Tailor-made inflammation: how neutrophil serine proteases modulate the inflammatory response

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Abstract Neutrophil granulocytes are important mediators of innate immunity, but also participate in the pathogenesis of (auto)inflammatory diseases. Neutrophils express a specific set of proteolytic enzymes, the neutrophil serine proteases (NSPs), which are stored in cytoplasmic granules and can be secreted into the extra- and pericellular space upon cellular activation. These NSPs, namely cathepsin G (CG), neutrophil elastase (NE), and proteinase 3 (PR3), have early been implicated in bacterial defense. However, NSPs also regulate the inflammatory response by specifically altering the function of cytokines and chemokines. For instance, PR3 and NE both inactivate the anti-inflammatory mediator progranulin, which may play a role in chronic inflammation. Here, we provide a concise update on NSPs as modulators of inflammation and discuss the biological and pathological significance of this novel function of NSPs. Mounting evidence support an important proinflammatory function for PR3, which may have been underestimated in the past.

Keywords Neutrophil serine proteases · Inflammation · Neutrophil elastase · Proteinase 3 · Cathepsin G · Progranulin

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Introduction

Neutrophil granulocytes make up the majority of all leukocytes in peripheral, human blood, where they circulate through the body and swiftly react on danger signals from injured or infected tissues. Once infection or wounding occurs, neutrophils are one of the first cell types that infiltrate the harmed tissue, where they release antimicrobial factors and reactive oxygen species (ROS) to kill pathogens and disinfect the injured tissue. Neutrophils contribute to inflammation and bacterial defense mainly via three distinct cell biological mechanisms, namely by phagocytosis, degranulation or the formation of neutrophil extracellular traps (NETs) [1]. Bacterial infections are usually battled via phagocytosis of the pathogens, i.e., the receptor-mediated uptake of bacteria followed by degradation of the pathogen inside the phagolysosome. Neutrophils are pre-charged with antimicrobial factors such as neutrophil serine proteases (NSPs), which are stored in cytoplasmic organelles, the so-called azurophilic granules, ready to be released into the extracellular space by degranulation. When neutrophils encounter bacteria, they can also undergo an alternative cell death program leading to the extrusion of their own nuclear chromatin to form NETs, extracellular chromatin fibers decorated with antimicrobial proteases that trap and kill bacteria even after neutrophil cell death [2]. However, uncontrolled neutrophil activation can lead to chronic inflammatory conditions such as autoimmune diseases or hypersensitivity reactions. For instance, neutrophil activation by autoantibodies in autoimmune small-vessel vasculitis promotes the formation of NETs, which are deposited in the kidneys of these patients and are likely to drive the disease process and the autoimmune response against myeloperoxidase and PR3 present on these NETs [3]. Recent studies have clarified that the physiological role
of NSPs is not limited to the degradation of microbes, but also involves specific immunoregulatory functions that affect the cascade of events during the inflammatory reaction [4]. In this minireview, we provide a concise update on the role of NSPs as inflammatory mediators and discuss the biological and pathological significance of these events.

Neutrophil serine proteases

Neutrophils express several classes of proteases including matrix metalloproteinases such as matrix metalloproteinase (MMP)-8 and -9, cysteine proteases such as caspases or calpain, and NSPs. Serine proteases are a class of proteases with the amino acid serine as the key residue of their enzymatic center, which initiates the proteolytic cleavage of specific target proteins, the protease substrates. The serine proteases of neutrophils cathepsin G (CG), neutrophil elastase (NE), and proteinase 3 (PR3) are crucially involved in bacterial defense [5], but also mediate regulatory functions that shape the inflammatory response [4]. While PR3 and NE are very similar in their substrate specificity supporting a potentially redundant function for these enzymes, CG usually targets a different set of substrate proteins.

Unregulated extracellular proteolysis would be detrimental to the entire organism. Hence, the liver constantly produces copious amounts of physiological serine protease inhibitors, such as α1-antitrypsin (α1-AT), α1-antichymotrypsin, and α2-macroglobulin, which are distributed throughout the body in the plasma and interstitial fluids and efficiently remove NSPs and other proteases after their release by complexation and delivery to lysosomes of phagocytic cells. Moreover, there are tissue specific inhibitors of NSPs such as elastin or secretory leukocyte protease inhibitor (SLPI), which control serine protease activity in interstitial fluids. Disruption of the balance between protease and their inhibitors as it occurs in patients with genetic α1-AT deficiency leads to the development of lung emphysema and certain autoimmune disorders [6]. This reflects the involvement of NSPs in the progression of inflammatory conditions and tissue breakdown.

