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Monitoring Human Neutrophil Activation by a Proteinase 3 Near-Infrared Fluorescence Substrate-Based Probe

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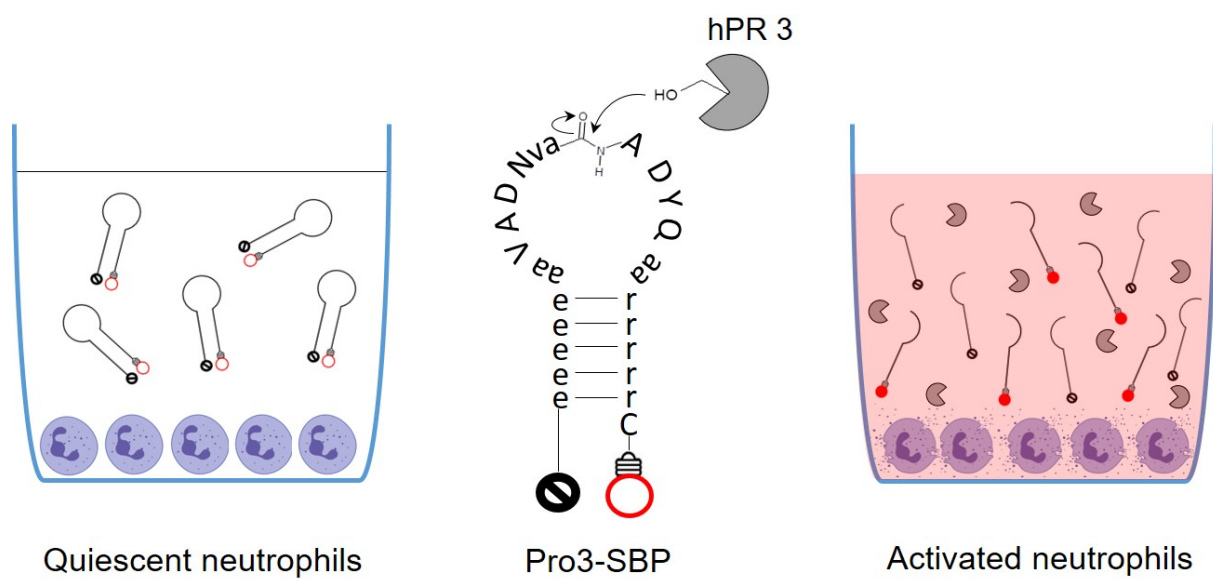
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Monitoring of human neutrophil activation by a proteinase 3 near-infrared fluorescence substrate-based probe

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Complete List of Authors:	Saidi, Ahlame; Université de Tours, INSERM UMR1100 - Centre d'études des pathologies respiratoires (CEPR) Wartenberg, Mylène; Université de Tours, INSERM UMR1100 - Centre d'études des pathologies respiratoires (CEPR) Madinier, Jean-Baptiste; CBM, Centre de Biophysique Moléculaire Ilango, Guy; Université de Tours Seren, Seda; Université de Tours Korkmaz, Brice; Université de Tours, INSERM UMR1100 - Centre d'études des pathologies respiratoires (CEPR) Lecaille, Fabien; Université de Tours, INSERM UMR1100 - Centre d'études des pathologies respiratoires (CEPR) AUCAGNE, Vincent; CBM, Centre de Biophysique Moléculaire Lalmanach, Gilles; Université de Tours, INSERM UMR1100 - Centre d'études des pathologies respiratoires (CEPR)

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10 **Ahlame Saidi,^{†, ‡, §} Mylène Wartenberg,^{†, ‡, §} Jean-Baptiste Madinier,[¶] Guy Ilango,[#]**
11
12 **Seda Seren,^{†, ‡, f} Brice Korkmaz,^{†, ‡} Fabien Lecaille,^{†, ‡} Vincent Aucagne,[¶] and**
13
14 **Gilles Lalmanach^{†, ‡, *}**
15
16
17
18
19

