REVIEW



The intricate relationship between autoimmunity disease and neutrophils death patterns: a love-hate story

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Abstract

Autoimmune diseases are pathological conditions that result from the misidentification of self-antigens in immune system, leading to host tissue damage and destruction. These diseases can affect different organs and systems, including the blood, joints, skin, and muscles. Despite the significant progress made in comprehending the underlying pathogenesis, the complete mechanism of autoimmune disease is still not entirely understood. In autoimmune diseases, the innate immunocytes are not functioning properly: they are either abnormally activated or physically disabled. As a vital member of innate immunocyte, neutrophils and their modes of death are influenced by the microenvironment of different autoimmune diseases due to their short lifespan and diverse death modes. Related to neutrophil death pathways, apoptosis is the most frequent cell death form of neutrophil non-lytic morphology, delayed or aberrant apoptosis may contribute to the development anti-neutrophil cytoplasmic antibodies (ANCA)-associated vasculitis (AAV). In addition, NETosis, necroptosis and pyroptosis which are parts of lytic morphology exacerbate disease progression through various mechanisms in autoimmune diseases. This review aims to summarize recent advancements in understanding neutrophil death modes in various autoimmune diseases and provide insights into the development of novel therapeutic approaches for autoimmune diseases.

 $\textbf{Keywords} \ \ Neutrophil \cdot Cell \ death \cdot Pyroptosis \cdot Neutrophil \ extracellular \ traps \cdot Autoimmune \ diseases \cdot Inflammation$

Abbreviations

ANCA	Anti-neutrophil cytoplasmic antibodies
RA	Rheumatoid arthritis
SS	Sjogren's syndrome
UC	Ulcerative colitis
MS	Multiple sclerosis
NE	Neutrophil elastase
GM-CSF	Granulocyte colony-stimulating factor
NETs	Neutrophil extracellular traps
DAMPs	Damage-associated molecular patterns

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TNF-α	Tumor necrosis factor-α
GPA	Granulomatosis with polyangiitis
TRAIL	TNF-related apoptosis-inducing ligand
LRP	Lipoprotein receptor-related protein
HA	Hyaluronic acid
PD-L1	Programmed cell death-ligand 1
PBMC	Peripheral blood mononuclear cell
ANA	Antinuclear antibodies
HNP	Human neutrophil peptide
TLR9	Toll-like receptor 9
IC	Immune complexes
MPA	Microscopic poly-vasculitis
HMGB1	High mobility group protein 1
ACPAs	Anti-citrullinated peptide antibodies
SF	Synovial fluid
TEN	Toxic epidermal necrolysis
LPMC	Lamina propria mononuclear cells
TNBS	2,4,6-Trinitrobenzene sulfonic acid
pfCD	Perianal fistulising Crohn's disease
μPs	Microparticles
aPL	An-tiphospholipid antibody
FAP-a	Fibroblast-activated protein-a
MRSA	Methicillin-resistant staphylococcus aureus

AS	Ankylosing spondylitis
AIH	Autoimmune-mediated hepatitis
HSCs	Hematopoietic stem cells
LMWH	Low molecular weight heparins
PMN	Polymorphonuclear neutrophils
EGPA	Eosinophilic granuloma with poly-vasculitis
Hif-1α	Hypoxia-inducible factor-1α
LAMP-2	Lysosomal membrane protein-2
RNP IC	Immunocomplexes containing
	ribonucleoproteins
TIM-3	T cell immunoglobulin and mucin domain-
	containing protein 3
MCPIP-1	Monocyte chemotactic protein-inducible
	protein-1
AAV	ANCA-associated vasculitis
SLE	Systemic lupus erythematosus
SSC	Systemic sclerosis
CD	Crohn's disease
MPO	Myeloperoxidase
ROS	Reactive oxygen species
Cyt-C	Cytochrome C
CTL	Cytotoxic T lymphocytes
NK	Natural killer cells
PR3	Proteinase 3
CRT	Calreticulin
APS	Antiphospholipid syndrome
ECM	Extracellular cell matrix
IFN-I	Type I interferon
MIF	Migration inhibitory factor
PAD4	Peptidylarginine deiminase 4
LDG	Low-density granulocytes
pDC	Plasmacytoid dendritic cells
MMP-9	Matrix metalloproteinase-9
EC	Endothelial cells
mDCs	Myeloid dendritic cells
TF	Tissue factor
IL-17A	Interleukin-17A
SJS	Stevens Johnson Syndrome
FPR1	Rormyl peptide receptor 1
HOCI	Oxidant hypochlorous acid
RRMS	Relapsing-remitting MS
ANETA	Anti-NET antibodies
PSGL-1	P-selectin
aCL	An-ticardiolipin antibody
MSU	Monosodium urate
	Plasma membrane
PM AOSD	
AOSD	Adult Still's disease (AOSD)
Atg	Autophagy related gene
CQ	Chloroquine
MA ET 1	3-Methyl-adenine
ET-1	Endothelin-1

Introduction

Autoimmune disease is a complex group of conditions marked by the breakdown of self-tolerance to various autoantigens. These multifaceted pathologies arise from a complicated interplay between genetic and environmental factors. In cases of pathological autoimmunity, the intrinsic immune system of the organism is mobilized against self-tissues, leading to the recruitment of autoantibodies, self-reactive lymphocytes, and other innate immune cells. This immune response culminates in a cascade of inflammatory processes, resulting in the infliction of tissue damage and the development of chronic inflammation [1]. Autoimmune disease is a multifaceted and heterogeneous illness with a wide array of clinical manifestations, characterized by the immune system of body attacking its own tissues and organs. This disease can be classified into two major categories: systemic and organ-specific autoimmune diseases [2, 3]. The systemic autoimmune diseases, including rheumatoid arthritis (RA), systemic lupus erythematosus (SLE), Sjogren's syndrome (SS) and systemic sclerosis (SSC) are distinguished by their pathologic impact on multiple organs and tissues. Conversely, organ-specific immune diseases, such as psoriasis, ulcerative colitis (UC), Crohn's disease (CD), and multiple sclerosis (MS), display a localized manifestation of autoantibodies accumulation in specific systems and organs. The pathogenesis of autoimmune diseases is intricate, involving a complex interplay between genetic, environmental, and psychological factors, leading to the activation of both innate and adaptive immune responses.

Within the immune system, macrophages, dendritic cells, and neutrophils are all considered to be integral players in the development and progression of autoimmune diseases [4, 5]. Specifically, neutrophils have garnered considerable attention in recent years due to their ability to impact the autoimmune process through both direct and indirect mechanisms. For example, in response to pathogenic invasion, such as bacterial infections, neutrophils detect and recognize the pathogens through specific receptors. Upon recognition, neutrophils utilize their toxic products, such as myeloperoxidase (MPO), neutrophil elastase (NE), reactive oxygen species (ROS), and cytokines, to perform their phagocytic function to eradicate the invading pathogens directly [6]. Nevertheless, excessive infiltration of neutrophils in sites of tissue injury can lead to chronic inflammation by releasing cytokines and proteases, thereby causing tissue damage. Furthermore, the infiltration of neutrophils can regulate the activity of adaptive immune responses by activating B cells and T cells, thereby leading to immune system imbalance indirectly [7]. Therefore, the relationship between neutrophil death and inflammation and autoimmunity deserves further exploration.

As a type of white blood cell, neutrophils are derived from hematopoietic stem cells and undergo maturation within the confines of the bone marrow. This process of granulopoiesis is marked by a series of distinct stages, each of which is characterized by a unique morphology and granular content. From promyelocytes to myelocytes to metamyelocytes, the neutrophilic precursors gradually adopt the classic segmented appearance and develop the characteristic intracellular granules that confer their potent antimicrobial properties. This complex process of neutrophil differentiation and maturation is crucial for the maintenance of immune homeostasis [8]. The intricate regulation of neutrophil development, production, and release is orchestrated by a multifaceted network of signaling molecules, but perhaps none is more central than the granulocyte colony-stimulating factor (GM-CSF). Acting through a cascade of downstream effectors, GM-CSF finely tunes the delicate balance of neutrophil homeostasis by suppressing the expression of CXR4, a receptor that promotes retention of immature neutrophils in the bone marrow, and simultaneously enhancing the activity of CXCR2, a receptor that stimulates the release of mature neutrophils into the bloodstream. This exquisite modulation of neutrophil trafficking and differentiation is essential for the proper functioning of the immune system and serves as a prime target for pharmacological intervention in a host of hematological disorders and inflammatory diseases [9]. Actually, neutrophils are the sentinel guardians of the innate immune system which are characterized by their remarkable ability to rapidly migrate to sites of infection or injury and unleash a potent arsenal of antimicrobial effector molecules. However, this prodigious performance comes at a cost, as these cells are inherently short-lived and have a limited lifespan in the circulation, with a half-life of a mere 8 h in humans. The physiological range of neutrophil count in a healthy individual is estimated to be within the range of 1500-8000 per microlitre of blood, which contributes to the remarkable surge of millions of these cells each day. The high turnover rate of neutrophils reflects the tremendous metabolic demands required for their effector function, as well as the need to prevent the accumulation of damaged or dysfunctional cells that could cause tissue damage or autoimmunity. While this tight regulation of neutrophil turnover is critical for immune homeostasis, it also poses a major challenge for therapeutic targeting, as any intervention that alters neutrophil survival or turnover could have far-reaching consequences for immune function and host defense [10].

