



Review article

NETosis – Dr. Jekyll and Mr. Hyde in inflammation

Lucia Lauková*, Barbora Konečná

Comenius University in Bratislava, Faculty of Medicine, Institute of Molecular Biomedicine, Bratislava, Slovakia



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ABSTRACT

Neutrophils play an important role as the central mediators of the innate immune defence response, providing the first line of host protection. It was shown that these cells can trap and kill various microorganisms through different ways. One of them is a release of neutrophil extracellular traps (NETs) composed of chromatin fibrils and antimicrobial proteins. There is the evidence that the release of NETs does not have only a beneficial effect. NETs can trap and kill microorganisms and pathogens, however on the other hand the same pathway can also cause the damage of the organism by various mechanisms. NETs participate in the pathogenesis of a lot of inflammatory and autoimmune disorders, such as thrombosis, atherosclerosis, cystic fibrosis, periodontitis, lupus, rheumatoid arthritis and others. The aim of this review is to summarize information about the release of NETs and their beneficial, but also detrimental effect during various diseases. The better characterization and understanding of the dual role of NETosis during these diseases is necessary for the early diagnosis and more effective treatment.

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Introduction

NETosis is a specific form of cell death performed by neutrophils which is characterized by release of chromatin fibrils into the extracellular space referred to as neutrophil extracellular traps (NETs) (Yang et al., 2016). This process was firstly described by Brinkmann et al. (2004). Since then, mechanism and function of NETosis have been intensively studied.

Neutrophils play a pivotal role in inflammatory response. Variety of neutrophil cell death including apoptosis, necrosis, autophagic cell death and NETosis coexist in response to inflammation. Distribution of cell death phenotypes of neutrophils may be determined by specific factors expressed by individual pathogens, the degree of insult, the duration of stimulation, and the response of the host immune system (Iba et al., 2013). NETosis as well as other types of cell death are mechanisms releasing DNA to the circulation. However, NETosis has several specificities. While apoptotic cells display eat-me signals, such as translocation of phosphatidylserine to the extracellular monolayer of plasma membrane, cells after NETosis do not, which prevents them from clearance by phagocytes. In contrast to apoptosis or necrosis,

granular and nuclear membranes demerge during NETosis, but the integrity of plasma membrane can be retained (Luo and Loison, 2008).

Nowadays, three types of NETosis are known. Suicidal NETosis is the well characterized and most often observed type. This process is oxidant-dependent and includes the cell membrane rupture and the development of cell lysis, which indicates neutrophil death (Kaplan and Radic, 2012). In contrast, in vital NETosis neutrophils release DNA without a loss of nuclear and plasma membrane. That means that after the vital NETosis neutrophils stay alive and are able to phagocytose (Clark et al., 2007). Vital NETosis is divided into the two groups dependently on oxidant formation and origin of the DNA in NETs (Yousefi et al., 2009) (Table 1).

NETs are formed in response to a variety of pro-inflammatory stimuli as well as to several microorganisms in order to trap and kill them (Brinkmann et al., 2004). Today the relationship between NETosis and inflammation is not fully understood. On the one hand, NETosis is the mechanism of immune response and can help to reduce the number of pathogens. On the other hand, NETosis can be connected with inflammatory pathway as an inductor. This dual character can be explained by the fact that NETs are composed of DNA and different types of proteins, which all can induce immune response by their own way (Kaplan and Radic, 2012). The limit, when NETosis is changed from beneficial to detrimental is unknown but NETs showed to be involved in pathogenesis of various inflammatory diseases.

* Author for correspondence: Lucia Lauková, Comenius University in Bratislava, Faculty of Medicine, Institute of Molecular Biomedicine, Sasinkova 4, 811 08 Bratislava, Slovak Republic.

E-mail address: luci.laukova@gmail.com (L. Lauková).

Table 1
Characterization of different types of NETosis.

| Type of NETosis | Suicidal | Vital |
|-----------------------------|-------------|----------|
| Duration | 120–240 min | 5–60 min |
| Source of DNA | nucleus | nucleus |
| ROS-dependent | yes | no |
| Rupture of plasma membrane | yes | no |
| Rupture of nuclear membrane | yes | yes |

ROS=reactive oxygen species.

