

How anti-neutrophil cytoplasmic autoantibodies activate neutrophils

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R. Kettritz

*Nephrologie und Internistische Intensivmedizin
Charité Virchow Klinikum and Experimental and
Clinical Research Center, a joint co-operation
between the Charité Medical Faculty and the
Max-Delbrück Center for Molecular Medicine,
Berlin-Buch, Berlin, Germany*

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Correspondence: R. Kettritz,
Universitätsmedizin Berlin, Nephrologie und
Internistische Intensivmedizin Charité Virchow
Klinikum und Experimental and Clinical
Research Center, a joint co-operation between
the Charité Medical Faculty and the
Max-Delbrück Center for Molecular Medicine,
Lindenberger Weg 80, 13125 Berlin, Germany.
E-mail: kettritz@charite.de

Summary

Neutrophils are pivotal to host defence during infectious diseases. However, activated neutrophils may also cause undesired tissue damage. Ample examples include small-vessel inflammatory diseases (vasculitis) that are associated with anti-neutrophil cytoplasmic autoantibodies (ANCA) residing in the patients' plasma. In addition to being an important diagnostic tool, convincing evidence shows that ANCA are pathogenic. ANCA–neutrophil interactions induce important cellular responses that result in highly inflammatory necrotizing vascular damage. The interaction begins with ANCA binding to their target antigens on primed neutrophils, proceeds by recruiting transmembrane molecules to initiate intracellular signal transduction and culminates in activation of effector functions that ultimately mediate the tissue damage.

Keywords: ANCA, neutrophils, pathogenesis, signal transduction, vasculitis

Anti-neutrophil cytoplasmic autoantibodies (ANCA) and the expression of ANCA target antigens by the neutrophil

ANCA must recognize and bind their target antigens, proteinase 3 (PR3) or myeloperoxidase (MPO), in order to initiate signalling events and to subsequently activate the neutrophil. Thus, ANCA must either be internalized by the neutrophil or the antigens must be accessible on the cell surface, or both may occur. Many studies exploring the membrane expression of ANCA antigens have been performed. MPO and the vast majority of PR3 antigens reside in azurophilic granules, which can be mobilized during activation *in vitro* and *in vivo* [1,2]. In contrast to MPO, PR3 is also stored in specific granules and in secretory vesicles that are mobilized more easily [3]. Moreover, significant PR3 amounts are already expressed on the surface of resting cells with a strong increased expression after activation. Thus, there are major differences in PR3 and MPO membrane expression. Notably, and in contrast to PR3, MPO is not detected on the plasma membrane of resting neutrophils. Furthermore, the membrane MPO that increases after cell activation is small compared to PR3.

Neutrophils must be primed for subsequent ANCA-induced activation. Priming includes ANCA antigen translocation and can be achieved *in vitro* by various mediators, including tumour necrosis factor (TNF)- α , interleukin (IL)-1, IL-6, IL-18, N-formyl-Met-Leu-Phe (fMLF) and complement 5a (C5a) [4–7]. *In-vivo* priming may occur during infections that frequently precede the clinical manifestation of ANCA vasculitis. Indeed, patients with active disease show increased neutrophil ANCA antigen membrane expression [5,8,9]. A synergistic effect for increased mPR3 expression by cytokines, adhesion and anti-PR3 antibodies was demonstrated that could become relevant when neutrophils leave the circulating blood [10]. Recently, α 1-antitrypsin polymers have been described to prime the neutrophil for ANCA activation, indicating that additional priming mechanisms exist [11].

An important observation established that PR3, but not MPO, has a bimodal membrane expression pattern. mPR3^{low}- and mPR3^{high}-expressing neutrophils can be distinguished with a percentage of mPR3^{high} neutrophils ranging between 0 and 100% [12]. The expression pattern is extremely stable in a given individual, does not change with activation and correlates strongly in monozygotic twins and

human leucocyte antigen (HLA)-matched siblings [13,14]. The clinical significance of the mPR3 phenotype was established in independent cohorts showing that a large subset of mPR3^{high} neutrophils is a risk factor for ANCA vasculitis. The risk factor has a negative effect on clinical patient outcomes [13,15–17]. Compared to the mPR3^{low} cells, mPR3^{high} neutrophils generate more superoxide and degranulate more strongly to PR3–ANCA, but not to other stimuli. This provides a potential explanation for the clinical observation on risk and outcome [18].

