in SLE patients express pro-inflammatory genes driven by type 1 interferons and undergo more rapid cell death than healthy control neutrophils [20, 21]. This cell death, possibly triggered by immune complexes, results in the extracellular release of both nuclear DNA and oxidised mitochondrial DNA [22] in conjunction with neutrophil proteins including LL-37 and HMGB1. These structures in turn are bound by autoantibodies, commonly found in the blood of SLE patients, and this leads to their uptake by plasmacytoid dendritic cells and stimulation of interferon production. This vicious cycle of interferon-driven neutrophil activation is proposed to be a potential target of therapeutic intervention in SLE [23].

LDGs have also been found in the circulation of patients with rheumatoid arthritis (RA). In the case of RA, LDGs are an immature subset with prolonged survival but relatively low rates of NETosis and lower response to cytokine stimulation. Of note, exogenous G-CSF as described above induces LDGs, and G-CSF may play an important role in RA, with antagonism to G-CSF being considered as a therapeutic target [24].

These studies suggest that the phenotype of LDGs may vary depending on the disease context. Describing LDGs in other contexts will be important in order to understand their role and whether they may serve as therapeutic targets in inflammation and other diseases. Future study will be facilitated by finding other markers of LDGs, which can currently only be assessed using Ficoll gradient separation.

Age as a Source of Neutrophil Heterogeneity Young Neutrophils

Release of immature neutrophil forms into the circulation during infection leading to the appearance of band forms during haematological testing is well known, termed left shift by Joseph Arnet in 1904 [25]. Some recent studies have characterised these cells. A study of bone marrow neutrophils in mice using advanced cell cytometric techniques identified a dedicated pre-neutrophil population capable of differentiating into mature and immature neutrophils (based on CXCR2 status) [26]. This pre-neutrophil responded to systemic stressors of inflammation and malignancy by proliferating and providing a sustained neutrophil response in blood and tissues. This population were also present in the spleen and

markedly expanded under the same stressors. Notably no immature neutrophils were present in the blood under steady-state conditions, but they were present in malignancy. CD101 was identified as a positive marker of immature murine neutrophils. Immature neutrophils were also mobilised with G-CSF.

Another study employed a human *in vivo* model of sepsis coupled with deuterium labelling of neutrophils to establish the timing of release of neutrophils from the bone marrow following cell division, at which time the deuterium is incorporated into DNA. With lipopolysaccharide treatment, an immature banded neutrophil population was identified and two separate mature populations were released into the circulation [27]. The immature cells released under systemic lipopolysaccharide treatment were demonstrated to be 2 days younger than the mature population. The mature neutrophil subsets were separated based on CD62L labelling, and the differences were validated by demonstration of significantly different proteomic profiles, pointing towards important functional differences that remain to be demonstrated.

Old Neutrophils

Ageing neutrophils are also present in the circulation, and increasing evidence suggests they behave differently from their younger counterparts. A paper by Casanova-Acebes et al. [28] demonstrated this ageing of neutrophils in mice *in vivo*. Ageing neutrophils expressed high levels of CXCR4, previously identified as being upregulated during *in vitro* ageing of neutrophils. The neutrophils were also CD62^{lo} and are eliminated from the circulation in a circadian fashion by bone marrow macrophages. This clearance in turn provides the impetus to the further release of neutrophils.

Physiologic neutrophil ageing appears to be governed by the microbiome [29]. In sickle cell disease, a chronic inflammatory condition, numbers of circulating aged neutrophils were significantly elevated in both a mouse model and in affected patients. These aged neutrophils unexpectedly possess a pro-inflammatory phenotype, in contrast to neutrophils aged *in vitro*, which exhibit a downregulated functionality [30]. Aged neutrophils have enhanced integrin activation and release neutrophil extracellular traps under inflammatory conditions. Of note, antibiotic treatment abrogated the increase in aged neutrophils under these conditions and consequently reduced the impact of inflammation reflected by less vascular occlusion and fibrosis.