NSPs control inflammatory reactions

The role of NSPs in inflammation has been controversial for several years. Most knowledge about the function of NSPs has been gained from in vitro studies using purified enzymes. Initial findings that NSPs cleave extracellular matrix components suggested that these enzymes might facilitate the egress of leukocytes from blood vessels and pave the way towards the sites of inflammation. More recent reports have provided interesting in vitro results showing that NSPs cleave several chemokines and cytokines, which leads to enhancement or abolishment of the function of these immune mediators [7, 8]. Some of these studies analyzed ex vivo specimens by comparing chemokine or cytokine cleavage in specimens from patients suffering from inflammatory conditions with those from healthy control individuals (summarized in Supplementary Table 1). However, the biological significance of these data remains to be addressed using adequate experimental in vivo models in order to elucidate if a substrate is in fact targeted by the protease under physiological conditions.

The recent generation of NSP-deficient mice has formed the basis for the in vivo evaluation of the role of these proteases during inflammation. For instance, NE-deficient animals have been used to determine whether this enzyme is critical for neutrophil extravasation during inflammation as suggested by previous in vitro results. However, these papers have yielded conflicting results regarding the effect of NE-deficiency on neutrophil extravasation to sites of inflammation, since NE may or may not be important for leukocyte migration across the basement membrane [9, 10]. One of the first in vivo findings that supported an immunoregulatory role of NSPs were made in mice deficient in dipeptidyl peptidase I (DPPI), another protease that mediates the conversion of NSPs from their inactive precursors (zymogen) into the enzymatically active protease by removal of an N-terminal pro-peptide [11]. Neutrophils of DPPI-deficient mice showed very little NSP enzymatic activity and demonstrated strongly decreased infiltration to the sites of inflammation in an experimental arthritis model, although neutrophil migration in vivo appeared generally normal in these mice. Of importance, this was accompanied by decreased local production of inflammatory cytokines. These findings suggest that NSPs modulate cellular events such as neutrophil activation and chemokine production, while at least in this setting they are not directly necessary for the crossing of the endothelial basement membrane during neutrophil diapedesis.

Cellular activation of neutrophil granulocytes during the inflammatory response leads to the substantial release of NSPs into the extracellular space, where the proteolytic activity of these enzymes is tightly regulated by local protease inhibitors (Fig. 1a). There are several ways how NSPs can be active despite the presence of protease inhibitors. For instance, previous studies have revealed that NSPs compartmentalize to the cell surface of neutrophils where they may be shielded from inhibitor binding and therefore fully active [12]. NSPs are also present on extracellular chromatin fibers called NETs [2, 3]. It remains to be determined whether NSPs on NETs are enzymatically active and whether their interaction with endogenous inhibitors is altered. Moreover, inhibitors of NSPs can be degraded by other proteases released from activated neutrophils such as MMP-9, which degrades the serpin

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α1-PI and thus prolongs local NE proteolytic activity [13]. Hence, the pericellular microenvironment around a neutrophil may be depleted from local inhibitors leading to increased NSP activity at the site of inflammation.

But how do NSPs regulate the course of an inflammatory response on the molecular level? In the following parts, we will highlight selected actions that involve NSP function during the inflammatory response.

Regulation of chemokine function

The coordination of leukocyte migration to a damaged tissue is essentially mediated by a concentration gradient of chemokines, which is sensed by the migrating cell using G protein-coupled receptors. Extracellular proteases modulate chemokine function in several ways, which may lead to abrogation or enhancement of chemokine function, and sometimes alters the bioavailability of the chemokine.

Proteolytic abrogation of chemokine function often leads to an inflammation-dampening effect or changes the subsequent recruitment of immune cells. The stroma-derived factor 1α (SDF-1α) is an important lymphocyte-attracting chemokine. The proteolytic cleavage of SDF-1α by NE on the endothelium during neutrophil extravasation inactivates the chemokine and leads to decreased secondary recruitment of lymphocytes to the site of inflammation.
Therefore, inactivation of SDF-1α by NE seems to be an important step that controls the temporal pattern of immune cell trafficking to inflammatory sites [14].