Because MPO and PR3 are not transmembrane molecules, elucidating how ANCA antigens are anchored in the plasma membrane is another important step in understanding how signal transduction may begin. PR3 presentation on the neutrophil membrane occurs by at least two different mechanisms. PR3 can be inserted directly into the plasma membrane, as predicted by molecular dynamics simulations using a membrane model [19]. This model suggested that PR3 associates strongly with anionic membranes, whereby basic residues mediate the orientation of PR3 at the membrane and hydrophobic amino acids mediate anchoring of the molecule. Kantari *et al.* mutated the basic and the hydrophobic amino acids and observed that the modified PR3 preserved its enzymatic activity. However, the mutated protein lost its plasma membrane expression in a myeloid rat basophilic leukaemic (RBL) cell model [20]. Another way of expressing PR3 on the neutrophil membrane is its presentation by a glycosylphosphatidylinositol (GPI)-linked receptor, namely CD177 (or human neutrophil antigen B1, NB1) [21,22]. Although all neutrophils contain intracellular PR3, only those cells that express NB1 protein on the neutrophil plasma membrane show high mPR3 surface expression. Studies have demonstrated further that PR3 and NB1 were not only co-expressed on the same neutrophil subset, but that both molecules co-localize and co-immunoprecipitate. Co-transfection experiments in human embryonic kidney 293 (HEK293) cells showed that NB1 was a sufficient receptor for PR3, but not for pro-PR3 [23]. Future experiments need to elucidate the control mechanisms of NB1 expression and why only a subset of neutrophils generates NB1 protein. Korkmaz *et al.* showed that a unique hydrophobic patch, present on human and absent from gibbon and murine PR3, enabled binding to NB1 [24]. Choi *et al.* performed high-throughput screening using a small molecule library and identified compounds that inhibited the interaction between NB1 and PR3 [25]. Kuhl *et al.* characterized conformational PR3 epitopes recognized by monoclonal anti-PR3 antibodies or PR3–ANCA from patients. These epitopes are distinct from the catalytic site and from the hydrophobic patch that allowed binding to membranes and NB1 [26]. All these studies have improved our understanding with respect to PR3 presentation on the neutrophil membrane and its accessibility for anti-PR3 antibodies. Consistent with the co-expression of NB1 and PR3 on the same cell, a larger percentage of mNB1-expressing neutrophils was a risk factor

for ANCA vasculitis [27]. The role of the lacking PR3–NB1 interaction in mice could be one reason for the difficulty in generating an anti-PR3 antibody-mediated disease model, and needs further study.

We have reviewed the data describing modes of ANCA antigen expression on the neutrophil membrane and how ANCA can bind to their targets on the plasma membrane to initiate activation. Also conceivable is the possibility that ANCA internalization by the neutrophil contributes to activation. In fact, ANCA penetration into neutrophils has been observed by different investigators; however, the mechanisms and significance of this observation for the activation process are not yet understood [9,28,29]. Furthermore, reactivation of PR3 and MPO transcription has been observed and epigenetic mechanisms that control this process are beginning to be characterized [30,31]. It will be interesting to see if this process results in a protein or cellular localization distinct from those of the 'original' PR3 antigen.

An additional ANCA target is the lysosomal membrane glycoprotein lysosomal-associated membrane protein 2 (LAMP-2) that was implicated in pauci-immune necrotizing glomerulonephritis by Kain *et al.* [32,33]. LAMP-2 is a heavily glycosylated protein expressed in many cell types, including neutrophils and endothelial cells. Lysosomal membrane proteins were detected in membranes of different cellular compartments such as lysosomes, multi-vesicular bodies, the trans-Golgi and plasma membranes [34]. LAMP-2 was found mainly in granule membranes of resting neutrophils and its plasma membrane expression was increased with fMLF treatment [32]. The clinical significance of LAMP-2 as an ANCA antigen in small vessel vasculitis was challenged by the Chapel Hill group. The investigators found much lower anti-LAMP antibody titres compared with antibodies to PR3 and MPO, no correlation with vasculitis disease activity and no disease induction by passive antibody transfer into rats [35]. Kain *et al.* were able, very recently, to repeat their findings in different European patient cohorts [36]. The conflicting data have no obvious explanation, but may be related to methodological and population differences as discussed by Flint *et al.* [37]. Major findings with respect to ANCA antigens are summarized in Fig. 1.