Recent research has revealed that neutrophils possess multiple programmed cell death pathways, which can be categorized into lytic and non-lytic morphologies based on the final cell death morphology [11]. Apoptosis, the most common non-lytic morphology, is characterized by 'a dance of death' and is typically immunologically silent [12]. Conversely, lytic morphology pathways such as necroptosis, pyroptosis, neutrophil extracellular traps (NETs) and ferroptosis release cellular molecules by rupturing the plasma membrane, resulting in the formation of damage-associated molecular patterns (DAMPs) that promote inflammatory reactions. Consequently, autoimmune diseases and chronic inflammation can develop [13, 14]. The importance of studying neutrophil death in autoimmune disease mechanisms and treatment modalities is emphasized by these findings.

However, the significance of survival mechanisms cannot be overlooked even in short-lived cells such as neutrophil. Autophagy plays a crucial role in maintaining the normal state and functionality of neutrophil. Unlike the diverse modes of cell death, autophagy is activated in response to accumulated cellular damage, oxidative stress, and nutritional starvation. Its primary function is to degrade damaged organelles, unwanted proteins, and other substances, thereby preserving overall cellular homeostasis. Recent studies have revealed that autophagy not only acts as a conserved and self-protective mechanism in neutrophil but also exhibits intricate interactions with various cell death pathways, including apoptosis, NETosis, pyroptosis, and the release of inflammatory factors. Consequently, autophagy becomes interdependent and plays a crucial role in the development of numerous autoimmune diseases. Therefore, the relationship between autophagy and the mode of neutrophil death in autoimmune diseases has also attracted interest.

Here, we reviewed recent progress in understanding the pathogenesis, abnormalities of different neutrophil death pathways and relationship between survival and death mechanism in autoimmune diseases, with the goal of developing targeted therapeutic agents to modulate neutrophil death.

Apoptosis: the king of non-lytic neutrophil cell death

Apoptosis is a regulated morphology that widely exists in the neutrophil cell death kingdom. During the apoptosis, no intracellular components are released outside the cell, the cell shape becomes round and the volume decreases. What's more, the nucleus shrinks and breaks, chromatin condenses along the nucleus, typical cytoplasmic organelles do not change in ultrastructure. However, the plasma membrane does not break. Eventually, the cell fragments were swallowed by phagocytes, and were degraded by proteases and nucleases in lysosomes after being transported to lysosomes, which prevented the inflammatory contents of cells from being released to the surrounding areas and was beneficial to tissue renewal [15]. Therefore, normal apoptosis as a nonlytic morphology of the neutrophil cell death can maintain the stability of the organism.

Currently there are two known pathways of apoptosis, endogenous pathway and exogenous pathway. Endogenous apoptosis refers to the imbalance of intracellular homeostasis caused by the changes of microenvironment such as stimulation of toxic substances, DNA damage and secretion of growth factors, which leads to the increase of permeability of mitochondrial outer membrane and entrance of Cytochrome C (Cyt-C) to the cytoplasm to activate the caspase cascade of apoptosis [16]. The endogenous pathway is usually controlled by Bcl-2 protein family. Some proteins (such as Bax and Bid) promote the occurrence of apoptosis while some proteins (such as Bcl-2) have the opposite effect. It is the balance between these pro-apoptotic proteins and anti-apoptotic proteins that ensures the normal progress of apoptosis [17]. Exogenous pathway refers to the pathway used by cytotoxic T lymphocytes (CTL) and natural killer cells (NK) to attack their target cells, which is initiated by the activation of tumor necrosis factor- α (TNF- α), Fas ligand and TNF-related apoptosis-inducing ligand (TRAIL) receptors DR4 and DR5 located on the cell surface.

As a non-lytic neutrophil cell death pathway, normal apoptosis is considered to be an effective strategy to avoid tissue damage which can eliminate neutrophil and prevent neutrophil from secreting large amounts of inflammatory cytokines [13, 18]. Nevertheless, recent studies show that abnormal apoptosis are associated with autoimmune diseases.

Some autoimmune diseases are associated with the untimely clearance of apoptotic neutrophil, where the phagocytic clearance of apoptotic neutrophil is impaired, thus triggering the autoantigenic clusters that maintain autoimmune disease. However, it has been also shown that some autoimmune diseases exhibit delayed apoptosis of neutrophil, resulting in the accumulation of neutrophil that can lead to the release of inflammatory factors and cytokines and the recruitment to other immune cells, which leads to inflammation eventually. In conclusion, abnormal neutrophil apoptosis is likely to be the underlying cause of some autoimmune diseases.

Disrupted clearance mechanisms of apoptotic neutrophils in autoimmune pathology

In recent years, many targets involved in neutrophil death have been extensively studied, which have been certified to be related to the development of autoimmune diseases, especially SLE. This section will explore the relationship between apoptotic neutrophil clearance and autoimmune diseases, focusing on the 'key players' in apoptotic neutrophil clearance.

Autoantigen protease 3 (PR3), also known as myeloblastin, is a kind of serine protease existed in neutrophil azurophilic granules, which can regulate granulocyte differentiation [19]. It degrades a variety of extracellular matrices such as elastin, haemoglobin, collagen type IV and many other tissue components. Generally speaking, the expression of PR3 on cell surface is increased in apoptotic neutrophil, which limits their phagocytic clearance by macrophages and promotes a pro-inflammatory microenvironment [20–22]. In recent research, PR3 is considered as the target for ANCA in granulomatosis with polyangiitis (GPA) [23, 24]. Studies have shown that PR3 enhances the degree of the occurrence of certain autoimmune diseases such as GPA and RA [19].

The efferocytosis of macrophages prevent the release of intracellular substances and the autoimmune response of body to its own antigens. If efferocytosis is impaired, it can lead to autoimmune diseases or tissue damage [25]. During apoptotic neutrophil death in GPA patients, increased expression of PR3 on the membrane induces abnormal phagocytosis of apoptotic neutrophils by macrophages thereby inducing an inflammatory response [19]. It has been shown that PR3 causes abnormal macrophage efferocytosis associated with calreticulin (CRT), an endoplasmic reticulum chaperone protein that releases a 'eat me signal' in the membrane of apoptotic neutrophils, thereby inducing macrophages to take up the apoptotic neutrophils. This promotes the regression of inflammation. During apoptotic neutrophil death in GPA patients, PR3 may block the binding of CRT to one of its major receptors, the low-density lipoprotein receptor-related protein (LRP), and the complex of PR3 and CRT may also impair the function of LRP and thereby disrupt the phagocytic function of macrophages [26], preventing the timely clearance of apoptotic neutrophils and ultimately amplify inflammation. Apoptotic neutrophils may also bind to cytokines at the site of inflammation, leading to autoimmune [23]. In patients with vasculitis, the combination between ANCA and PR3 also modulates neutrophil apoptosis and thus has an effect on phagocytosis. However, PR3-ANCA can lead to increased uptake by macrophages, resulting in TNF- α production to induce inflammation and autoimmunity.

Inflammation and autoimmunity through autoantibody modulation of apoptotic cells may be important not only in vasculitis but also in other autoimmune diseases [27]. Antiphospholipid syndrome (APS), SLE, may lead to inflammation through the same mechanism [28, 29]. Therefore, PR3 and its conjugates with other substances can not only modulate neutrophil apoptosis, but also have an impact on the clearance of neutrophils by macrophages and can be an important therapeutic target for certain autoimmune diseases (Table 1).

In fact, it can be seen from the relationship between PR3 and apoptotic neutrophil clearance that phagocytosis of macrophages plays an important role in the process of apoptotic cells clearance. Phagocytosis removes altered self-cells, especially apoptotic cells, and complement protein C1q is one of the mediators of phagocytosis which plays an important role in preventing the formation of

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Type of neutrophil cell death	Potential drug intervention Mechanism	Mechanism	Study design	Effect on neutrophil cell death	Autoimmune diseases	Effect on disease clinical endpoint	References
Necroptosis	KE-86-QG38 GSK9843/ GW806742X	FAP-a RIPK3/ MLKL inhibitor	Peripheral blood neu- trophils from healthy normal individuals and patients suffering from RA	Respectively blocked CD44-triggered ROS production and cell death of joint fluid RA neutrophils	RA	Not tested	[108]
	Ptpn6	Autoimmune and interleu- kin-1 (IL-1) receptor- dependent, caspase-1 independent inflamma- tory disease preventive medicine	Ptpn6 ^{APMN} mice	Ptpn6 regulate negatively the activation of p38 mitogen- activated protein kinase to maintain Ripk1 function to prevent caspase-8- and Ripk3-MLKL-dependent cell death	Inflammation	Not tested	[163]
Apoptosis	DNase1 DNase1 like 3(DNase1L3)	DNA degradation	SLE mouse model	Degraded the DNA produced by apoptotic cell buildup, Decreased NETs formation	SLE	Not tested	[164–166]
	Milk fat globule-EGF fac- tor 8 (MFG-E8)	Macrophage phagocytosis stimulant Necrosis and NETs inhibi- tors	Pristane-induced lupus mouse model	Enhanced the uptake of apop- totic neutrophils, inhibited the migration of neutrophils	SLE	Not tested	[167, 168]
	Pycnogenol	ROS inhibitor	Phase I study	Decreased ROS production which reduces neutrophil apoptosis and autoantibody production	SLE	Enhance immune and hematopoietic function in the body	[169]
	IgA	Apoptosis agonist	Peripheral blood neu- trophils from patients suffering from CD	Induced neutrophil apoptosis in a PI3K-, P38 MAPK- and JNK- dependent manner under cytokine stimulation	CD	Not tested	[170]
	Fe (II)-IVIG	Apoptosis agonist	Peripheral blood neu- trophils from patients suffering from CD	Induced neutrophil apoptosis in a Fas dependent manner under low PH environment	CD	Not tested	[171]
NETs	Thrombomodulin (TM)	NETs formation inhibitor	Experimental autoimmune vasculitis rat model	Binded to neutrophils via macrophage-1 antigen (mac- 1),resulting in a suppression of ANCA-induced NETs formation	AAV	Not tested	[172]
	IVIG-S	Unknown	MPO-AAV rat model	Reduced NETs formation and ANCA titles	AAV	Not tested	[173]
	MitoQ	Mitochondria- targeted Antioxidant	Lupus prone MRL- lpr mice treated with MitoQ	Reduced neutrophil ROS and NETs formation	SLE	Not tested	[174]