A common mechanism of chemokine alteration that frequently involves NSP function is the proteolytic truncation at the N-terminus of the chemokine. In many cases, this removes several amino acids N-terminal to the chemotactic motif without destroying its biological function, but rather leading to enhanced chemotactic potential of the molecule [8]. Interleukin 8 (IL-8) is one prominent neutrophil-attracting chemokine that is cleaved by all three members of NSPs with differential effects on its biological function. Interestingly, while CG and NE mediated N-terminal truncation of IL-8 leads to inactivation of the chemokine, PR3 proteolytically converts IL-8 into a more potent neutrophil activating version [15]. This may have important implications during the recruitment of inflammatory cells to the site of inflammation (Fig. 1b). However, the biological significance of this proteolytic conversion of chemokines remains to be evaluated in vivo. While there is no direct homolog of IL-8 in mice, the respective IL-8 receptor homolog is known to be activated strongest by macrophage inflammatory protein 2 (MIP-2), and by the approximately tenfold less potent chemokine KC [16]. Hence, analyzing the cleavage of MIP-2 and KC in mice, for instance in NSP-depleted mice, will provide a useful tool to address the biological significance of proteolytic modulation of IL-8.

Conversion of inflammatory mediators

The posttranslational conversion of cytokine precursors by extracellular proteases determines the activation of cytokines and the time frame of their activity. For instance, all three members of the NSPs have been shown to inactivate the soluble IL-6 receptor, thus interfering with IL-6 signaling, which may contribute to the inflammatory condition in patients suffering from cystic fibrosis [17]. Another key proinflammatory mediator is tumor necrosis factor α (TNFα) that is produced as a membrane-attached precursor and activated by proteolytic conversion after which the active form of TNFα is solubilized. This is mostly carried out by a metalloproteinase, named TNFα-converting enzyme (TACE, ADAM17) [18]. Interestingly, also PR3 seems to contribute to TNFα conversion for example in pneumonia, as indicated by experiments using alveolar macrophages, which, although they show no expression of PR3, seem to carry significant amounts of PR3 on their cell surface [19]. This might be an important function of PR3 during the establishment of an inflammatory milieu (Fig. 1c). Moreover, PR3 proteolytically activates further inflammatory cytokines including IL-1β [20], IL-18 [21], as well as IL-32, a recently discovered cytokine that may play a crucial role during lung inflammation by triggering the release of neutrophil-attracting IL-8 from epithelial cells [22]. Although caspase-1 is probably the most important IL-1β-activating protease, results from DPP1-deficient mice, which lack NSP enzymatic activity, argue for an important role of PR3 in the conversion of this potent inflammatory mediator in a mouse model for rheumatoid arthritis [23]. The fact that all these cytokines are selectively converted by PR3, but not by CG or NE, points towards a distinct proinflammatory role of PR3 during the establishment of an inflammatory environment. However, these findings are mainly based on in vitro studies and remain to be corroborated by in vivo experiments to determine the biological relevance of PR3-mediated conversion of these cytokines.

Progranulin (PGRN; also known as granulin-epithelin precursor) is an important regulator of the wound-healing response [24]. Zhu and co-workers have described that PGRN potently suppresses the adhesion-dependent oxidative burst and release of proteases by neutrophils in the presence TNFα, however, it loses its anti-inflammatory activity after cleavage by NE [25]. Recent studies using PR3/NE double-deficient mice revealed that PGRN is degraded by both PR3 and NE during neutrophil activation in vitro and immune complex-mediated inflammation in vivo [26]. In the absence of PR3 and NE, anti-inflammatory PGRN was not degraded any more leading to significantly higher PGRN levels during the inflammatory response. This may inhibit the activation of infiltrating neutrophils and most likely leads to significantly decreased feedback recruitment of inflammatory cells, which explains the impaired inflammatory response to immune complex deposition in PR3/NE-deficient mice (Fig. 1d). This mechanism may play an important role in the establishment and control of the chronic inflammation. Interestingly, loss-of-function mutations in the PGRN gene are associated with the development of neurodegenerative disorders such as dementia, Alzheimer, and Parkinson disease [27]. Indeed, recent studies using PGRN-deficient mice suggest that decreased levels of PGRN in these patients leads to a chronic, subtle inflammatory condition, which may finally manifest in age-related neurodegenerative disease [28]. The details about the exact molecular pathway of the anti-inflammatory effects linked with PGRN yet remains to be elucidated.