The ANCA antigen-binding site and the Fc part co-operate in neutrophil activation

Once ANCA have bound their neutrophil-expressed antigens, signalling and activation are initiated. Several investigators have characterized the part of the ANCA molecule that is important for neutrophil activation. Conflicting data exist, but the emerging picture is that both the antigen-binding part and the Fc part are needed. We found that ANCA Fab bind to their antigens expressed on the neutrophil, but did not trigger activation. In contrast, ANCA

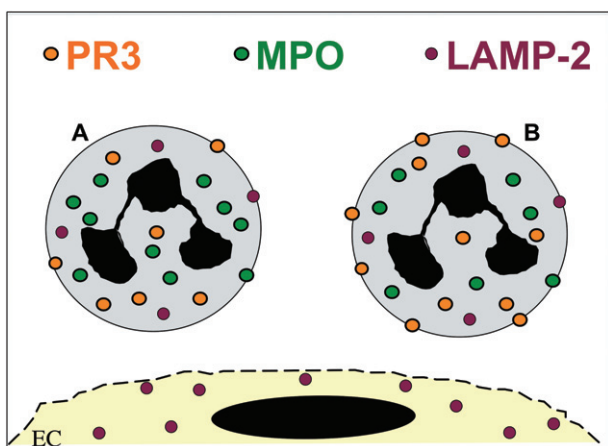


Fig. 1. Anti-neutrophil cytoplasmic autoantibodies (ANCA) antigen expression. Myeloperoxidase (MPO) and proteinase 3 (PR3) are expressed by neutrophils and monocytes only, whereas lysosomal-associated membrane protein 2 (LAMP-2) is also found in endothelial cells. PR3 has a bimodal expression pattern on the neutrophil membrane in that a mPR3^{low}- and mPR3^{high}-expressing subset can be distinguished.

F(ab)2, or ANCA Fabs that were cross-linked, did activate superoxide generation. In the same study, and in contrast to these human ANCA data, F(ab)2 from a murine monoclonal antibody (mAb) had no activating capacity [38]. A PR3- and MPO-ANCA F(ab)2-induced respiratory burst was confirmed in another study [39], but not observed by other investigators [40–42]. The use of human *versus* murine antibodies, the strength of the activation response, assaying intra- or extracellular oxidant generation and the antigen specificity of the antibodies that were employed may, at least in part, explain some of the differences in the results. Williams *et al.* observed that ANCA F(ab)2 induced p21ras activation that was necessary, but not sufficient, for the respiratory burst [43]. Moreover, gene arrays showed that both ANCA F(ab)2 and ANCA immunoglobulin (Ig)G induce leucocyte gene transcription [44]. Interestingly, some of the transcribed genes were unique to intact ANCA IgG and some to the F(ab)2, whereas others were induced by both fragments. Thus, ANCA F(ab)2 bind to the neutrophil and trigger several neutrophil responses that do not depend on FcγR engagement. Few studies investigated this issue in monocytes. Weidner *et al.* showed that human ANCA also activated respiratory burst in monocytes and that ANCA F(ab)2 triggered a similar response compared to the complete ANCA IgG [45].

In addition to the antigen-binding fragments, the Fc part of the ANCA molecule is also important. ANCA IgG bind to FcγRIIa (CD32A) and FcγRIIIb (CD16B). FcγRIIa blockade abrogated ANCA-induced activation, whereas the role of the FcγRIIIb blockade is somewhat more controversial [38,40–42,46]. The FcγRIIa has two allelic variants with either a histidine or an arginine at amino acid position 131,

resulting in a high-responder and low-responder receptor form. Neutrophils with the high-responder variant showed a stronger response to anti-PR3 and anti-MPO IgG1 mAbs *in vitro* [40]. This FcγRIIa also has high affinity to the IgG3 subclass, which is the dominant ANCA subclass in patients with active disease, and had the strongest capability to induce neutrophil adhesion *in vitro* [47,48].