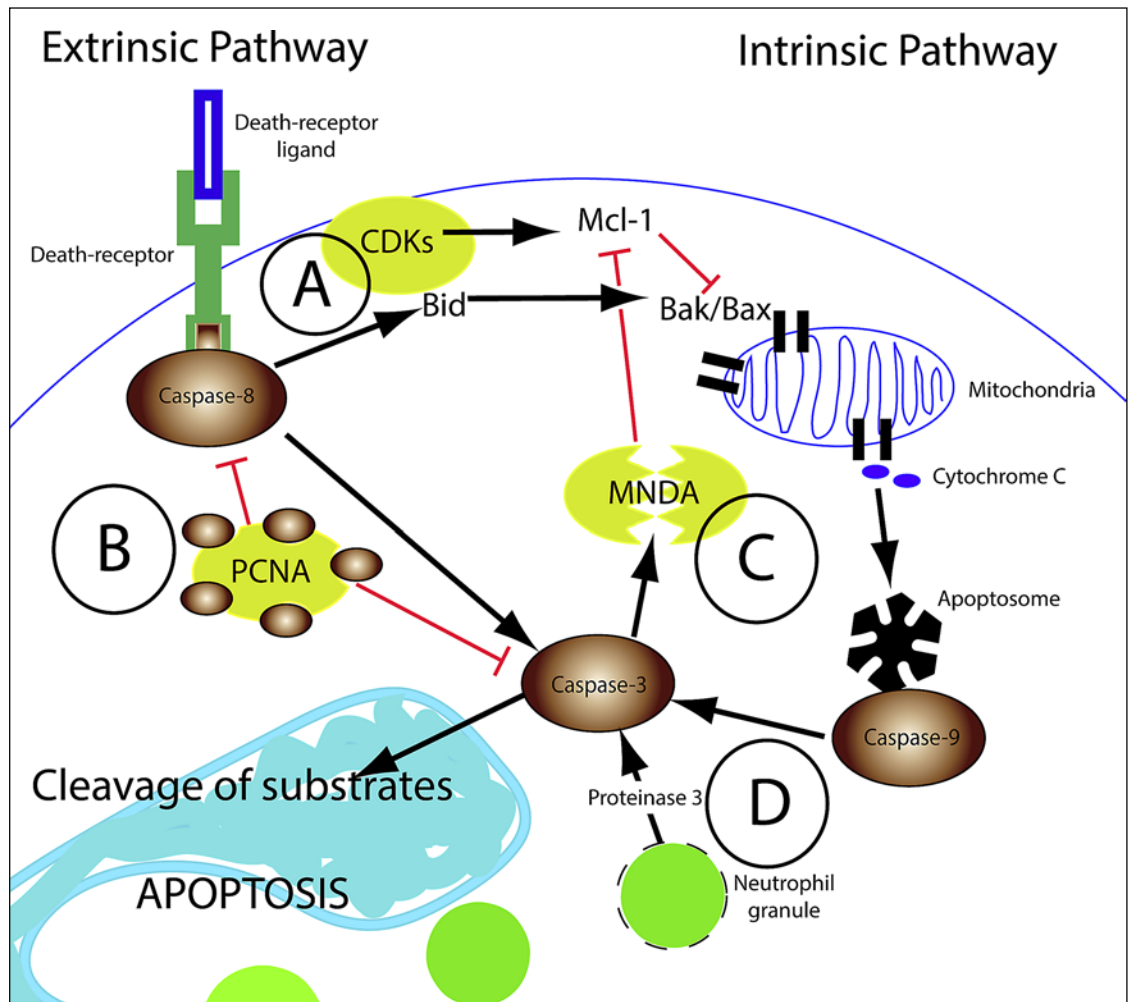


Fig. 2. Neutrophil apoptotic pathways. A simplified schematic of the intrinsic and extrinsic pathways of apoptosis with specific emphasis on the main factors involved in neutrophil apoptosis. Novel mediators of neutrophil apoptosis are: (A) cyclin-dependent kinases (CDKs) that inhibit Mcl-1 degradation and reduce apoptosis; (B) proliferating cell nuclear antigen (PCNA), which sequesters

pro-caspases to prevent their activation; (C) myeloid cell nuclear differentiation antigen (MNDAs), which is cleaved by caspase-3 and the cleaved product promotes Mcl-1 proteasomal degradation; and (D) proteinase 3, which leaks from the granules of aging neutrophils to directly cleave caspase-3 and activate the apoptotic programme.

New Ways to Die – Neutrophil Cell Death

Neutrophils have a relatively short lifespan in the circulation, estimated at 5 days [31]. The rate and mode of neutrophil death has been studied intensively over the last decade. New insights have been gained into the mechanisms of neutrophil apoptosis and the potential anti-inflammatory consequences of promoting accelerated apoptosis. Furthermore, non-apoptotic forms of neutrophil cell death have been identified that may lead to pro-inflammatory consequences in both infective and autoimmune diseases [32].