NSPs engage surface receptors

As mentioned earlier, NSPs often bind to surface receptors, once they are externalized by activated neutrophils. This can create an auto- or paracrine effect, since several receptors transduce signals that promote neutrophil activation or alter epithelial cell function. For instance, CG
appears to trigger integrin clustering on neutrophils interacting with immobilized immune complexes, which leads to cytoskeletal rearrangements, increased ROS production, and secretion of chemokines such as MIP-2 [29]. These findings suggest an important role of CG during immune complex-induced inflammation (Fig. 1e). However, the exact molecular mechanism as to how CG modulates integrin clustering on the cell surface remains elusive.

Another intriguing pathway of NSP-mediated inflammation is based on the engagement of protease-activated receptors (PARs), which are G protein-coupled receptors mostly expressed on platelets and epithelial cells that are activated by proteolytic cleavage of the extracellular domain. All three NSPs have been implicated in activating PAR-2 on human oral epithelial cells and gingival fibroblasts, which resulted in the release of proinflammatory mediators such as IL-8 [30], suggesting that PAR-activation by NSPs may contribute to the inflammatory process for example during periodontitis. Moreover, NE and PR3, but not CG, seem to be involved in activating PAR-1 and PAR-2 on epithelial cells leading to downregulation of epithelial barrier function and increased leukocyte transmigration, a process that may be important in the pathogenesis of inflammatory bowel disease [31]. Although NSPs may be important activators of PARs, many aspects of this relationship remain elusive. Here again, most findings are based on in vitro studies and would require further confirmation in the living animal. For instance, PAR-2 is known to be expressed on neutrophils, however, it is still unclear whether NSPs externalized by stimulated neutrophils promote an autocrine feedback on neutrophil activation via PAR-2.

The toll-like receptors (TLRs) represent another class of intra- and extracellular membrane-anchored molecules that sense inflammatory stimuli such as bacterial DNA or lipopolysaccharide (LPS) and mediate cellular activation via activation of the nuclear factor κB pathway. Interestingly, NE has been found to activate TLR-4, the surface receptor for LPS, leading to release of IL-8 by a human embryonic kidney cell line [32]. This process appears to depend on proteolytic activity as it was inhibited by phenylmethyisulphonyl fluoride, a broad spectrum serine protease inhibitor. However, the exact molecular pathway of TLR-4 activation by NE needs yet to be addressed and, although NSPs can modulate the function of integrins, PARs, or TLRs, there is yet no direct evidence that NSPs actually bind to these classes of cell surface molecules.

Synergistic effects

Many of the inflammatory pathways modulated by NSPs are interconnected, so that, hypothetically, NSPs may work synergistically to accelerate the defense reaction. For instance, NSPs mediate the release of IL-8 by engaging several surface receptors such as TLRs, PARs, or integrins [29, 30, 32]. The bioactivity of IL-8 may then be enhanced by PR3-mediated truncation of the chemokine N-terminus [15], leading to increased recruitment of more neutrophils to the site of inflammation. Moreover, the generation of active TNFα by PR3 triggers the externalization of further NSPs by neutrophil granulocytes. Likewise, PGRN inhibits the release of NSPs by neutrophils [25], thus the degradation of PGRN by PR3 and NE may drive neutrophils to release additional NSPs. Furthermore, NSPs often trigger increased production of ROS by neutrophils, e.g., via the promotion of integrin clustering or by degrading anti-inflammatory PGRN [26, 29]. Given that these oxidants inactivate physiological protease inhibitors [33], NSPs indirectly prolong the time frame of their actions by promoting ROS production.

Conclusion and outlook

Although NSPs sometimes inactivate certain inflammatory cytokines and chemokines, the function of NSPs generally promotes rather than suppresses the inflammatory response. Especially the relevance of PR3 as a proinflammatory mediator may have been underestimated in the past. Most of the target molecules have been identified by in vitro studies, hence the biological significance of these pathways often remains to be confirmed using in vivo using appropriate animal models. The three neutrophil enzymes CG, NE, and PR3 seem to synergize with each other during the establishment of an inflammatory milieu. Thus, the generation of NSP inhibitors specific for all three NSPs may be a potent inflammation-suppressing drug that could be used to treat patients with chronic inflammatory disorders.

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