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Table 1 (continued)	nued)						
Type of neutrophil cell death	Potential drug intervention Mechanism	Mechanism	Study design	Effect on neutrophil cell death	Autoimmune diseases	Effect on disease clinical endpoint	References
	PA-dPEG24	Inhibitor of complement C1	Human neutrophils stimu- lated by PMA, MPO	Inhibited complement activa- tion, limiting the generation of proinflammatory complement effectors and inhibited NETs formation	SLE	Not tested	[175]
	Celastrol	Unknown	Healthy donors SLE and RA patients neutrophils stimulated by TNFα, Ova IC, total serum IgG	Inhibited neutrophil oxidative burst and NETs formation induced by different inflam- matory stimuli and citrullina- tion of histones	RA/ SLE	Not tested	[176]
	Polydatin (PD)	Unknown	Neutrophils from SLE/ RA patients and healthy people stimulated with PMA Pristane induced lupus mice collagen- induced arthritis mice	Inhibited ROS mediated NETs formation	SLE/RA	Not tested	[177, 178]
NETS	Tofacitinib	Janus kinase (JAK) inhibitor	MRL/lpr lupus-prone mice	Modulated NETs formation and significantly increased endothelium dependent vasorelaxation and endothelial differentiation	SLE	Lead to significant improvement in meas- ures of disease activity including nephritis, skin inflammation, and autoantibody production	[621]
	Cl-amidine BB-Cl-amidine	PAD inhibitors	MRL/lpr mice	Decreased histone citrullination and reduced release of NETs	SLE	Not tested	[59]
	Rituximab (RTX) and Belimumab (BLM)	Blocking IC formation	Phase II study	Interfered with ICx formation and decreased NETs forma- tion	SLE	Decreased lupus disease activity	[180]
	Emodin	Unknown	Murine adjuvant induced arthritis (AA) model of RA	Promoted neutrophil apoptosis thereby reducing proinflam- matory cytokines and inhib- ited autophagy and NETs	RA	Not tested	[43]
	Infliximab Tocilizumab	TNF-α inhibitor IL-6r inhibitor	Phase II study	Inhibited NETs extrusion	RA	Abridged the endothelial dysfunc- tion and the activation of immune cells, thus influencing the global activity of the vascular system	[181]

Table 1 (continued)	inued)						
Type of neutrophil cell death	Potential drug intervention Mechanism	Mechanism	Study design	Effect on neutrophil cell death	Autoimmune diseases	Effect on disease clinical endpoint	References
	Quercetin	Neutrophil inhibitors of inflammatory activity	AA model	Promoted the apoptosis of activated neutrophils and inhibited NETs formation by suppressing autophagy	RA	Inhibited neutrophil infil- [182] tration and reduced the plasma levels of inflam- matory cytokines	[182]
	Tanshinone IIA	Antioxidant	AA model	Promoted neutrophil apoptosis and inhibited NETs formation	RA	Alleviated cartilage ero- sion and neutrophil infiltration in the ankle joints and reduced pro- inflammatory cytokines expression levels in sera	[183]
	Recombinant human ADAMTS-13 (rhAD- AMTS- 13)	Von Willebrand factor (VWF) nickase	Mouse model of collagen- induced arthritis (CIA)	Inhibited The formation of NETs and deposition of VWF in synovial membrane	RA	Not tested	[184]
Ferroptosis	LPX-I	Ferroptosis and ROS inhibitor	Lupus-prone MRL/lpr mice	Significantly mitigated disease progression, and reduced the production of auto-antibodies and various inflammatory cytokines	SLE	Not tested	[145]

antigen-antibody complexes and in the activation of the complement pathway [30].

C1q binds tightly to CD91 and CRT on the surface of apoptotic neutrophils, recognising apoptotic neutrophils and enabling phagocytes to perform their phagocytosis. In the presence of CRT, the addition of extracellular C1q accelerates the uptake of cellular debris. However, in patients with SLE, C1q and the bound proteins and receptors are absent or blocked, and anti-CRT and anti-C1q antibodies have been found in the serum [31]. The recognition of apoptotic cells by C1q is also affected by the aforementioned PR3, which is involved in the clearance of apoptotic neutrophils and the regulation of the immune response, and shares ligands such as CRT. It has been found that PR3 binds to C1q, thereby impairing C1q recognition of apoptotic neutrophils and resulting in defective clearance of apoptotic neutrophils [32]. In conclusion, C1q acts by promoting the clearance of apoptotic neutrophils, and when C1q is deficient or genetically mutated, it can lead to the development of autoimmune diseases such as SLE and RA [33]. The abnormal interaction between C1q, PR3 and CRT leads to the development of autoimmune diseases by delaying the clearance of apoptotic neutrophils, providing a new idea and therapeutic target for the treatment of autoimmune diseases characterized by defective clearance of apoptotic neutrophils.

In addition, CD44 is an adhesion molecule involved in extracellular cell matrix (ECM) interactions [34] and is widely expressed in a variety of tissues and involved in cell adhesion and migration, lymphocyte activation and proliferation [35]. The interaction of CD44 with hyaluronic acid (HA) has been extensively studied [36] and this action is thought to be closely related to the regulation of the immune response [37–39]. In SLE, reduced expression of CD44 on the surface of neutrophils leads to failure of macrophages to clear apoptotic neutrophils, resulting in secondary necrosis and ultimately the release of autoantigenic clusters that can trigger autoimmunity [40, 41]. The ability of CD44 to promote cell clearance may be due to its association with cytoskeletal molecules, whereas in SLE, changes in the cytokine milieu such as the release of IL-10 lead to the downregulation of CD44, which results in impaired clearance of apoptotic neutrophils and promotes the development of autoimmune disease. It is easy to see that although CD44 plays an important role in immune regulation, the mechanisms that promote inflammation in different autoimmune diseases vary [35]. Therefore, understanding the role of CD44 in the development of inflammation and autoimmune diseases is essential for the use of this molecule as a drug target.

Overlived neutrophils make contribution to autoimmune disease

Over-activation or delayed apoptosis of neutrophils can lead to continued inflammation and amplification of the inflammatory response, which can lead to tissue damage [42, 43]. These processes occur in a variety of autoimmune diseases, of which RA is the most representative.

Programmed cell death-ligand 1 (PD-L1) is a ligand for programmed death 1 (PD-1) and is thought to be one of the key factors that inhibits T cell signaling, mediate tolerance mechanisms and provide immune homeostasis [44, 45]. Recent studies have revealed that PD-L1 can regulate neutrophil apoptosis and thus influence inflammation [46]. PD-L1 upregulates and interacts with the p85 subunit of PI3K to phosphorylate downstream Akt and delay neutrophil apoptosis, which in turn causes inflammatory responses and tissue damage [46]. As in RA, upregulation of PD-L1 causes delayed apoptosis of neutrophils, which results in massive infiltration of neutrophils in the synovium and contributes to inflammation. Monocyte chemotactic protein-inducible protein-1 (MCPIP-1), also known as Regnase-1, is a potent inhibitor of the inflammatory response [29], which appears to play an important role in neutrophil death. It can limit tissue damage by specifically targeting genes in different cell types through its ribonuclease activity and thereby reducing inflammatory factor storms [47]. MCPIP-1 controls the balance between anti- and pro-apoptotic proteins in neutrophils, thereby regulating their lifespan. MCPIP-1 is upregulated in dead neutrophils, but this effect is counteracted by GM-CSF, which causing a delay in neutrophil apoptosis, which leads to an excessive accumulation of neutrophils, resulting in inflammation and tissue damage, and even the development of autoimmune diseases [48]. In SLE, elevated type I interferon (IFN-I) expression promotes the expression of MCPIP-1, which in turn promotes reduced production of miR-146a, which has a suppressive autoimmune function [49]. However, therapeutic approaches targeting MCPIP-1 need to be cautious, as its complete inhibition or over-activation may produce detrimental immunopathological outcomes, and the search for therapeutic strategies with modulatory effects on MCPIP-1 could lead to promising results in autoimmune diseases. In addition, the chemokine-like inflammatory cytokine macrophage migration inhibitory factor (MIF) has been shown to be associated with neutrophil apoptosis, which contributes to the development of RA, MIF binds to its receptor CXCR2 and releases the survival factor CXCL8 from peripheral blood mononuclear cell (PBMC), thereby causing delayed neutrophil apoptosis and resulting in neutrophil infiltration, leading to inflammation [50]. It has also been shown that high intracellular K⁺ concentration or inhibition of K⁺ efflux inhibits neutrophil apoptosis by

suppressing cytochrome c-dependent apoptotic vesicle production [51]. In patients with RA, neutrophils are exposed to pro-inflammatory mediators in the synovial fluid, thereby delaying apoptosis and contributing to the amplification and continued production of inflammation [52]. It follows that in some autoimmune diseases such as RA, inhibition of delayed neutrophil apoptosis and thus inflammation may be the key to reducing the symptoms of autoimmune disease (Fig. 1).