Neutrophil Apoptosis

The exact mechanisms behind spontaneous neutrophil apoptosis have not yet been defined, although a number of recent studies have highlighted unexpected pathways to apoptosis (Fig. 2). Neutrophils do not appear to employ the same cell death pathways characterised by other cell types [33]. The anti-apoptotic protein Mcl-1 is abundantly expressed in neutrophils and may play a role in governing the timing of apoptosis [34], whereas Bcl2, the founding member of this family of anti-apoptotic proteins and widely expressed in other cell types, is not expressed in mature neutrophils [35].

Caspases are important in the execution phase of spontaneous apoptosis, but the mechanism by which they become cleaved and activated in neutrophils is not clear. A recent paper highlighted a novel role for the neutrophil serine protease proteinase 3 (PR3) as a mediator of spontaneous neutrophil apoptosis. With neutrophil ageing, the study found that PR3, normally sequestered within neutrophil granules, is released into the cytoplasm, where it is able to cleave caspase 3, inducing apoptosis. The role of PR3 was demonstrated in vivo using adoptive transfer of PR3-deficient and -sufficient neutrophils in the inflammatory context of thioglycolate peritonitis. PR3-deficient neutrophils survived longer than wild-type PR3-sufficient neutrophils [36]. In contrast, our recent study using novel transgenic mice expressing human PR3 (whose enzymatic activity is different from its murine counterpart) showed a defect in the resolution of inflammation associated with an increased survival of neutrophils expressing human PR3. Furthermore, transgenic mice expressing human PR3 have an increased mortality during sepsis [37].

Ingestion of apoptotic neutrophils by macrophages has the ability to downregulate the inflammatory response through modification of macrophage function [38]. However, when PR3 is expressed at the neutrophil membrane in an inflammatory context, it alters the nature of phagocytosis by macrophages, promoting a pro-inflammatory rather than anti-inflammatory response [39]. This mechanism appears to be of relevance in anti-neutrophil cytoplasmic antibody-associated vasculitis, a severe inflammatory disorder characterised by the presence of circulating autoantibodies against neutrophil granule enzymes including PR3 [40].

Along with PR3, a number of other unexpected mediators of neutrophil apoptosis have been identified that had not previously been characterised in other cell types. Myeloid cell nuclear differentiation antigen (MNDNA) is a protein with roles in cell differentiation that has been discovered to influence neutrophil apoptosis [41]. Nuclear MNDNA is cleaved by caspases early during apoptosis and translocates to the cytoplasm, where it promotes proteasomal degradation of Mcl-1, accelerating apoptosis. Knockdown of MNDNA in HL-60 granulocytic cells decreased susceptibility to apoptosis.

Another protein better characterised as a mediator of cell proliferation, proliferating cell nuclear antigen (PCNA), has been identified as a novel mediator of neutrophil apoptosis [42]. Normally expressed primarily in the nucleus in the majority of cell types, PCNA is expressed in the cytoplasm of mature neutrophils, where it

sequesters pro-caspases, limiting their activation and delaying apoptosis [43]. Peptides derived from the endogenously expressed cyclin-dependent kinase (CDK) inhibitor p21 displace the pro-caspases from PCNA and facilitate activation to trigger neutrophil apoptosis [44].

A related group of cell proliferation proteins, the CDKs (in particular CDK9 [45]), have been identified to modulate neutrophil apoptosis. The CDK inhibitor roscovitine induces neutrophil apoptosis and has been demonstrated to have an anti-inflammatory effect in vivo, harnessing the pro-resolving capabilities of apoptotic neutrophils [46, 47]. Finally, the long non-coding RNA *Morrbid* has been identified as a novel regulator of myeloid cell lifespan, including that of neutrophils [48]. The mechanism involves regulation of the transcription of Bim, a pro-apoptotic molecule.

Following apoptosis, neutrophils that are not phagocytosed in a timely manner may undergo secondary necrosis, spilling the neutrophil contents into the surrounding tissues and possibly contributing to exacerbated inflammation. One recent study found that during apoptosis, neutrophils processed the pro-form of the cytokine IL-16C through the action of caspase 3 but were only able to release this cytosolic protein once the membrane had been compromised [49]. Secondary necrosis also leads to release of other pre-formed molecules such as pleiotropic migration inhibitory factor [49]. This parallels work on macrophages describing the release of IL-1 β , which is cytosolic and does not possess export sequences, necessitating cell death for release [50].

Immune complexes, implicated in the pathogenesis of a number of inflammatory diseases, have been found to induce neutrophil apoptosis through a non-canonical PI3K-Cdc42-Pak-Mek-Erk pathway, resulting in alterations in the proportions of pro- and anti-apoptotic molecules [51].