Kocher *et al.* observed that ANCA IgG also bind to the FcγRIIIb on neutrophils that is expressed approximately 10 times higher than the FcγRIIa [46]. Distinct patterns of CD11b increase and CD62L shedding suggested that FcγRIIIb is involved in ANCA-induced neutrophil activation. FcγRIIIb has two common genetic variants named NA1 and NA2, the former triggering a stronger neutrophil activation than the latter. A recent study on a large cohort of patients with granulomatosis with polyangiitis (GPA, also known as Wegener's granulomatosis) demonstrated a similar NA1 allele frequency in patients compared to controls. However, the presence of NA1 was associated with more severe renal disease [49]. The FcαR (CD89) also has genetic variants characterized by a single nucleotide polymorphism (SNP) that changes the amino acid sequence in the coding region. A serine (A) is associated with less inflammatory cytokine release and a glycine (G) with more phagocytosis and cell activation [50]. Kelley *et al.* studied also IgA ANCA and the SNP variants of the FcαR in their GPA patient cohorts [49]. IgA ANCA were present in 27% of the GPA patients, and were less frequent in those patients who developed end-stage renal disease and more frequent in those with upper airway manifestation. The G allele was, however, found more frequently in patients with renal disease and less frequently in those with upper airway manifestation. Neutrophils with the proinflammatory allelic FcαR variant triggered a stronger activation response to IgA ANCA *in vitro*. Thus, the data indicate that FcγR and FcαR genotypes influence manifestation patterns and disease severity in patients with ANCA-induced vasculitis.

Post-translational modifications such as sialylation might be an additional mechanism to change the activating capability of ANCA. It has been shown that the PR3-ANCA sialylation ratio was significantly lower in patients with active disease correlating with the Birmingham Vasculitis Activity Score (BVAS) score. Moreover, the *in-vitro* respiratory burst was correlated inversely with sialylation of the PR3-ANCA IgG [51]. All these findings suggest an important interplay between the ANCA antigen-binding fragment, the Fc part with its isotype and class characteristics and post-translational ANCA modifications as well as important genetic variants in the corresponding Fcα and Fcγ receptors on the neutrophil that may determine the mechanisms and strength by which ANCA interact with the neutrophil. The bacterial enzyme endoglycosidase S resulted in hydrolysis of ANCA IgG glycans and attenuated ANCA-induced neutrophil activation necrotizing crescentic glomerulonephritis

(NCGN) in an anti-MPO antibody-mediated mouse model [52].

ANCA antigens exist in a larger signalling complex allowing neutrophil activation

MPO and PR3 are not transmembrane molecules, and therefore need to co-operate with other molecules to start intracellular signal transduction. Previous data using blocking antibodies had implicated β 2-integrins in ANCA-induced neutrophil activation [42]. David *et al.* characterized a direct interaction between PR3 and CD11b/CD18 (Mac-1) on the neutrophil membrane and suggested that PR3 modulates neutrophil adhesion by activating Mac-1 [53]. The same group described later that PR3 was present in lipid rafts together with the GPI-linked Fc γ RIIb and p22phox, an essential component of nicotinamide adenine dinucleotide phosphate-oxidase (NADPH) oxidase complex [54]. An interesting finding in their study was that using phospholipase D to cleave GPI-linkers resulted in a reduction of both PR3 and Fc γ RIIb, suggesting that a GPI-anchored receptor indeed mediates mPR3 presentation. As discussed above, the NB1 is also a GPI-linked protein and is a sufficient receptor for mPR3 presentation [23]. However, even knowing that PR3 exists in a complex with GPI-linked molecules could not explain how neutrophils become activated after PR3–ANCA binding, as none of these components have transmembrane domains. Precipitating CD177 from the neutrophil membrane and performing mass spectrometry, we found that several molecules co-precipitated with CD177. Among those proteins were the Fc γ RIIR as well as Mac-1 [55]. CD177 and Mac-1 co-localized, co-precipitated and showed direct protein interactions by plasmon-resonance analysis and when Mac-1 transfected cells interacted with immobilized NB1. We subsequently established that Mac-1 was a functionally important transmembrane component of the PR3 membrane complex, allowing subsequent PR3–ANCA-induced activation predominantly of mPR3^{high}/NB1^{positive} neutrophils (Fig. 2). However, we observed that degranulation and extracellular superoxide generation, but not intracellular hydrogen peroxide formation depended on the mPR3 phenotype. Interestingly, PR3–ANCA were equally potent in inducing DHR oxidation in mPR3^{high}/NB1^{positive} and mPR3^{low}/NB1^{negative} cells an observation also made by Hu *et al.* [27]. The underlying mechanism for this finding still needs to be elucidated.