NETosis: neutrophil-specific way of death

Neutrophils as the most abundant innate immune effector cells in the human immune system are equipped with a wide range of effective antibacterial agents and are mainly stored in specialized granules. Neutrophils normally protect the host against the spread of bacteria and fungi, but if they over defend, they can damage the host tissue. Therefore, its deployment is regulated by three strategies: phagocytosis, degranulation and NETs release. NETs are fibrous assembly networks of nuclei and particles that are released in activated neutrophil-cell membranes and can inhibit and kill bacteria outside the cell. NETs consist of citrulline group proteins and highly decompressed chromatin fibers released in a process known as NETosis [53]. Most of the DNA in NETs comes from the nucleus and also contains mitochondrial DNA. NETs appear to be faceless: normally, they can capture and kill pathogens, but beneath this mask may be the evil face of autoimmune disease.

There are many important roles in the formation of NETs. The DNA that makes up NETs does not have active transcription levels but condenses into heterochromatin in the nucleus. Heterochromatin decompression is mediated by peptidylarginine deiminase 4 (PAD4), which catalyzes the conversion of histone arginine to citrullines, reducing the strong positive charge of histones and consequently weakening histone-DNA binding. This attenuated interaction then opens up the nucleosome and constitutes a prerequisite for NETs formation [54]. In addition to PAD4, NE is an essential member of NETosis, which cleaves histones during the formation of NETs. In addition, MPO can accept the stimulation of ROS and trigger the activation and transport of NE [53, 55]. Therefore, the search for treatments for NETsrelated diseases can start from these closely related roles and explore new and effective treatments with NETs as the entry point.

As shown above, NETs contain histone and chromatin as well as important autoantigens, which increased NETs production, unique low density granulocyte populations and impaired NETs degradation leading to persistent inflammation and tissue damage related to autoimmune diseases. NETs can also activate other immune cells, and their components may amplify the inflammatory response by activating the complement pathway and inflammasome. NETs may also promote autoantibody formation in diseases such as RA, AAV and SLE by providing a constant source of autoantigens. Thus, NETs can be used as biomarkers to provide insights into disease diagnosis and treatment [56]. NETs are indeed strongly associated with a variety of autoimmune diseases, but the roles they play in the pathogenesis of various autoimmune diseases are changeable. We can understand that the same person plays a different role in different circumstances. Roles are different, but they can lead to seemingly the same result. In this review, we will focus on the role of NETs in several major autoimmune diseases. In addition, new drugs for the treatment of autoimmune diseases targeting NETs are summarized (Table 1).

NETs in SLE: chief culprit

SLE is a systemic autoimmune disease whose early pathological manifestations are mostly manifested in the skin and joints and the kidney, cardiovascular and other important organ systems are also affected. SLE patients will produce antinuclear antibodies (ANA) targeting a variety of nuclear components, such as anti-dsDNA antibodies, anti-ribosome antibodies, etc. Anti-dsDNA antibodies are highly specific and can form immune complexes with exposed chromatin to cause inflammation and promote disease [57]. Under normal conditions, a variety of dsDNA clearance mechanisms exist in cells. For example, dsDNA is rapidly degraded by DNA enzymes in the cytoplasm. Therefore, the failure of these mechanisms, such as inefficient clearance of apoptotic cell debris and NETs that are essentially DNA, will lead to the accumulation of dsDNA, inducing the production of autoantibodies and becoming the trigger of disease [58]. The relationship between NETs and SLE is not unfounded but traceable. Some researchers explored the effect of NETs on SLE using MRL/lpr mice which is a classic SLE mouse model and found that PAD inhibitors could reduce albuminuria and the deposition of immune complex in the kidney, which reduce the severity of skin diseases at the same time [59]. Immunocomplexes containing ribonucleoproteins (RNP IC) in lupus patients have also been shown to induce NETs. Simultaneously low-density granulocytes (LDG), a subset of proinflammatory neutrophils characteristic of SLE patients exhibit enhanced idiopathic NETs on biopsy [60].

Further investigation reveals that the role of NETs in SLE is closely related to IFN-I production. Researches show that SLE serum contains immune complexes of its own DNA, antimicrobial cationic peptide LL37 and human neutrophil peptide (HNP), which are protected from nuclease degradation and interact with the FC- γ receptor IIa on pDC to activate plasmacytoid dendritic cells (pDC) to release

IFN-I [61]. However, NETs in SLE contain DNA and a large amount of LL37, which contribute to pDC production of high levels of IFN- α in a manner which is dependent on DNA and toll-like receptor 9 (TLR9) [62]. At the same time, there is a positive feedback regulatory mechanism for NETs in SLE, that is, mature SLE neutrophils are activated by IFN-I and die after exposure to SLE derived anti-ribonucleoprotein antibodies and release NETs [63]. The released NETs then activate pDC to produce IFN-I, which aggravates the progression of SLE. Meanwhile, the increased content of histone acetylation and methylation residues in NETs in SLE patients may indirectly increase the stimulative capacity of NETs by enhancing the binding of antimicrobial peptides in NETs [64]. In recent years, researchers have found that the mechanism by which NETs aggravate SLE is not limited to IFN-I. In SLE patients with endothelium dependent impaired vasodilation, matrix metalloproteinase-9 (MMP-9) and anti-MMP-9 immune complexes (IC) enhance NETs in LDG. Meanwhile, LDG-derived NETs are the source of MMP-9 activation in endothelial cells (EC), triggering endothelial dysfunction [65]. Although mitochondrial DNA and chromatin are present in NETs, other components such as RNA, especially siRNAs, may play a role in immune disorders and tissue damage. Studies have shown that extracellular LL37 can bind to extracellular endogenous RNA and transfer it to the DC endosomes to activate the IFN-I response via TLR7/ TLR8 [66]. Meanwhile, abnormal downregulation of miR-4512 in monocytes and macrophages has been confirmed to promote the formation of NETs by targeting CXCL2 and TLR4 in SLE [67].

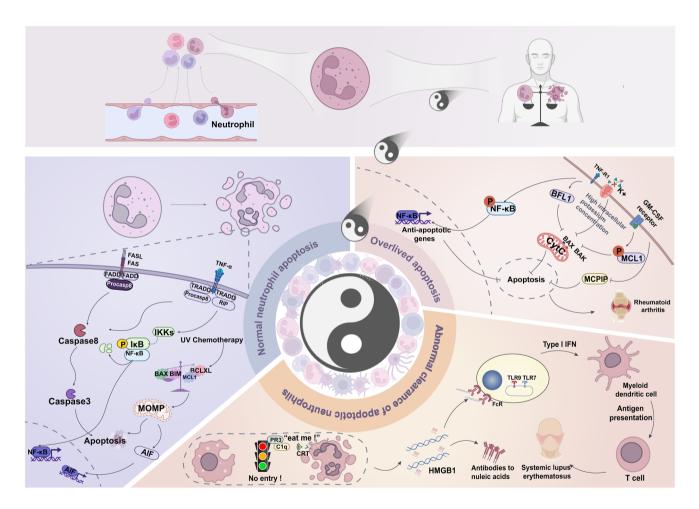


Fig. 1 Yin-yang balance of neutrophil apoptosis. In healthy individuals, the processes of normal neutrophil apoptosis and clearance of apoptotic neutrophil debris are balanced, and this equilibrium is crucial for maintaining homeostasis. Disruption of this balance can lead to autoimmune diseases. There are two known apoptotic pathways, the endogenous and exogenous pathways, which regulate neutrophil function. Abnormal clearance of apoptotic neutrophils can trigger autoimmune diseases, such as SLE. In SLE patients, binding of C1q to CRT on the surface of apoptotic neutrophils is disrupted by binding of PR3 to CRT, preventing macrophage recruitment and resulting in accumulation of apoptotic neutrophils. Delayed neutrophil apoptosis is also a trigger of autoimmune diseases, with RA being a typical example. Inhibition of neutrophil apoptosis can result in inflammation due to infiltration and activation of inflammatory cells. Factors such as high intracellular potassium concentration, activation of TNFR1 and GM-CSF receptor, and inhibition of potassium efflux can lead to inhibition of apoptosis and inflammation Delayed apoptosis of neutrophils and impaired clearance of NETs play similar roles in SLE on a macro level, both of which lead to the increase of anti-dsDNA antibodies and thus the occurrence of disease. This also reflects that although the mechanisms are different, different ways of dying of neutrophils may lead to the same outcome, and that there may be multiple ways of dying of neutrophils in the same autoimmune disease. And the same way of cell death will lead to different outcomes in different diseases through different pathways, perhaps this is also the significance of further research on the death of neutrophils.