Stimulation and inhibition of NETosis

Formation of NETs represents the response to different proinflammatory impulses, such as lipopolysaccharide (LPS), interleukine-8 and tumour necrosis factor α (TNF- α), as well as to various types of pathogens (Table 2). Beside these, NETosis can be stimulated by activated platelets and endothelial cells, nitric oxide, monosodium urate crystals and various autoantibodies or even chemical compounds such as phorbol-12-myristate-13-acetate (PMA) (Kaplan and Radic, 2012).

A physiological inhibitor of NETosis has not been known for a long time. Recent study showed that the blockade of the prostaglandin E2 signalling pathway inhibits NETosis (Domingo-Gonzalez et al., 2016) through the production of cyclic adenosine monophosphate (Shishikura et al., 2016). Modulation of NETosis

could be possible also by blocking of inducers or other members of the pathway (Table 2). These findings can contribute to the development of novel treatments for NETosis-related diseases.

NETs structure

Neutrophils have a nucleus consisting of two to five lobes joined together by hair-like filaments. Neutrophils contain granules in cytoplasm, which are classified into three groups based on the presence of proteins. Primary granules comprise mainly the myeloperoxidase (MPO), while secondary granules comprise the lactoferrin, and tertiary gelatinase. They also contain a number of proteins that are important in the initiation of the immune response, such as α -defensins, cathelicidin and others (Borregaard, 2010). NETs are formed by chromatin filaments composed of DNA and histones (Urban et al., 2009). Moreover, the proteins from primary, secondary and tertiary granules of neutrophils are also part of NETs. The granular NET-associated proteins are primarily cationic; therefore, they can effectively bind to DNA molecule. They include: neutrophil elastase (NE), MPO, cathepsin G, proteinase 3, bactericidal/permeability-increasing protein, calgranulin, α -defensins, lactoferrin, a fragment of the protein cathelicidin hCAP18—the peptide LL-37, pentraxin, matrix metalloproteinase-9 (MMP-9) and peptidoglycan recognition protein-S (Table 3) (Averhoff et al., 2008; Bianchi et al., 2011; Parker et al., 2012).

Table 2
Inductors and inhibitors of NETosis.

| Inductors | Article |
|-----------------------------------|--|
| Gram-positive bacteria | Malachowa et al. (2013), Ramos-Kichik et al. (2009) |
| Gram-negative bacteria | Juneau et al. (2011), Marín-Esteban et al. (2012) |
| Virus | Narasaraju et al. (2011) |
| Protozoa | Abi Abdallah et al. (2012), Ventura-Juarez et al. (2016) |
| Yeast | Byrd et al. (2013) |
| PMA | Keshari et al. (2013) |
| LPS | Brinkmann et al. (2004) |
| Cytokines | Alfaró et al. (2016), de Boer et al. (2013) |
| Chemokines | Rossaint et al. (2014) |
| Immune-complexes | Behnen et al. (2014) |
| Anti-neutrophil antibodies | Sha et al. (2016) |
| Activated platelets | Clark et al. (2007) |
| Uric acid | Arai et al. (2014) |
| Monosodium urate | Schorn et al. (2012) |
| Ca ²⁺ ionophore | Palić et al. (2007) |
| Oxidants | Keshari et al. (2013) |
| Heavy metals | Haase et al. (2016) |
| Glucose | Miyoshi et al. (2016) |
| Cholesterol crystals | Neumann et al. (2014a) |
| LL-37 | Tripathi et al. (2014) |
| High-mobility group Box 1 protein | Tadie et al. (2013) |
| Complement component 5a | Huang et al. (2015) |
| Inhibitors | |
| Deoxyribonucleases | Halverson et al. (2015) |
| Vitamin C | Mohammed et al. (2013) |
| Gallic acid | Haute et al. (2015) |
| Acetylsalicylic acid | Lapponi et al. (2013) |
| Metformin | Wang et al. (2015) |
| Antiproteases | Majewski et al. (2016), Zabieglo et al. (2015) |
| Ascomycin | Gupta et al. (2014) |
| Cyclosporine | Gupta et al. (2014) |
| Inhibitors of PAD4 | Knight et al. (2015), Kusunoki et al. (2016) |
| Inhibitors of MPO | Metzler et al. (2011) |
| Inhibitor of NF- κ B | Lapponi et al. (2013) |
| Antibodies against cytokines | Keshari et al. (2012) |
| Thrombomodulin | Shimomura et al. (2016) |
| Activated protein C | Healy et al. (2017) |

Note: PMA, phorbol-12-myristate-13-acetate; LPS, lipopolysaccharide; PAD4, peptidyl-arginine deiminase 4; MPO, myeloperoxidase; NF- κ B, nuclear factor-kappaB.