Non-Apoptotic Cell Death

The field of cell death has expanded greatly in the last decade to embrace newly described forms of non-apoptotic cell death including newly uncovered forms of “programmed” cell death. Whether or not neutrophils can undergo these forms of death, or indeed whether there are neutrophil-specific forms of cell death, is a field of ongoing study, with recent findings highlighted below.

NETosis – NET Release

One form of non-apoptotic neutrophil cell death has been termed NETosis or neutrophil extracellular trap release-associated cell death [52]. The central characteris-

Fluorescent image

Phase contrast image

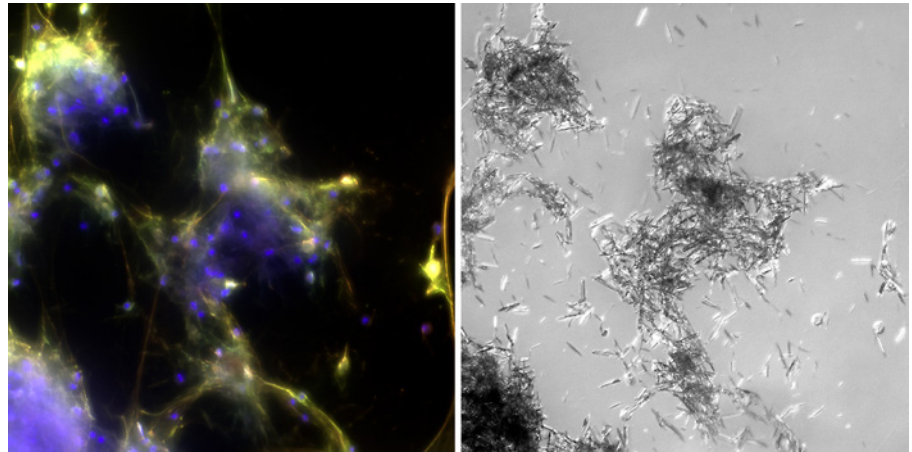


Fig. 3. Neutrophil death with DNA release and decondensation triggered by monosodium urate crystals. Healthy control neutrophils incubated with monosodium urate crystals (0.2 mg/mL) for 4 h. Maximum-intensity projection of a z-stack. The fluorescent image is a merge of blue (DAPI), green (neutrophil elastase) and red (myeloperoxidase). DNA colocalised with neutrophil granule proteins aggregates the crystals.

tics of this form of neutrophil demise are caspase-independent cell death and the release of DNA coated in neutrophil granule proteins. The exact mechanisms and clinical relevance of this form of cell death have been the focus of a large amount of research, with many questions remaining [53]. Key questions about NETosis and NET release remain, including whether DNA release can occur without triggering neutrophil death, as has been described *in vitro* for the release of mitochondrial DNA [54] and *in vivo* using two-photon microscopy in mice [55]. The physiologic and pathologic stimuli that induce NETosis need to be better defined, as a number of studies describing the mechanistic pathways of NETosis have been performed using the potent non-physiologic cell stimulant PMA [53, 56].

One scenario that appears to trigger this process is the attempted phagocytosis of large pathogens such as fungal hyphae [57]. As a consequence of the size of the hyphae, instead of forming a discrete pathogen-containing phagosome, the neutrophil granule enzymes are released into the cytoplasm in a ROS-dependent manner to enter the nucleus, degrade histones and allow the chromatin to decondense prior to cell membrane compromise. Intriguingly, neutrophils also respond to large particles by producing IL-1 β [58]. In comparison to the phagocytosis of small particles where intracellularly generated ROS leads to oxidation of NF- κ B and inhibition of IL-1 β generation, ROS are instead directed extracellularly. These two neutrophil responses create a strategy of amplified neutrophil recruitment and local deposition of DNA and neutrophil granule proteins to ensnare the large hyphae and control growth.

As discussed above, one important consequence of apoptosis in neutrophils is that it triggers orderly dismantling of the cell and labelling of the cells for phagocytosis by macrophages, such that the potentially toxic neutrophil contents are sequestered during cell death. Furthermore, ingestion of apoptotic neutrophils triggers an anti-inflammatory response in the macrophages themselves [59]. Cell death with release of intracellular contents has the potential to circumvent this process. DNA release in the context of neutrophil cell death has been found to prime macrophages to release IL-1 β in the context of atherosclerotic disease [60] both *in vitro* and *in vivo*.