NETs in AAV: mouth-watering prey

AAV is the general term for a class of multi-system autoimmune small vasculitis, including microscopic polyvasculitis (MPA), eosinophilic granuloma with polyvasculitis (EGPA) and GPA [68]. It is characterized by granuloma formation, arterioles and venules formation, and inflammation of capillaries. When blood vessels are in an inflammatory state, they may rupture or block causing systemic inflammation such as purpura rash and segmental glomerular infarction [69]. ANCA is mainly IgG autoantibodies against cytoplasmic components of neutrophils, which can bind to MPO and PR3 from neutrophils, and is considered to be a key marker to trigger the reaction of this disease [70]. Studies have shown that neutrophils and their interaction with ANCAs play a key role in the pathogenesis of AAV. Firstly, ANCAs activates neutrophils, which bind to antigens and attach to the antigen and Fcy receptors on neutrophils activated by proinflammatory cytokines to promote respiratory burst and degranulation of neutrophils, aggravating AAV [71, 72]. NETs are released by ANCA-stimulated neutrophils and contain targeted autoantigens PR3 and MPO. The deposition of NETs and MPO-DNA complex in the inflammatory kidney after ANCA binding with PR3, MPO and other autoantigens suggests that the formation of NETs induces vasculitis and promotes the autoimmune response in patients with vasculitis [73]. Fragments of NETs were also found in the microvascular system of patients with small vasculitis, suggesting that NETs are pathogenic in AAV [74]. In MRL/lpr mice, myeloid dendritic cells (mDCs) loaded with NETs-derived neutrophil-extracellular DNA led to the production of MPO-ANCA and PR3-ANCA with vasculitis-like kidney injury. The pathogenicity of NETs is also demonstrated by NETs-mediated damage to the vascular endothelium and its surrounding tissues [75]. Meanwhile, citrullinated histone H3 (Cit-H3) is also a key component of NETs, and the presence of Cit-H3 positive neutrophils is a specific marker for NETs formation in AAV [76]. Further study revealed that high mobility group protein (HMGB1) was strongly associated with the

NETs in AAV. HMBG1 increases ANCA translocation to regulate ANCA-mediated respiratory burst and neutrophil degranulation. In neutrophils treated with HMGB1 and ANCA positive IgG, HMBG1 can interact with TLR2 and TLR4 to mediate increased release of NETs in a NOX-dependent manner [77]. From the above description, it can be found that ANCA may promote the formation of NETs, which in turn aggravate the progression of AAV.

In fact, the specific mechanisms by which NETs facilitate the progress of AAV are not yet fully defined and comprehensive. It has been shown that ANCA-induced NETs activate alternative complement pathways to initiate inflammatory amplification cascades. Neutrophil cell membrane and neutrophil-derived particles can activate the alternative complement pathway, leading to C5a production. C5a can interact with C5a receptors of neutrophils to attract and activate neutrophils, forming an amplifying loop of ANCA activation of neutrophils [78]. At the same time, studies have shown that patients with AAV have a high incidence of venous thromboembolism events and are in a hypercoagulable state. C5a activates the coagulation system, and more importantly, both MPO-ANCA IgG and PR3-ANCA IgG stimulate C5A-modified neutrophils to produce NETs expressing tissue factor (TF), thus activating the coagulation system and aggravating AAV [79]. At the same time, multiple studies have shown that cellular immunoglobulin can affect NETs and thus aggravate the progression of AAV. For example, T cell immunoglobulin and mucin domain-containing protein 3 (TIM-3) have been reported to be important regulatory factors of T cells. Studies have found that the expression of TIM-3 in DCs of active MPO-AAV patients is significantly decreased, while the expression of TLR4 is significantly increased. Blocking TIM-3 can further enhance the expression of NETs mediated DC cytokines [80]. It has also been shown that platelets stimulated by the TLR pathway can induce the formation of NETs. TLR9-stimulated platelets release CXCL4 significantly in AAV and subsequently increase the formation of NETs. In addition, neutralizing anti-CXCl4 antibodies significantly inhibited AAV plateletenhanced NETs formation. TLR9 signal transduction and CXCL4 release underlie the critical role of platelets in NETs formation in the pathogenesis of AAV [81].

Therefore, if the role of NETs in SLE is the 'culprit', the role of NETs in AAV is a prey precisely targeted by autoantibodies. PR3, which constitutes NETs, binds to MPO and ANCA to activate neutrophilic degranulation and promote the production of NETs. At the same time, NETs directly cause vascular inflammation by destroying endothelial cells and activating the complement system, and indirectly act as a link between the innate and adaptive immune systems by producing PR3-ANCA and MPO-ANCA, creating a vicious cycle [82].

NETs in RA: danger maker

RA is an autoimmune disease and a chronic joint disease characterized by synovial inflammation, destruction of articular cartilage and production of anti-citrullinated peptide antibodies (ACPAs). The specific mechanism leading to the development of RA is not fully understood, but the formation of ACPAs is considered to be the key pathogenic factor. PAD2 and PAD4 may produce these citrulline antigens and are detected in the synovium of RA, which is closely associated with neutrophil infiltration [83]. Neutrophils as a member of innate immune cells infiltrate extensively in the joint cavity of patients with RA, in which neutrophils occupy a position that can not be ignored. NETs as an inflammatory mode of neutrophil death undoubtedly exacerbate the progression of RA. The histone citrullination catalyzed by PAD4 is a key step in NETs, in which citrullinated histones are neutralized.

It has been demonstrated that neutrophils externalize citrulline autoantigens associated with RA pathogenesis during the development of NETs which strongly induce the formation of NETs. In fact, NETs are in a positive feedback mechanism in RA. The inflammatory cytokines interleukin-17A (IL-17A) and TNF-ainduce the production of NETs in RA, and NETs induce the production of IL-6, IL-8, chemokines, and adhesion molecules, which aggravate the inflammatory response through the externalization of citrulline autoantigens and immune-stimulating molecules, which may promote abnormal joint and peripheral adaptations and innate immune responses and perpetuate the pathogenesis of the disease. In summary, the primary role of NETs in RA is that of the primary provider of autoantigens, in other words, NETs is a danger maker. Not only does PAD4 promote NETs participation in positive feedback regulatory mechanisms in RA, but other NETs-related proteases also show their skills. Neutrophils in blood were abnormally activated in RA, which increased the production of ROS and cytokines. At the same time, the expression levels of MPO, NE and lactoferrin related to NETs increased in synovial fluid (SF). These granule proteins can divide and activate cytokines and chemokines, who cause the degradation of cartilage and promote the production of RA [84]. Joanna Clarke et al. studied the direct impact of NETs on RA. They believe that NE can not only degrade cartilage, but stimulate the formation of downstream inflammation mediated by fibroblast like synovial cells (FLS), and promote the release of PAD2 in FLS. Autoantibodies such as anti-citrulline protein antibodies in RA will activate macrophages to release proinflammatory cytokines and activate neutrophils in synovium, so as to release more NE and aggravate joints inflammation. The NE of NETs aggravates the joint inflammation, and the joint inflammation in turn promotes the production of NE, the two promote and influence each other [85].

In recent years, researchers have made a detailed analysis of the types and proteomics of NETs in RA, which provides a basis for the pathological diagnosis of RA. Elinor A. Chapman and others found that ROS required for the production of NETs come from two ways, one from the activation of NOX2 and the other dependent on mitochondria. The occurrence of RA is NOX2 independent NETs. At the same time, it was found that many NETs proteins increased in RA synovial fluid, including cathepsin, MPO, MMP8, MMP9, Lcn2 and PADI2. These proteins contribute to the cellular infiltration and degradation of collagen in synovial joint. It provides a basis for predicting the severity and cure of RA patients [86].

NETs emerge as multi-faceted adversaries: unleashing a cascade of autoimmune disorders

Focusing on the skin, psoriasis is a chronic systemic inflammatory disease, which affects not only the skin, but various internal organs. One of the important signs of psoriasis is the infiltration of epidermal neutrophils, which will lead to the occurrence of NETs [87]. There have been many studies on the occurrence and pro-inflammatory mechanism of NETs in psoriasis. Neutrophils amplify skin inflammation by releasing NETs and activating IL-36 and TLR4 signals, and further increase the infiltration of neutrophils into the skin [88]. Other studies have shown that neutrophils and NETs can promote inflammation through several different mechanisms, such as the activation of inflammasome, triggering TLR7 and TLR9 through self-antigen complexes (such as LL-37-DNA), promoting macrophage apoptosis, or processing and activating IL-36 cytokines. At the same time, NETs also contain RNA. In previous studies, DNA research in NETs is more extensive. However, it is ignored that RNA is also a rich part of mammalian NETs. The RNA associated with NETs may be the effective component of RNA-LL37 and PMA triggering NETs [89]. In addition, for Stevens Johnson Syndrome (SJS) and toxic epidermal necrolysis (TEN), Manao Kinoshita et al. found that CD8⁺ T cells that trigger skin infiltration in the early stage of inflammation produce lipocalin-2, which triggers the formation of NETs in the early damaged skin, and then releases LL-37, which induces keratinocytes to express formyl peptide receptor 1 (FPR1), resulting in cell necrosis [90].