Table 3

NET-associated proteins and their origin and function.

| Protein name | Localization in cell | Function | Publication |
|--|----------------------|--|---|
| Histones | Nucleus | Bactericidal function | Hirsch (1958); Wang et al. (2011) |
| Myeloperoxidase | Primary granules | Synthesis of HOCl and chromatin decondensation | Papayannopoulos et al. (2010) |
| Neutrophil elastase | Primary granules | Degradation of the histone H1 and modification of other histones | Korkmaz et al. (2008), Massberg et al. (2010) |
| Peptidyl-arginine deiminase 4 | Primary granules | Citrullination of histones | Li et al. (2010), Neeli et al. (2008) |
| Cathepsin G | Primary granules | Antimicrobial activity and stimulation of coagulation | Korkmaz et al. (2008), Massberg et al. (2010) |
| Proteinase 3 | Primary granules | Antimicrobial activity | Korkmaz et al. (2008) |
| α-defensins | Primary granules | Antimicrobial activity | Brook et al. (2016) |
| Bactericidal/permeability-increasing protein | Primary granules | Bactericidal activity | Skopelja et al. (2016) |
| Lactoferrin | Secondary granules | Antimicrobial and anti-inflammatory activity | Baker and Baker (2005) |
| Pentraxin 3 | Secondary granules | Antimicrobial activity and control of autoimmunity | Jaillon et al. (2007) |
| LL-37 | Secondary granules | Antimicrobial activity | Neumann et al. (2014b) |
| Calprotectin | Cytoplasm | Antimicrobial activity and recruitment of inflammatory cells | Urban et al. (2009) |
| Catalase | Peroxisomes | Catalyzation of decomposition of hydrogen peroxide | Palmer et al. (2012a) |

Mechanism of NETs formation in suicidal and vital NETosis

The process of suicidal NETosis, which is the most common type of NETosis, starts with the activation of neutrophils after recognition of an inductor (Figs. 1 and 2). The most effective stimulator of this pathway is PMA; therefore, it is frequently used as an activator of NETosis. Activation of neutrophils leads to the stimulation of protein kinase C activity and this induces formation of NADPH oxidase (NOX) complex on plasma and granule membranes (Fuchs et al., 2007; Guimarães-Costa et al., 2012). Then NOX enzymes ensure the transport of electrons across the plasma membrane and to molecular oxygen, which leads to reactive oxygen species (ROS) production. Under the impact of ROS, plasma and granule membrane break and the content of a nucleus, granules and cytoplasm is mixed (Yang et al., 2016). The next activating step is chromatin decondensation, in which activity of enzymes NE, MPO and peptidyl-arginine deiminase 4 (PAD4) is crucial as proved in murine model (Hemmers et al., 2011; Li et al., 2010; Metzler et al., 2011; Papayannopoulos et al., 2010). NE degrades linker histone H1 and modifies core histones, while MPO intensifies chromatin decondensation by the synthesis of hypochlorous acid (Papayannopoulos et al., 2010). PAD4 deiminases core histone proteins while arginine is converted to citrulline. Citrullination of histones results in their weakened binding to DNA (Neeli et al., 2008). The process of suicidal NETosis is finished after

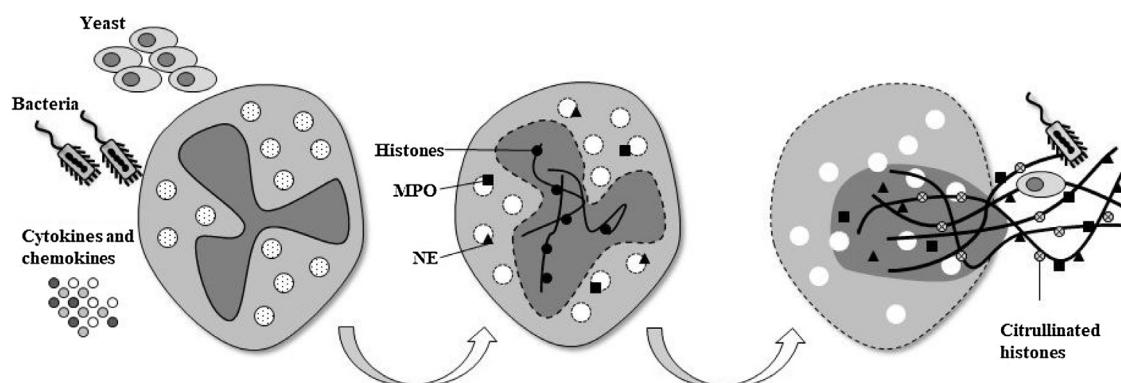
the rupture of plasma membrane and release of NETs into extracellular space.