As mentioned, MPO membrane expression by neutrophils is somewhat scarce and much less is known as to how signalling is initiated after MPO–ANCA bind their target. Hess *et al.* found that large amounts of MPO can be acquired by resting neutrophils from supernatants of activated neutrophils. This acquired surface MPO allowed MPO–ANCA binding and neutrophil activation [56]. Others showed that MPO is presented by CD11b promoting

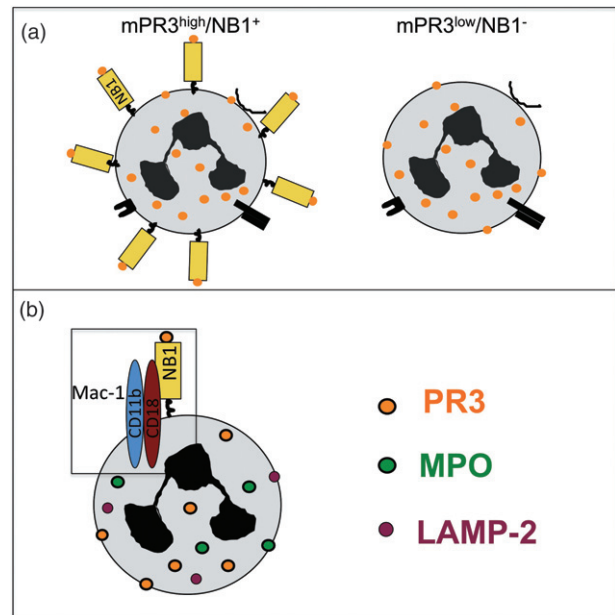


Fig. 2. Neutrophil antigen B1 (NB1) is a receptor for proteinase 3 (PR3) presentation on the neutrophil membrane. (a) A mPR3^{high}/NB1^{positive} and a mPR3^{low}/NB1^{negative} neutrophil subset exist. (b) PR3 and NB1 are part of a larger signalling complex in the lipid rafts. The transmembrane β 2-integrin Mac-1 is part of this complex and mediates PR3–anti-neutrophil cytoplasmic autoantibodies (ANCA)-induced neutrophil activation.

neutrophil activation even in the absence and presence of anti-MPO antibodies [57,58].

ANCA activate signalling pathways and neutrophil functions that participate in the vasculitic damage

Initial studies on ANCA-induced signalling events showed that distinct intracellular signalling events mediated ANCA-induced neutrophil activation. Tyrosine kinase and protein kinase C activation by ANCA, but not by control IgG, was observed by Radford *et al.* [59]. Blocking both kinases using pharmacological inhibitors abrogated ANCA-induced superoxide generation. These experiments encouraged further characterization of the signal transduction cascade involved in ANCA-induced neutrophil activation. The implication was to block important key elements specifically and thereby identify novel and more specific treatment targets.

p38 mitogen-activated protein kinase (MAPK) and extracellular regulated kinase (ERK) are important during both priming and the ANCA-induced neutrophil activation. Priming increases the amount of membrane-expressed antigens, but also sparks signalling pathways that are needed for a subsequent ANCA-induced full-blown activation. Both p38 MAPK and ERK are initiated during TNF- α priming and their blockade abrogates subsequent ANCA-induced activation. However, both pathways show differential effects in that p38 MAPK, but not ERK, controls

the ANCA-antigen translocation [60]. Interestingly, the 3-hydroxy-3-methylglutaryl co-enzyme A reductase inhibitors cerivastatin and simvastatin, which are known to exert actions above and beyond lipid lowering, abrogated ANCA-induced respiratory burst [61]. The effect was independent of the mevalonate pathway and involved ERK, but not p38 MAPK inhibition. Activated p38 MAPK was detected in glomerular neutrophils and intrinsic cells in biopsies from ANCA patients [62]. The importance of p38 MAPK for ANCA-induced NCGN was demonstrated recently in a disease mouse model [63] and is discussed in the adjacent review by Robson.