With a specific focus on the digestive system, IBD represents an autoimmune condition characterized by inflammation and damage to the gastrointestinal tract, where NETs also play a role. A recent investigation demonstrated elevated levels of NETs in both the plasma and colon tissue of individuals diagnosed with IBD, thereby indicating the potential involvement of NETs in the onset and progression of this disease. Notably, IBD is associated with an increased risk of thrombosis and heightened hypercoagulability. To mitigate disease progression, the administration of DNase has been shown to impede the formation of NETs, reduce the release of thrombogenic cytokine mediators by macrophages, and directly inhibit platelet activation [91]. Histones, constituents of NETs, possess the ability to directly impair the permeability and structural integrity of the intestinal epithelium. Experimental evidence using monolayers of Caco-2 intestinal cells demonstrates that PAD4 mediates the disruption of epithelial permeability by causing the degradation of zonula occludens protein 1 (ZO-1) and adhesion junction protein E-cadherin. This process ultimately triggers apoptosis and contributes to the progression of IBD. Nevertheless, the precise underlying mechanism of this phenomenon remains to be fully elucidated [92]. In the context of UC and CD, the involvement and specific functionality of NETs exhibit subtle distinctions. Investigations have revealed significantly elevated expression levels of NETsassociated proteins, including Cit-H3, PAD4, MPO, and NE in the colonic tissue of UC patients when compared to both healthy individuals and CD patients. Notably, the production of NETs in UC patients is stimulated by the presence of TNF- α that activates circulating neutrophils. Moreover, the activation of extracellular signal-regulated kinase 1/2 (ERK1/2) in lamina propria mononuclear cells (LPMC) in UC patients ultimately amplifies the production of TNF- α and IL-1 β , thereby exacerbating the disease process [93]. The use of PAD4 inhibitors can reduce the production of NETs and improve DSS induced acute enteritis [94]. Intriguingly, emerging evidence suggests that achieving a balanced state rather than complete elimination of NETs may be beneficial in patients with UC. Rectal bleeding has long been recognized as a prominent clinical manifestation in UC patients; however, the formation of fibrin and neutrophil-generated immune clots provides a protective mechanism by stabilizing the compromised mucosal tissue and effectively halting bleeding. In this intricate process, PAD4 facilitates the condensation of neutrophil chromatin, whereby the outer chromatin of NETs impedes blood vessel patency and further aggregates to fulfill a hemostatic function. Hence, NETs play a vital protective role in the context of UC-related bleeding [95]. While numerous proteins associated with NETs are more commonly observed in UC than in CD, there is evidence indicating a specific correlation between NETs and CD. Essential constituents of NETs, including Cit-H3, NE, and MPO, can indeed be detected in stool samples from CD patients, suggesting their potential involvement in the pathogenesis of CD. Furthermore, additional studies have demonstrated enhanced expression of Cit-H3 and PAD4 in mice with 2,4,6-trinitrobenzene sulfonic acid (TNBS)-induced colitis, further supporting the role of NETs in CD pathogenesis. Notably, the administration of PAD4 inhibitor Cl-amidine effectively mitigated the clinical colitis index and reduced the expression of pro-inflammatory factors in mice, indicating its therapeutic potential in CD [96]. The production of NETs is positively correlated with the severity of CD, which may be caused by the excessive production of bactericidal oxidant hypochlorous acid (HOCl) in the extracellular region mediated by MPO, which leads to host tissue damage and intensification of CD. Perianal fistulising Crohn's disease (pfCD) is a disabling phenotype of CD. Studies have shown the presence of NETs in perianal fistulas, increased production of NETs is associated with enhanced TNF- α production and adverse fistula remission [97]. In conclusion, the excessive generation of NETs worsens the onset and progression of CD and hampers its recovery. Furthermore, while MPO and NE have been identified as serum biomarkers for diagnosis and prognosis in patients with CD and UC, ANCAs targeting PR3 are more commonly detected in UC compared to CD, making them a reliable means of distinguishing between the two diseases.

Regarding the nervous system, emerging evidence indicates the involvement of neutrophils in the pathogenesis of MS. However, unlike in other diseases, neutrophils do not prominently appear in central nervous system tissue biopsies of MS patients. This discrepancy may be attributed to the transient nature of neutrophils as short-lived cells, which primarily respond to specific disease stages such as disease onset or relapse [98]. Due to this characteristic, the investigation of NETs in MS remains inadequate. Nevertheless, several studies have identified correlations between neutrophil-associated factors, such as NE, CXCL1, and CXCL5 in plasma, and the severity of the disease. Additionally, elevated levels of MPO have been significantly observed in the serum of individuals with MS [99]. Moreover, studies have shown that the serum NETs level in relapsing-remitting (RR)-MS (RRMS) patients is higher than that in other patients with MS pressure type, and the frequency in males belonging to this subtype is significantly higher than that in females, suggesting that NETs may be related to some sex differences in this disease. At the same time, NETs may affect the process of RRMS through T cell immune regulation [100].

Irrespective of individual organs and systems, the investigation of systemic autoimmune diseases has increasingly focused on the physiological activity and impact of NETs in SSC. Apart from RA and SLE, substantial levels of anti-NET antibodies (ANETA) and fragments of NETs have also been identified in the serum of SSC patients. Notably, a gradual reduction in NETs formation was observed during the early stages of the disease, whereas patients with severe and chronic vascular complications exhibited persistently elevated levels of NETs throughout the course of the disease. Additionally, DNA filaments have been detected in skin biopsies of SSC patients, potentially representing NETs. These NETs are thought to undergo modifications by CXCL4, leading to the formation of CXCL4-DNA complexes that contribute to immune activation and the production of IFN-I in individuals with early active SSC [101]. The interaction between platelets and neutrophils plays a crucial role in SSC, enhancing the ability of white blood cells to extravasate from blood vessels and migrate to inflamed tissues, while also promoting the generation of NETs. When phagocytosis is impaired, activated platelets evade clearance mechanisms, leading to the buildup of microparticles (µPs), triggering NETs formation and contributing to vascular damage. The substantial decrease in P-selectin (PSGL-1) expression observed in SSC increases the likelihood of phagocytes being unable to recognize and engulf platelets, thus resulting in sustained µPs release. Therefore, the elimination of activated platelets via PSGL-1 may serve to restrict µPs accumulation and NETs production, ultimately reducing the progression of the disease [102].

The impact of NETs on APS is also considered impossible to ignore. APS is an autoimmune thromboinflammatory disease, which is characterized by the formation of arteriovenous or microvascular thrombosis and poor pregnancy. There are a variety of autoantibodies, such as antiphospholipid antibody (aPL), an-ticardiolipin antibody (aCL). APL makes the normal steady-state mechanism on the surface of neutrophils destroyed and directly triggers the release of NETs [103]. High levels of anti-NETs IgG can damage the clearance of NETs, which is also closely related to the formation of venous thrombosis [104]. Pregnancy complications are also the main clinical manifestation when APS occurs. Yuan Lu et al. found that the markers of NETs increased in pregnant women with APS, and NETs may lead to placental defects. This direction may provide a new target for the treatment of the disease [105].

Other autoimmune diseases, such as autoimmune diseases in the kidney, can also see the accumulation of NETs. At the same time, studies have shown that NETs can lead to a vicious circle [106]. It can be seen that NETs are closely related to autoimmune diseases, and the treatment of autoimmune diseases for NETs is worthy of in-depth study (Fig. 2).

Neutrophil necroptosis: another potential mechanism contributing to autoimmune pathogenesis

Neutrophil necroptosis is a form of regulated necroptosis that is triggered by RIPK3-MLKL-dependent necroptosis when cells are stimulated by TNF, adhesion receptors, monosodium urate (MSU) crystals or staphylococcus aureus [107]. Necroptosis is similar to other forms of death such as NETs, which are capable of causing inflammation and immune damage. RIPK3-MLKL signalling can work with p38 MAPK and RIPK3-MLKL signalling can activate NOX together with p38 MAPK and PI3K, resulting in ROS production and non-apoptotic cell death [107]. Neutrophil necroptosis has been implicated as a mode of death that triggers inflammation because this mode of cell death results in the release of toxic cellular contents, namely DAMP, which cause tissue damage and induce an inflammatory response [107, 108]. Recently, neutrophil necroptosis has been found in autoimmune diseases such as psoriasis and RA [109]. This suggests that the study of neutrophil necroptosis may provide new ideas for the treatment of autoimmune diseases. The following section will focus on the contribution of neutrophil necroptosis to inflammation in patients with RA.

Neutrophil necroptosis triggers RA through different pathways

Neutrophil necroptosis is essentially a form of programmed death mediated by RIPK1, RIPK3, and MLKL that involves the release of intracellular toxic contents that mediate the onset of inflammation in RA [110].

In RA patients, neutrophil exposure to GM-CSF activates RIPK1, RIPK3, and MLKL results in necroptosis [111], followed by attachment of adhesion receptors [112]. It was found that CD44 attached to GM-CSF-stimulated neutrophils induced necroptosis, and interestingly, CD44 attached to neutrophils isolated from the synovial fluid of RA patients in the absence of GM-CSF stimulation also triggered necroptosis, which could indicate that CD44 can trigger necroptosis in neutrophils in the presence of inflammation. granulocyte necroptosis under inflammatory conditions [113].

Fibroblast-activated protein-a (FAP-a) also plays an important role in the development of necroptosis in neutrophils from RA patients [114]. Neutrophils isolated in the joints of RA patients undergo GM-CSF non-dependent necroptosis after CD44 clearance and this effect can be blocked by the FAP-a inhibitor, MLKL. It was found that FAP-a was activated in parallel with RIPK3-MLKKL-p38 MAPK signalling and that its activity was required for CD44 to trigger ROS production and subsequent cellular necroptosis. Thus, FAP-a is potentially a key target for the inhibition of neutrophil necroptosis in RA [108].

Indeed, in addition to RA, neutrophil necroptosis has been identified in patients with a variety of autoimmune diseases, and because neutrophil adhesion molecules can trigger necrosis under inflammatory conditions, vacuolisation of the neutrophil cytoplasm has been found in inflamed tissue from patients with psoriasis [112]. In AAV, ANCA induces necroptosis in neutrophils, and necroptosis inhibitors can also prevent AAV altogether [115]. Currently, RIPK1 inhibitors are in phase II clinical trials and are being investigated for the treatment of autoimmune diseases such as psoriasis, UC and RA [116, 117].

Neutrophil necroptosis and NETs: intertwined patterns of neutrophil death

Though necroptosis and NETs are non-apoptotic deaths which lead to the release of toxic contents from neutrophils into the extracellular space causing damage to surrounding tissues, the mechanisms of necroptosis and NETs are not the same. However, there appears to be an intrinsic link between them.