Contrary to suicidal NETosis, neutrophils stay alive and are able to phagocytose after vital NETosis (Table 1). This process occurs in response to some types of pathogens and is rapid (5–60 min). Vital NETosis consists of a growth of nuclear envelope, chromatin decondensation and disruption of nuclear membrane (Pilsczek et al., 2010; Yipp and Kubes, 2013). However, neutrophils do not lose nuclear and plasma membrane during this type of NETosis. Induction of vital NETosis is through the stimulation of toll-like receptors (TLR) and the complement receptor for C3 protein (Delgado-Rizo et al., 2017). Despite the loss of nuclear DNA, neutrophils maintain their chemotactic ability and are able to chase and phagocytose bacteria.

The special type of vital NETosis is mitochondrial NETosis, which is ROS-dependent contrary to nuclear vital NETosis. In this case NETs are created from mitochondrial DNA (McIlroy et al., 2014). It is still unclear why the neutrophils choose the particular type of NETosis and whether it is dependent on the type or intensity of stimulating factor.

Importance of NETs

Capturing of the pathogens is important for the limitation of their extension from the initial site of infection. The key

**Fig. 1.** Schematic representation of NETosis.

Neutrophil activation by biological or chemical stimuli (cytokines, chemokines, bacteria, yeast . . .) is followed by the nuclear chromatin decondensation and the release of enzymes (MPO = myeloperoxidase, NE = neutrophil elastase) from granules. Histone proteins are citrullinated and neutrophil extracellular traps are released to extracellular space where they trap microorganisms.

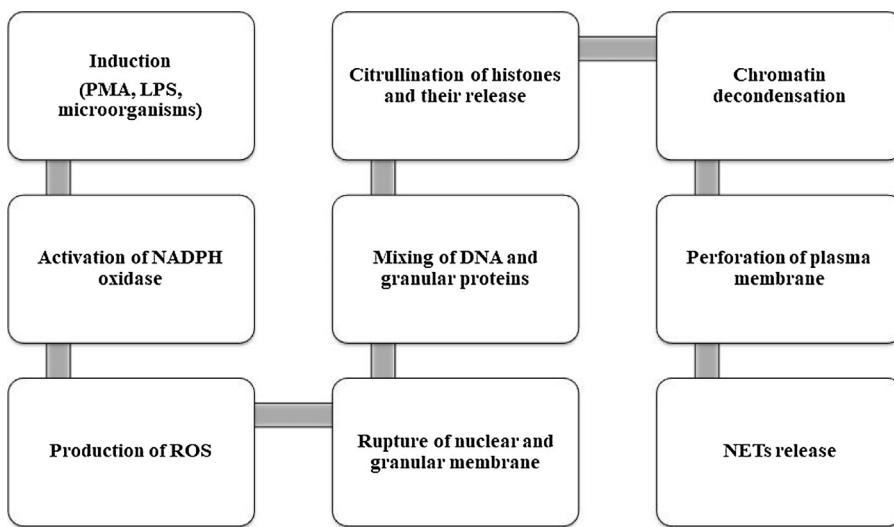


Fig. 2. Mechanism of release of neutrophil extracellular traps (NETs) during suicidal NETosis.