The interaction of dead neutrophils with clearing phagocytes may be further perturbed by bacteria. *Staphylococcus aureus* was found to degrade the extracellular DNA structures released by neutrophils, not only dismantling their net-like properties, but converting the nucleic acids into deoxyadenosine, which resulted in caspase-mediated cell death of macrophages in proximity [61].

Another scenario where neutrophil cell death and DNA release may be of physiologic importance is in the human inflammatory joint disease gout. *In vitro*, monosodium urate (MSU) crystals are phagocytosed by neutrophils and, unlike hyphae, are encased in a maturing phagosome. Neutrophils subsequently undergo rapid cell death with post-mortem decondensation of the chromatin [62] (Fig. 3). This raises the question whether this form of cell death is programmed or a form of unregulated cell disruption occurring in the context of neutrophil granule release, resulting in dismantling of the chromatin. Extracellular DNA structures associated with neutrophil granule proteins are found to be associated with MSU crystals taken from the joints of gout patients, sug-

gesting the process is occurring in this context. Using an in vivo model of MSU peritonitis, release of DNA leads to the formation of large aggregates of crystals that not only sequester the crystals but also actively degrade pro-inflammatory cytokines, leading to downregulation of inflammation [63]. This is in keeping with the clinical observations that gout attacks are self-limited and large aggregates of MSU crystals form around the joints of gout patients but remain largely uninfamed.

Other Forms of Non-Apoptotic Cell Death

Other forms of programmed, non-apoptotic cell death have been characterised in the last decade, including necroptosis and pyroptosis. The majority of studies on these forms of death have been undertaken in cells other than neutrophils. In one study [64], *Staphylococcus aureus* triggered a non-apoptotic form of lytic cell death that involved RIPK3, a key component of the necroptotic death pathway, but did not involve RIPK1, the upstream component, and the death was not accompanied by MLKL membrane translocation, the critical downstream step leading to cell lysis in necroptosis. This study cautioned against relying on the results of experiments using chemical inhibitors of the necroptosis pathway, such as necrostatin-1, as evidence of necroptosis.

Neutrophil phagocytosis-induced cell death has been difficult to characterise and may depend on ROS generation and rapid caspase activation [65]. The phagocytosed pathogen plays an active role in the death pathway, exemplified by bacteria such as *Francisella novicida* that inhibit phagocytosis-induced cell death. This adaptation may serve as a survival advantage for this intracellular organism [66]. Another organism, *Salmonella*, induces macrophage pyroptosis, a form of programmed non-apoptotic cell death mediated by caspase 1 cleavage of gasdermin and accompanied by IL-1 β release [67]. Neutrophils, on the other hand, despite containing the machinery to respond to *Salmonella* including the inflammasome NLRC4, resist pyroptosis whilst retaining the capacity to release active IL-1 β . This heterogeneity in cell death and response to different pathogens reveals the complexity of neutrophil function [68].

Neutrophil-Derived Microvesicles – Novel Messengers That Modulate Inflammation

Microvesicles are tiny membrane-bound structures released by leukocytes or endothelial cells that can circulate throughout the body, carrying with them various im-

munomodulatory molecules that allow them to remotely modify inflammation [69]. Microvesicles are surrounded by a membrane containing high levels of phosphatidylserine, a lipid that is normally located inside the cell. Interestingly, using a unique hydrophobic patch, neutrophil-derived PR3 can bind to phosphatidylserine, a major component of microvesicles, thereby affecting both their production and their function to ultimately promote inflammation [70]. Conversely, microvesicles can have potent anti-inflammatory effects. As an example, a recent report has shown that neutrophil-derived microvesicles can enter cartilage and protect the joint in inflammatory arthritis [71]. In this setting, the anti-inflammatory protein annexin A1, which is also a phosphatidylserine-binding protein, bound to microvesicles. Annexin A1 could next interact with its receptor FPR2 (formyl peptide receptor 2)/ALX, increasing transforming growth factor- β production by chondrocytes, ultimately leading to cartilage protection. Microvesicles are emerging as a novel means for neutrophils to deliver proteins remotely to other cells with an impact on inflammation. Harnessing the pro-resolving effects of microvesicles may serve as a future therapeutic opportunity.

Conclusions

The last decade has uncovered some surprising aspects to neutrophil biology. Neutrophil heterogeneity at multiple levels, from neutrophil maturity, functionality and ultimately demise, has led to a reconsideration of the role of neutrophils in many diseases ranging from inflammatory disorders over cardiovascular and thrombotic diseases to cancer. The future of neutrophil research looks bright, but challenges remain. Translating discoveries into neutrophil-targeting therapeutics is the next critical step.

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Disclosure Statement

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