Studies have shown that PMA or MSU crystals can trigger the production of NETs in the presence of neutrophil necroptosis, while the absence or inhibition of necroptosis by PIPK3 also inhibits the induction of NETs by MSU crystals [118, 119], and that PIPK3-dependent necroptosis is an upstream activator of NETs. However, necroptosis does not inhibit all NETs, it may be able to control the formation of NETs induced by MSU crystals, but it remains controversial whether there is a role for NETs formed by PMA induction [115]. Outside of this phenomenon, however, NETs have been identified through studies independent of RIPK3 and MLKL, and one possible explanation is that the so-called NETs that occur during necroptosis are actually passive releases of chromatin associated with necrosis, whereas the formation of true NETs is not dependent on the occurrence of necroptosis [107]. It is undeniable that there is an underlying link between NETs and necroptosis, and it is the intertwining and balancing of these death pathways that determines the ultimate fate of neutrophils. Perhaps a deeper understanding of the love-hate relationship between neutrophil death modalities may also provide a novel way to explore the mechanisms of autoimmune disease development.

Neutrophil pyroptosis: a novel pathological mechanism in autoimmune disorders

Pyroptosis is a programmed death process associated with inflammation and has recently been extensively studied by researchers. Cells undergoing pyroptosis form pyroptosis

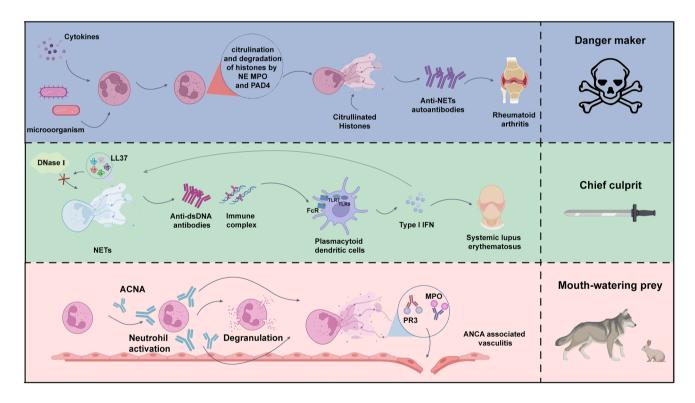


Fig. 2 The role of NETs in various autoimmune diseases. NETs are closely related to a variety of autoimmune diseases, but the *role* they play in the pathogenesis of various autoimmune diseases is variable. In SLE, immune complexes such as LL37 and HNP protect the DNA in NETs from nuclease degradation, which induces anti-dsDNA autoantibodies and releases I IFN with pDC to trigger SLE. In AAV, NETs are released by ANCA-stimulated neutrophils and contain targeted autoantigens PR3 and MPO. After ANCA binds to autoantigens

such as PR3 and MPO, NETs and MPO-DNA complexes are deposited in inflammatory sites and promote the autoimmune response in patients with vasculitis. The formation of NETs is catalyzed by PADs, in which the arginine residues of the protein are deaminated to citrullinated proteins. Therefore, the production of NETs promotes the production of ACPAs and leads to the production of inflammatory factors and adhesion proteins in RA. If NETs are the chief culprit in SLE, NETs are prey and danger maker in AAV and RA vesicles similar in size to apoptotic vesicles and are accompanied by chromatin sequestration and damage and the formation of numerous voids in the cell membrane, ultimately causing lysis of the cell membrane and the release of intracellular contents leading to inflammation [13, 120]. NLRP3 inflammasome, caspase-1, GSDMD, IL-1 β and IL-18 play key roles in the development of pyroptosis [121]. NLRP3 inflammasome are sensors that activate caspase-1, which can be activated by factors such as ROS, and when caspase-1 is activated by inflammatory vesicles, the caspase-1 then cleaves pro-IL-1 β and pro-IL-18 to produce IL-1 β and IL-18, which in turn cause inflammation to occur [122, 123].

Pyroptosis was firstly detected in macrophages and related diseases, followed by numerous reports confirming that neutrophils can also undergo pyroptosis, which is closely related to the immunity of body. In the early stages of sepsis, neutrophil pyroptosis can act as an amplifier of inflammatory signals and thus recruit immune cells, which may have a positive impact on the treatment of sepsis. During infection, neutrophil pyroptosis is particularly important for bacterial clearance [120]. In but some studies have indicated that neutrophils have a specific resistance to pyroptosis and that the pathway for IL-1ß production is somewhat different from the classical pyroptosis pathway. While macrophages, for example, undergo pyroptosis mainly by rupture of the plasma membrane (PM) via GSDMD, the cleavage of N-GSDMD during NLRP3 inflammasome signalling in neutrophils is mainly associated with abundant intracellular organelle membranes. In neutrophils activated by NLRP3, instead of being transported to the plasma membrane, N-GSDMD protein is transported to azurophilic granules and autophagosomes, which ultimately release IL-1 β via an autophagy-dependent pathway [124]. Therefore, the mechanism of pyroptosis in neutrophils is not clear, which provides an important direction for research. Instead, it has recently become clear that neutrophils can undergo this form of death when lacking the functional NADPH oxidase Nox2 [121, 123]. However, recent studies have found that methicillin-resistant staphylococcus aureus (MRSA) uses neutrophil necroptosis with pyroptosis to trigger IL-1ß production. Neutrophil pyroptosis may not act alone [13].

The role of inflammasome activation in the pathogenesis of rheumatic autoimmune diseases

A variety of typical inflammatory vesicles such as NLRP1, NLRP3, AIM2 and NLRC4 have been identified in a variety of rheumatic autoimmune diseases. In RA, NLRP3 inflammasome mediate the activation of IL-1β production, which

leads to inflammation [125]. In SLE, anti-U1-snRNP autoantibodies are able to activate NLRP3 inflammasome in human monocytes, leading to IL-1 β production [126, 127]. In macrophages from SLE patients, NETs-mediated activation of NLRP3 inflammasomes is significantly enhanced [128]. Also, in ankylosing spondylitis (AS), IL-1 β production was highly induced, suggesting that inflammatory vesicles may be activated and induce IL-1 β production in AS [129]. It is not difficult to speculate on the important role of pyroptosis in autoimmune disease in these studies[121], it is also worthwhile to explore in depth the role of pyroptosis as a neutrophil critical for inflammation in the development of autoimmune disease.

Pyroptosis-related elements in other modes of neutrophil death

It has been found that GSDMD, NLRP3, which plays an important role in pyroptosis, also contributes to inflammation and autoimmune diseases by participating in, or being activated by, multiple modes of neutrophil death [125, 130]. In adult Still's disease (AOSD), DNA released by NETs activates NLRP3 inflammasome and increases the expression levels of IL-1 β and IL-18 [131].

GSDMD brings together two death pathways, pyroptosis and NETs, and GSDMD is cleaved by NE to drive NETs [124, 132]. the formation of NETs requires GSDMD, while in pyroptosis GSDMD is cleaved, resulting in the formation of membrane pores [133]. In summary, GSDMD is the proinflammatory cell pivotal to death. Also, GSDMD is thought to promote neutrophil death and GSDMD deficiency may enhance host defence by delaying neutrophil death [134]. In fact, neutrophil pyroptosis has been associated with autoimmune diseases. Clinical experiments showed that the expressions of NLRP3 and apoptosis associated speck-like protein containing a CARD (ASC) and pro-caspase-1 in neutrophils of RA patients were significantly decreased. The expression of activated caspase-1 protein was significantly increased. Overactivation of caspase-1 in RA neutrophils may mediate IL-18 activation, thereby contributing to RA progression in an NLRP3 inflammasome-independent manner [135].

Currently, death modalities such as pyroptosis and necroptosis are considered to be inflammation-associated death modalities. Perhaps, by inhibiting these inflammationrelated death modalities or converting these death pathways to non-inflammatory death, such as apoptosis, may also be a strategy to treat autoimmune diseases. It is therefore particularly important to explore the mechanisms that trigger the various modes of death of neutrophils in autoimmune diseases.

Neutrophil ferroptosis in autoimmune diseases: a new type of neutrophil death mode has emerged

Ferroptosis is a recently discovered form of non-apoptotic cell death that causes disease through iron-catalyzed oxidation of unsaturated fatty acids and thus the accumulation of ROS [136]. Several studies have found a close association between ferroptosis and the development of tumors and other degenerative diseases [137-141], as well as in autoimmune diseases [142]. Abnormalities in cellular ferroptosis have been seen in autoimmune-mediated hepatitis (AIH), but the exact mechanism of occurrence remains unclear. In the GPX4[±] mouse high glucose and high fat diet model, INOS is significantly more upregulated than in WT mice [143], which indirectly suggests that ferroptosis promotes the production of INOS and thus contributes to AIH [144]. Further, neutrophil ferroptosis is also closely associated with autoimmune diseases. IFN-a and SLE IgG inhibit the transcription of GPX4 by promoting the binding of CREM to the GPX4 promoter, thereby increasing the production of lipid peroxides and inducing the development of neutrophil ferroptosis, thus contributing to the development of SLE. the development of SLE. This places neutrophil ferroptosis at the centre of SLE pathogenesis and highlights the importance of targeting GPX4 transcription and neutrophil ferroptosis in the treatment of SLE [145, 146].

At present, research on ferroptosis in autoimmune diseases is uncommon and in-depth, and the role of neutrophil ferroptosis in autoimmune diseases is not yet clear. However, from the above-mentioned article on neutrophil ferroptosis in SLE published in Nature, it appears that this newly discovered mode of death of neutrophils, which play an important immune role in autoimmune diseases, may indeed exist. This clearly provides us with new research and therapeutic ideas, and the investigation of neutrophil death patterns in autoimmune diseases continues.