Inductors such as phorbol 12-myristate 13-acetate (PMA) and lipopolysaccharide (LPS) are recognised by receptors and activate NADPH oxidase, which leads to the production of reactive oxygen species (ROS). ROS generation results in the rupture of nuclear and granular membranes and mixing of contents. Citrullination of histones by peptidyl arginine deiminase (PAD4) as well as their degradation with proteases leads to chromatin decondensation and formation of NETs.

mechanism, by which bacteria are trapped by NETs, are the electrostatic interactions between positively charged bacterial membrane and negatively charged DNA of chromatin fibrils (Brinkmann and Zychlinsky, 2007). However, some types of bacteria which have a capsule on their surface or that can change their surface charge, are protected from being caught (Wartha et al., 2007). Another way to avoid entrapping is the production of nucleases and their expression on the surface of the bacteria. This ability to synthesize nucleases protecting them from NETs was demonstrated just in *Streptococcus pyogenes*, *Staphylococcus aureus* and *Pneumococcus* species (Beiter et al., 2006).

The next way, how NETs eliminate the infection is the inactivation of virulence factors that can modify the features of host cells. NE, proteinase 3 and cathepsin G have this ability (Averhoff et al., 2008; Weinrauch et al., 2002). Several components of NETs can stop and kill pathogens in other ways. These include proteases, lysozymes, antimicrobial peptides, ion chelators and histones. The action of these factors and their concentration in chromatin fibrils determine antimicrobial activity of NETs. MPO effectively kills microorganisms during antimicrobial oxidative bursts by converting diffusible long-lived hydrogen peroxide into highly reactive and locally confined hypochlorous acid (Schürmann et al., 2017). Other components of NETs include calgranulin and cathelicidin. Calgranulin as a chelator of zinc has fungicidal activity (Bianchi et al., 2011), while antimicrobial peptide cathelicidin binds to extracellular DNA to keep itself from the endonucleases, thus to prolong NETosis (Neumann et al., 2014b). Histones are bactericidal and antiprotozoal. Indeed, they are also able to kill the cells of a host and are closely related to the pathogenesis of sepsis (Saffarzadeh et al., 2012; Xu et al., 2009) and addition of antibodies against histones result in the inhibition of the killing pathogens (Brinkmann et al., 2004).

Involvement of NETosis in pathophysiologic processes

In contrast to antimicrobial function of NETosis is its involvement in noninfectious, sterile inflammation (Fig. 3). There is evidence that NETosis has an ability to amplify and under specific conditions also overstimulate the immune response. It is unclear whether all NETs are equally harmful for the host and whether NETs from various diseases are phenotypically and functionally identical or not (Kaplan and Radic, 2012).

Alarmins, also called danger-associated molecular patterns (DAMPs), are endogenous molecules recognised by the immune system of a host, able to induce the sterile inflammation. These molecules include heat shock proteins, high-mobility group Box 1 protein, but also extracellular DNA and RNA. Many structural compounds of NETs are alarmins; i.e. DNA as a main component of NETs can be sensed by endosomal TLR9 in plasmacytoid dendritic cells (pDCs) and monocytes, as well as by various cytosolic receptors in macrophages (Lande et al., 2007). Activation of these receptors leads to the synthesis of pro-inflammatory cytokines. The next detrimental side of excessive NETosis is the fact that it leads to tissue damage due to the epithelial and endothelial cell death (Xu et al., 2009). Level of damage and organ dysfunction depends on the intensity of NETs release (Caudillier et al., 2012; Narasaraju et al., 2011). Therefore, NETosis is involved in the pathogenesis of various disorders including cardiovascular and autoimmune diseases. Below, recent discoveries on the potential role of NETs in the pathogenesis of some of them are summarized.

Cardiovascular diseases

NETosis was observed in association with several cardiovascular diseases, such as thrombosis and atherosclerosis. Thrombosis is a formation of a blood clot inside a blood vessel, obstructing the flow of blood through the circulatory system. Neutrophils are accumulated and tightly adhere to the affected endothelium, where they produce NETs. NETs work as a scaffold for capturing of platelets and erythrocytes, thus contributing to the thrombus formation. Due to their negative charge, NETs are able to bind plasma proteins such as fibrin and von Willebrand factor and fix the clot (Fuchs et al., 2010). Moreover, NE and cathepsin G are proteases, which cleave a tissue factor pathway inhibitor and also anticoagulants such as antithrombin and activated protein C, thereby stimulate coagulation, as proved in murine model (Massberg et al., 2010). The next way how NETs can contribute to the thrombus formation is the activation of factor XII and factor VII (von Brühl et al., 2012). Histones are also able to activate platelets and the generation of thrombin (Carestia et al., 2013). The manner, in which histones stimulate coagulation, is the interaction with thrombomodulin and protein C, which leads to the inhibition of activation protein C and results in thrombin generation. Role of