Several studies explored the role of phosphatidylinositol 3-kinase (PI3K) in ANCA-induced neutrophil activation. PI3K generates phosphatidylinositol-3,4,5-triphosphate (PIP3) and phosphatidylinositol-3,4-diphosphate (PIP2). Both substances recruit the serine/threonine kinase Akt. Ben-Smith *et al.* observed that ANCA induced PIP3, but did not activate p85/p110 PI3K. This PI3K isoform was, however, activated by simple Fc γ R cross-linking, again underscoring the fact that ANCA-induced activation is not merely a consequence of Fc γ R cross-linking and that other transmembrane molecules are required [64]. In contrast, ANCA activated the p101/110 γ PI3K. Inhibition of all PI3K isoforms by LY294002 blocked ANCA-triggered superoxide generation. We confirmed the functional importance of PI3K. In addition, we investigated activation of the downstream kinase Akt by ANCA. Akt is phosphorylated by phosphoinositide-dependent kinase 1 (PDK1) and by PDK2. P38 MAPK can function as PDK2, and we showed that both the p38 MAPK and PI3K participate in Akt activation by ANCA [65]. TNF- α priming also resulted in Akt phosphorylation by both upstream kinases and promoted the association of Akt with the actin regulatory protein PAK1. ANCA patients frequently suffer from febrile infections that complicate immunosuppressive therapy. During these events neutrophils are exposed to increased temperatures. Antipyretics are distributed generously to fight fever, although its biological role is not so clear. We observed two interesting effects of short fever-like temperature spikes on neutrophils that could be clinically relevant in ANCA patients. Heat exposure abrogated PI3K/Akt activation and respiratory burst in primed neutrophils challenged by ANCA [66]. ANCA-induced phosphorylation of p38 MAPK and ERK was not affected. However, heat exposure prevented the increase in ANCA antigen expression in neutrophils that were treated with lipopolysaccharide (LPS) overnight [67]. This effect was mediated, at least in part, by diminishing TNF- α that was released from LPS-treated neutrophils. TNF- α required p38 MAPK to up-regulate ANCA antigen expression on the neutrophil surface, and heat accelerated p38 MAPK protein degradation in LPS-treated neutrophils. These data suggest that fever-like temperatures could modulate ANCA-mediated inflammatory responses via PI3K/Akt and p38 MAPK pathways.

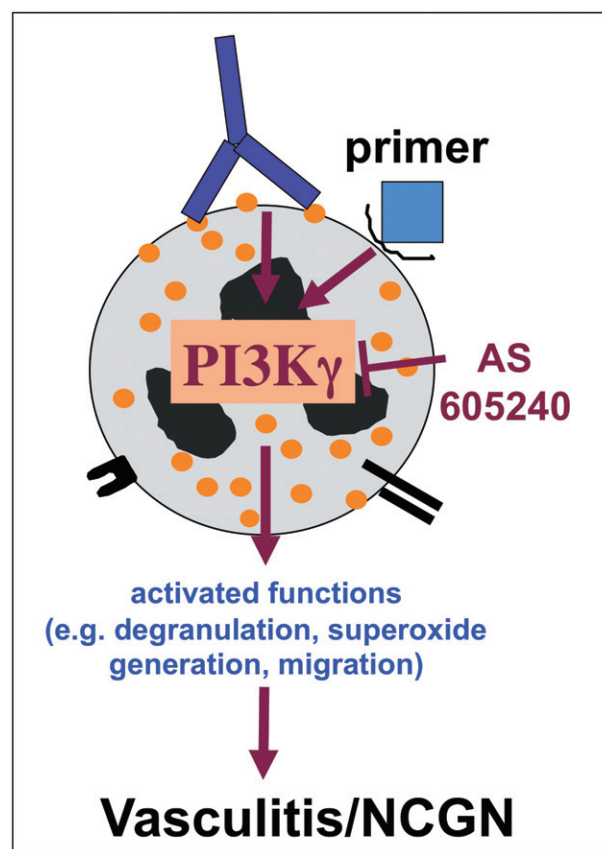


Fig. 3. Anti-neutrophil cytoplasmic autoantibodies (ANCA)-induced neutrophil activation is controlled by intracellular signalling pathways. phosphatidylinositol 3-kinase γ (PI3K γ) is depicted as an example for a signal molecule that was characterized as a key molecule for ANCA-induced neutrophil activation. PI3K γ can be targeted therapeutically to prevent necrotizing crescentic glomerulonephritis (NCGN).