Autophagy: the entanglement between survival and death mechanisms

In addition to various forms of cell death, investigating the normal physiological function and survival mecihansms of neutrophils is crucial for understanding neutrophil death. Autophagy is a pivotal mechanism that plays a role in multiple aspects of neutrophil biology and pathophysiology. Not only does it contribute to cellular homeostasis, but its interplay with different modes of neutrophil death also influences neutrophil survival and its potential association with autoimmune diseases.

Autophagy is a process in which eukaryotic cells use lysosomes to degrade cytoplasmic proteins and damaged organelles under the regulation of autophagy related gene (Atg) [13]. Autophagy exhibits significant upregulation in response to oxidative stress, nutritional deficiencies, and diverse antigenic stimuli, including pathogens, in order to ensure cellular survival. While autophagy provides energy and supports cell survival, dysregulated or uncontrolled autophagy can also result in autophagy-dependent cell death, leading to pathological conditions. In summary, autophagy plays a critical role in cellular survival [147]. Autophagy is a mechanism of survival whose function is momentous in the birth and death of neutrophils. It has been found that autophagy can dynamically regulate different stages of neutrophil differentiation in vivo. The accumulation of immature neutrophils in mice with conditionally knocked out Atg7 in hematopoietic stem cells (HSCs) and progenitor cells demonstrated that inhibition of autophagy delayed neutrophilic differentiation [148]. At the same time, in the Atg7 deficient neutrophil precursors, the glycolytic activity increased, but the mitochondrial function impaired, ATP production decreased, and lipid droplets accumulate, suggesting that free fatty acids provided by autophagy are particularly important for the differentiation of neutrophil [149]. However, autophagy can only regulate the number of neutrophils at the production level, but not by regulating the apoptosis of mature neutrophils.

In addition to the growth and development of neutrophils, autophagy is also essential for major neutrophil functions, including degranulation, reactive oxygen species production and release of neutrophil extracellular traps. Degranulation is a process in which neutrophils regulate the secretion of granulocytes, and it is one of the main mechanisms of neutrophilic inflammatory response. It was found that mice with myeloid specific autophagy deficiency showed reduced degranulation both in vivo and in vitro, and showed reduced severity of neutrophil-mediated inflammation and autoimmune diseases. At the same time, NADPH oxidationmediated ROS production is also reduced in autophagy deficient neutrophils, and inhibition of NADPH oxidase reduces neutrophils degranulation, which not only suggests that autophagy can control neutrophils degranulation, but also proves that NADPH oxidase is the intersection of autophagy and degranulation [150].

In addition to its role in neutrophil degranulation, autophagy exhibits a profound interplay with NETs, forming a complex relationship that contributes to the development of various autoimmune diseases, acting as a critical tipping point. Notably, ROS, serving as a trigger for NETs formation, displays an intricate and reciprocal dependency on autophagy. ROS induction can stimulate autophagy, while autophagy, in turn, becomes a crucial prerequisite for ROS generation. Consequently, inhibition of either autophagy or NADPH oxidase effectively hampers chromatin condensation, ultimately impeding the formation of NETs [147]. The interconnection between autophagy and NETs is vividly exemplified by their convergence. It has been observed that activated platelets exhibit a robust capacity to transfer potent HMGB1 molecules to neutrophils, thereby acting as formidable inducers of NETs. Moreover, this potent NETs inducer simultaneously triggers the activation of autophagy, further highlighting the intricate relationship between these two processes [151]. This evidence not only proves that platelets can affect the formation of NETs, but also reveals the inevitable relationship between NETs and autophagy. Interestingly, heparin as a conventional anticoagulant and antithrombotic agent interferes with the interaction between neutrophils and platelets, thereby inhibiting their eventual activation. low molecular weight heparins (LMWH) in prophylactic dose can inhibit the autophagy of peripheral neutrophils and NETs produced under inflammatory stimulation [152]. Meanwhile, chloroquine (CQ), a traditional treatment for malaria, is also a classic autophagy inhibitor, which can impair autophagosome lysosome fusion and degradation of autophagosome contents. It was found that CO significantly reduced serum free DNA and Cit-H3 in mice with pancreatitis as well as significantly inhibited the formation of NETs [153]. Autophagy inhibitor 5'-(4-fluorosulfonylbenzoyl) and knockout of Atg7 inhibit LC3 autophagy formation and also significantly reduce the production of NETs in promyelocytes [154]. The interplay between autophagy and NETs offers novel therapeutic prospects for addressing inflammation and autoimmune diseases, thereby underscoring the potential involvement of autophagy in other forms of neutrophil death, thereby showcasing its impact on autoimmune diseases. Initially, autophagy, serving as a survival mechanism, is inherently linked with apoptosis, recognized as an essential cellular pathway. Polymorphonuclear neutrophils (PMN), although short-lived, exhibit evidence of prolonged survival of Staphylococcus aureus within their confines, thereby facilitating pathogen persistence during infection. This survival mechanism employed by the bacterium within PMN primarily relies on the inhibition of neutrophil apoptosis through autophagy promotion and the upregulation of anti-apoptotic genes (mcl-1, bcl-2). Conversely, restoration of normal apoptosis levels in infected PMN is achieved through the inhibition of autophagy. Thus, a compelling association between autophagy and apoptosis in PMN undoubtedly exists [155]. At the same time, autophagy inhibitors 3-methyl-adenine (MA) and CQ can significantly delay TNF- α -induced neutrophils apoptosis and significantly down-regulate the expression of mcl-1 [156]. The relationship between granulocyte autophagy and pyroptosis seems to be more interesting and special. In fact, the pyroptosis of neutrophils is different from that of canonical and non-canonical macrophages, and autophagy is involved in the pyroptosis of neutrophils. In neutrophils, N-GSDMD does not accumulate in large quantities on the plasma membrane, but is transported to azurophilic (primary) granules and autophagosomes. At the same time, it was found that the secretion of IL-1 β was impaired in Atg7^{-/-} mouse neutrophils, and the inhibition of autophagy by VPS34-IN1 and the blocking of autophagy by BAFA1 significantly reduced the release of IL-1 β , suggesting that neutrophils release IL-1 β through an autophagy dependent pathway [124]. This process may be that IL-1 β enters lumen of a vesicle and then becomes autophagosome, and finally autophagosome containing IL-1 β can directly fuse with the plasma membrane and be secreted into the extracellular [157].

The primary objective of investigating the correlation between autophagy and neutrophil death lies in the quest for novel targets and therapeutic interventions for various diseases. The intricate interplay between autophagy and alternative forms of neutrophil death has been implicated in the pathogenesis of numerous autoimmune disorders. Notably, investigations have revealed elevated levels of autophagy in neutrophils of SLE patients. Among the pivotal players in this context, the REDD1 protein assumes a critical role in maintaining energy homeostasis and orchestrating multiple inflammatory pathways. Furthermore, endothelin-1 (ET-1) and hypoxia-inducible factor-1 α (HIF-1 α) act as inducers of autophagy, culminating in the release of NETs through the upregulation of REDD1, thus serving as inflammatory triggers that exacerbate the progression of SLE [158]. Similarly, REDD1 was highly expressed in intestinal neutrophils of UC patients and IL-1 β and tissue factor (TF) modified NETs were formed in an autophagy dependent manner [159]. In AAV, the lysosomal membrane protein-2 (LAMP-2) ANCA-induced NET formation is involved in initiation of autophagy mechanism. Anti-lamp-2 antibody can inhibit neutrophil spontaneous apoptosis, activate neutrophil autophagy and activate NETs. Apoptosis and necrosis inhibitors did not interfere with the formation of NETs induced by anti-LAMP-2 antibodies [160]. In SSC, activated platelets exhibit a robust release of SSC particles and HMGB1, thereby inducing neutrophils to undergo autophagy. Subsequently, the redistribution of vesicle contents triggers the generation of NETs, consequently exacerbating vascular pathology and tissue fibrosis associated with the disease [161]. Furthermore, the IL-23/17 axis plays a pivotal role not only in T cell differentiation but also in modulating neutrophil homeostasis and migration in psoriasis. Researches demonstrate a reduction in the activities of NETs, NE, and ROS in Wdfy3-deficient neutrophils, with Wdfy3 acting as a key regulator of macroautophagy. Hence, the interplay between autophagy and other forms of neutrophil death, including NETs, warrants significant attention in the context of autoimmune diseases [162].

Conclusion

Neutrophil death is a complex and multifaceted phenomenon that has significant implications for the immune system. While the normal non-lytic morphology of neutrophils is essential for maintaining the body homeostasis, the emergence of abnormal non-lytic morphology and lytic morphology that causes inflammation can lead to the development of autoimmune diseases that pose troublesome problems for the immune system. Unlike other cells, neutrophils exhibit unique mechanisms of death that involve the production of reactive oxygen and nitrogen species. These species play a critical role in the development of infections and autoimmune diseases, highlighting the importance of understanding the various modes of neutrophil death.

Furthermore, it is not uncommon for multiple modes of neutrophil death to be present in a given autoimmune disease, and these modes of death often interact or overlap with each other. This interplay between different modes of neutrophil death underscores the pivotal role played by these cells in the immune system and highlights the potential importance of their death in the development of autoimmune diseases. As such, continued research into the various modes of neutrophil death may provide new insights into the treatment of autoimmune diseases, and this area of study represents an exciting frontier for pharmacological research.

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Declarations

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