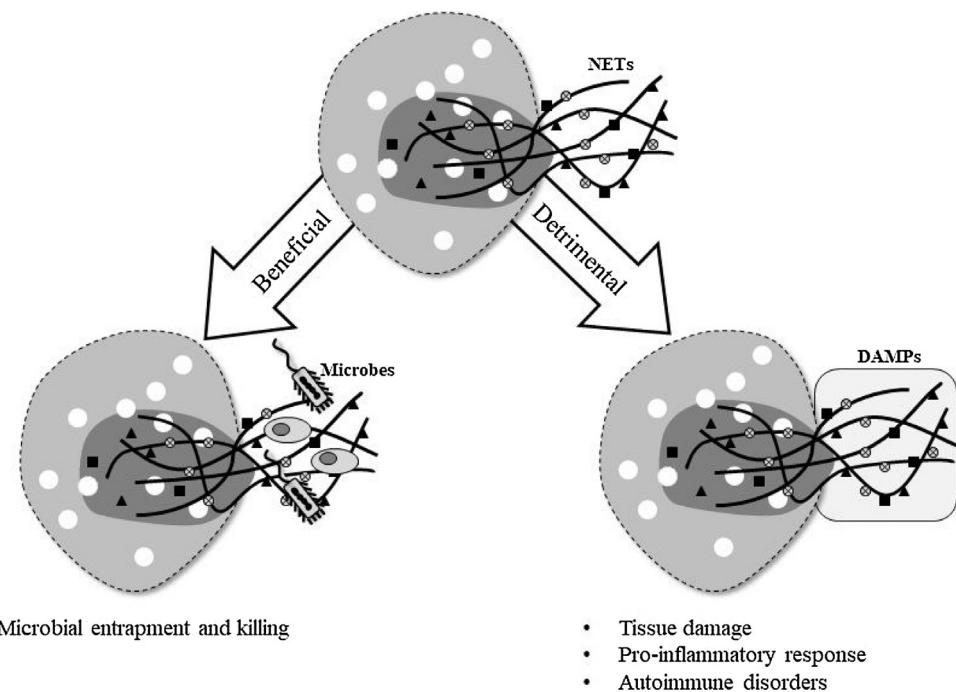


Fig. 3. Advantages and disadvantages of NETosis.

The formation of neutrophil extracellular traps (NETs), induced by pathogens is crucial for microbial trapping and killing. However the excessive generation of NETs results in tissue damage, activation of proinflammatory response and autoimmune disorders. (DAMPs = damage-associated molecular patterns).

NETs and their components in pathophysiology of thrombosis was proved also by deoxyribonuclease (DNase) and the anti-NET antibody treatment, which decreased clot formation in murine model (Brill et al., 2012; Massberg et al., 2010).

Atherosclerosis is a chronic inflammatory disease, in which nodules and plaques build up inside the arteries. Atherosclerotic plaques as well as the damaged vessel walls contain components of NETs, such as peptides LL-37 and its analogue cathelin-related antimicrobial peptide (Cramp) (Döring et al., 2012a, b). Cramp^{-/-} mice fed with high-fat diet exhibit reduced lesion sizes with lower macrophage and neutrophil numbers in comparison to controls (Döring et al., 2012a). Also complexes of citrullinated histones with DNA and MPO/DNA complexes were found in coronary vessel walls in patients with coronary artery disease. These DNA-protein complexes are also present in blood whereby their level increases in direct proportion to the severity of the general atherosclerosis. Moreover, the level of MPO/DNA complexes could serve as a predictive biomarker of the number of coronary vessels involved in atherosclerosis as was proved in the clinical study (Borisoff et al., 2013).

Immunopathogenesis of atherosclerosis is connected with increased level of pDCs in affected vessel walls and occurrence of immunostimulatory DNA-protein complexes. Components of NETs activate pDCs which results in high synthesis of interferons in vessel walls. The removal of pDCs had a positive effect and reduced the formation of atherosclerotic plaques in mice. Activation of pDCs and the production of interferon I resulted in an intensive development of atherosclerotic plaques (Döring et al., 2012b).