The functional role of the PI3K γ isoform was confirmed *in vivo* by a genetic approach in a murine bone-marrow (BM) transplantation model of anti-MPO antibody-induced NCGN [68]. NCGN occurred in mice that had received BM from wild-type, but not from PI3K γ gene-deleted mice. Moreover, a γ isoform-specific inhibitor abrogated ANCA-induced superoxide generation, degranulation and neutrophil migration *in vitro* and oral treatment with this compound prevented NCGN in mice, suggesting that specific PI3K γ inhibition could be used therapeutically (Fig. 3).

Several investigators have now implicated the participation of complement activation in ANCA-induced inflammation. In fact, animal studies narrowed the alternative pathway and particularly C5 as an important component in ANCA-induced NCGN [69,70]. *In-vitro* experiments elucidated that C5a is generated by ANCA-activated neutrophils and that this component further provides additional neutrophil priming for ANCA activation. Thus, ANCA-induced

C5a would then act as an acceleration loop, further enhancing inflammation. C5a is connected to the important PI3K pathway in that the C5a receptor belongs to the G protein-coupled receptors that signal via PI3K γ [71]. Importantly, mice lacking the C5a receptor in myeloid cells only were protected from anti-MPO antibody-induced NCGN [6]. These data imply that the C5a receptor may provide an additional treatment target in patients with ANCA vasculitis.

ANCA stimulation induces neutrophils and monocytes to produce and release cytokines [44,72–74]. Proinflammatory IL-1 β may be of particular clinical interest because it is increased by ANCA, the lack of IL-1 β R in renal cells protected from glomerular injury in murine anti-GBM model and an IL-1R blocker is available in the clinic [72,75,76]. Active IL-1 β is generated from inactive precursor pro-IL-1 β . The classical enzyme that mediates this process is caspase-1. Alternative IL-1 β converting enzymes were suggested. We showed recently that active neutrophil serine proteases (NSPs) are critical for IL-1 β generation in ANCA-stimulated monocytes and neutrophils. The IL-1 β amount produced by monocytes was clearly higher compared to neutrophils, but neutrophils outnumber monocytes *in vivo*, suggesting that both cell types are possibly important. Murine monocytes and neutrophils lacking dipeptidylpeptidase I (DPPI) and therefore lacking active NSPs produced significantly less IL-1 β in response to anti-MPO antibodies [77]. Preincubation of human monocytes with cell-permeable serine protease inhibitors or a caspase-1 inhibitor also diminished IL-1 β generation. NSPs consist of human neutrophil elastase (HNE), PR3 and cathepsin G (CG). Exogenous PR3 rescued IL-1 β generation in DPPI-deficient monocytes. DPPI- and PR3/HNE-deficient myeloid cells as well the IL-1R blocker Anakinra protected from NCGN in an anti-MPO antibody-mediated NCGN mouse model. These findings demonstrate that at least two mechanisms participate in IL-1 β generation, namely caspase-1 and PR3, and that PR3 alone or in combination with HNE is important for ANCA-induced NCGN. It is conceivable that additional NSP-mediated effects participate in ANCA-mediated tissue damage. These proteases may cleave extracellular matrix proteins and injure the endothelium. Lu *et al.* demonstrated that ANCA-activated neutrophils released serine proteases, but not superoxide when co-cultured with EC, and that serine proteases mediated EC damage resulting in von Willebrand factor (vWF) release [78]. Serine proteases that are packaged in ANCA-induced neutrophil microparticles or in neutrophil extracellular traps (NETs) possibly also participate in endothelial damage [79,80].

Together, ANCA induce a variety of neutrophil responses *in vitro*. Some of these were shown to be significant *in vivo*, such as p38 MAPK, PI3K γ , C5a and serine proteases. Others that are thought to be important await further *in-vivo* proof, including the role of ANCA-induced reactive oxygen generation.

Conclusion

The neutrophil is both the cell that expresses target ANCA antigens and a major effector cell in ANCA-induced small vessel vasculitis. The ANCA antigens PR3 and MPO differ substantially in their expression pattern on the neutrophil plasma membrane. ANCA bind to membrane expressed target antigens and initiate intracellular signalling events. The PR3–NB1–Mac-1 membrane complex is one example showing that larger signalling complexes with transmembrane molecules exist. Distinct signalling pathways triggered by ANCA F(ab)2 and the intact ANCA IgG molecule were identified and co-operate in neutrophil activation. Detailed characterization of the activation process will identify novel treatment targets that need to be tested in animal models and subsequently in patients.

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Disclosure

Nothing to declare.

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