Furthermore, NETosis causes damage to the vascular endothelium. The endothelial dysfunction is caused by MMP-9 as the next component of the NETs. This enzyme induces the cell death and causes the dysfunction of endothelium by activation of endothelial MMP-2. The repression of NETosis and MMP-2 leads to the restoration of normal function of endothelium (Carmona-Rivera et al., 2015), therefore the detection of NETs can be a good diagnostic tool.

Autoimmune diseases

Systemic lupus erythematosus (SLE) is a chronic autoimmune systemic disease characterized by the production of autoantibodies, deposition of immune complex in tissues, infiltration of tissues with T and B cells, neutrophils and macrophages, and subsequent inflammatory tissue injury (Liu and Davidson, 2012). Autoantibodies are specifically targeted not only against nuclear antigens such as histones, DNA and ribonucleoproteins, but also against granular proteins of neutrophils typical for NETs. Therefore NETs are considered as a potential source of autoantigens. The deficiency of NET clearance is connected with high titers of anti-NET antibodies and renal involvement in patients with SLE correlates with the lack of NET clearing (Hakkim et al., 2010). Moreover, the concentration of NET-associated proteins, such as defensins, high-mobility group box protein 1, LL-37 and other bactericidal proteins in blood of lupus patients is higher compared to healthy controls. These can then be the good target for autoantibodies and may worsen the disease activity (Ma et al., 2012). The other problem, which can help NETs in exacerbation of the disease, is a deficiency in DNase activity. This lack of DNA cleavage was shown especially in patients with inherited form of SLE, who had a mutation in gene for DNase 1 or DNase1-like 3 (Al-Mayouf et al., 2011; Yasutomo et al., 2001). DNase gene polymorphisms, DNase 1 inhibitors and high amount of anti-NET antibodies were found in SLE patients (Kim et al., 2008; Shin et al., 2004). All these reasons led to the lack of NETs degradation (Hakkim et al., 2010).

As mentioned previously, NETs are recognized by immune system as DAMPs by various receptors, including TLR9, which induces imbalanced immune homeostasis (Kahlenberg et al., 2013). Interestingly, mice lacking functional TLR9 had more serious disease and a shortened lifespan (Nickerson et al., 2013; Stoehr et al., 2011). Moreover, the deficiency of the NOX2, which should inhibit the pathogenesis of the disease in lupus-prone mice by the inhibition of suicidal NETosis, caused exacerbation of SLE

(Campbell et al., 2012). The clarification and characterization of the role of NETs in the pathogenesis of SLE is therefore necessary for the acquisition of new therapeutic approaches.

Rheumatoid arthritis (RA) is an autoimmune disease, which involves mainly the inflammation of joints, but similarly to SLE can also affect internal organs. Pathophysiology of RA as well as SLE is associated with NETs. Serum of RA patients contains a lot of various autoantibodies, the most of them targeted to citrullinated proteins, which are characteristic for NETs (Szekanecz et al., 2008). Neutrophils isolated from blood and synovial fluid of RA patients undergo NETosis more spontaneously compared to healthy controls (Sur Chowdhury et al., 2014). The amount of NETs in blood of RA patients correlated with the presence of anticitrullinated peptide antibodies and cytokine production (Khandpur et al., 2013). Contrarily, the PAD4-deficiency had no effect on the development or severity of the disease in mouse model of RA (Rohrbach et al., 2012). These opposing views on the impact of NETosis in this autoimmune disease can be elucidated just by extensive study and usage of more experimental models.

The next autoimmune disease, in which NETosis could play a role, is type 1 diabetes mellitus (T1DM). T1DM is characterized by the inability of pancreas to produce insulin due to the autoimmune destruction of beta cells resulting in hyperglycemia. Several studies showed that T1DM patients suffer from neutropenia, which may be attributed to infiltration of neutrophils in pancreatic tissue and increased NETosis (Harsunen et al., 2013; Valle et al., 2013). Neutrophils exposed to hyperglycemic conditions are stimulated to produce ROS and cytokines. A concentration of TNF- α is increased in plasma of diabetic patients and this induces NETosis. By this way NE, cathepsin G and proteinase 3 are freed while elevated concentration and activity of these enzymes are shown in T1DM patients (Wang et al., 2014). Moreover, these proteins are able to induce the production of cytokines and expression of TLRs, which are mediators of insulitis and beta cells destruction (Padgett et al., 2013).

Cystic fibrosis

Cystic fibrosis is a hereditary autosomal recessive disorder that causes severe damage to the lungs, digestive system and other organs in the body. Cystic fibrosis affects the cells that produce mucus, sweat and digestive juices and is characterized by the increased number of neutrophils as well as NETs in the lungs (Marcos et al., 2010). High level of extracellular DNA, some of which originates from the NETs, increases sputum viscosity in patients with cystic fibrosis (Yoo et al., 2014). In order to reduce the sputum viscosity, these patients are treated with recombinant DNase I. However, DNase I is not able to effectively cleave the complexes consisting of DNA and histones typical for NETs because a lot of cleavage sites are hidden. The characterization of the chromatin structures in sputum from patients who were not under DNase I therapy showed that previous treatment with NE enhances solubilisation of sputum by cleaving histones, increasing the access of DNases to DNA (Papayannopoulos et al., 2011). The appropriate adjustment of therapy targeted to reduce the NETs in lungs than can be useful for the treatment not just the cystic fibrosis patients but also patients with other diseases associated with sputum accumulation, such as chronic obstructive pulmonary disease (Grabcanovic-Musija et al., 2015).

Periodontitis

Periodontitis is one of the most common chronic inflammatory diseases, which causes serious gum infection that damages the soft tissue and destroys the bone that supports teeth and causes their loss (Farquharson et al., 2012; Kassebaum et al., 2014). The first

step of the initiation of the development of periodontitis is a disbalance within the accumulated plaque of biofilm followed by bacterial infections (Bascones-Martínez et al., 2009). Neutrophils are important for bacterial clearance during infection but in sensitive patients, they are not able to eliminate the pathogenic bacteria. Production of cytotoxic components including ROS during NETosis is important to destroy the bacteria, however neutrophils isolated from chronic periodontitis patients are hyper-reactive regarding ROS release, which may lead to the progression of the disease (Matthews et al., 2007).

Long-term periodontitis and hypercoagulability may cause the atherothrombosis (Demmer and Desvarieux, 2006). It is suggested that repeated injections with *Porphyromonas gingivalis*, which is commonly responsible for periodontitis, cause abdominal aortic aneurysms as proved in rat model (Delbosc et al., 2011). In this study, thrombi observed in rats after the infection contained high level of NETs (Delbosc et al., 2011). Determination of the concentration of mediators, which are able to stimulate the production of NETs, such as interferon- α in serum of periodontitis patients showed that NETosis may be an effective biomarker of periodontitis (Wright et al., 2008). On the contrary, reduced amount of NETs in serum of patients with periodontitis may increase their sensitivity to the colonization of bacteria. This effect may be due to production of DNase by bacteria. Palmer et al. suggested that many periodontal bacteria release DNases, which can help them to enter the tissue and prevent their abolishment (Palmer et al., 2012b). The same authors also demonstrated that the activation of neutrophils to release the NETs differed between specific periodontal bacteria *in vivo* (White et al., 2014), which can also help them with persistence of the infection. The role of NETosis in periodontitis is undoubtedly important, but further studies are needed for its explanation and better understanding.

Conclusion

The evidence that NETosis participates on pathophysiology of cardiovascular diseases, autoimmune diseases as well as other pathologies such as cancer and Alzheimer disease increases during last years. Regardless of intensive research and the development of this field of study, there are still lots of questions, which are important for better understanding of the pathophysiological impact of NETosis. It is necessary to clarify the critical boundary when NETosis changes from helpful to destructive as well as to characterize if these neutrophils and their NETs, which participate on diseases, have any composition differences. Understanding precisely if sterile inflammatory NETs differ structurally from NETs released during infection requires the progress in this field. More work in experimental models of diseases as well as prospective studies are also needed to assess if presence of NETs and their components can be used as biomarkers in patients with autoimmune and other diseases. Moreover, the lack of the standardized methods for the detection of NETs and NET-associated proteins is also a problem because there are no commercial kits for reliable detection. The simple visualization and quantification of NETs using DNA-intercalating dyes or immunofluorescence microscopy using antibodies against NET-associated proteins have limitations and it is time-consuming. Thus, there is a requirement for techniques, which are able of robust and rapid evaluation of NET-releasing cells, for example, by flow cytometry combined with imaging. The solving of these problems and ambiguities would be helpful for more accurate diagnosis, prognosis and for the target of the treatment of many diseases.

Conflicts of interests

The authors declare no conflict of interests.

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