20 [†] Université de Tours, Tours, 37032, France

21
22 [‡] INSERM, UMR 1100, Research Center for Respiratory Diseases (CEPR), Team: "Proteolytic
23 Mechanisms in Inflammation", Tours, 37032, France

24
25 [¶] CNRS UPR 4301, Center for Molecular Biophysics (CBM), Team: "Molecular, Structural and
26 Chemical Biology", Orléans, 45071, France

27
28
29 [#] IBiSA Electron Microscopy Platform, Université de Tours, Tours, 37032, France

30
31
32
33 [§] AS & MW: Equal contribution to the work

34
35
36
37 ^f Current address: Centre de Résonance Magnétique des Systèmes Biologiques, UMR 5536
38 CNRS, Université de Bordeaux, Bordeaux, France.

39
40
41 ^{*} Corresponding author : Prof. Gilles Lalmanach, INSERM UMR1100, Centre d'Etude des
42 Pathologies Respiratoires (CEPR), Equipe « Mécanismes protéolytiques dans l'inflammation »,
43 Université de Tours, Faculté de Médecine, 10 Boulevard Tonnellé, 37032 Tours cedex, France.
44 Tel: (+33) 2 47 36 61 51. Mail: gilles.lalmanach@univ-tours.fr
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Abstract

A near-infrared fluorescent (NIRF) substrate-based probe (SBP) was conceived to monitor secreted human proteinase 3 (hPR3) activity. This probe, called pro3-SBP, is shaped by a fused peptide hairpin loop-structure, which associates a hPR3 recognition domain (Val-Ala-Asp-Nva-Ala-Asp-Tyr-Gln, with Nva: norvaline) and an electrostatic zipper (consisting of complementary polyanionic (D-Glu)₅ and polycationic (D-Arg)₅ sequences) driving a tight vicinity of the N- and C- terminal FRET couple (fluorescent donor: sulfoCy5.5; dark quencher: QSY21). Beside its subsequent stability, no intermolecular fluorescence quenching was detected following its complete hydrolysis by hPR3, advocating that pro3-SBP could further afford unbiased imaging. Pro3-SBP was specifically hydrolyzed by hPR3 ($k_{\text{cat}}/K_m = 440,000 \pm 5,500 \text{ M}^{-1} \cdot \text{s}^{-1}$) and displayed a sensitive detection threshold for hPR3 (sub-nanomolar concentration range), while neutrophil elastase showed a weaker potency. Conversely, pro3-SBP was not cleaved by cathepsin G. Pro3-SBP was successfully hydrolyzed by conditioned media of activated human neutrophils, but not by quiescent neutrophils. Moreover, unlike unstimulated neutrophils, a strong NIRF signal was specifically detected by confocal microscopy following neutrophil ionomycin-induced degranulation. Fluorescence release was abolished in the presence of a selective hPR3 inhibitor, indicating that pro3-SBP is selectively cleaved by extracellular hPR3. Taken together, present data support that pro3-SBP could be a convenient tool allowing a straightforward monitoring of human neutrophil activation.

Keywords

Cathepsin, Elastase, Near-infrared fluorescence (NIRF), Neutrophil serine protease (NSP), Proteinase 3

Introduction:

Neutrophil elastase (NE), proteinase 3 (PR3), and cathepsin G (CatG) are three neutrophil serine proteases (NSPs) that are stored in the primary (azurophil) granules of polymorphonuclear neutrophils (PMNs).^{1,2} NSP4, a less abundant NSP which is also uncovered in the azurophilic granules, was recently identified.^{3,4} Maturation of NSPs occurs through cleavage of a N-terminal dipeptide by cysteine cathepsin C.⁵ NSPs are implicated in a variety of diseases, including ischemic-reperfusion injury, glomerulonephritis, arthritis, bullous pemphigoid, non-alcoholic fatty liver disease or lung diseases (COPD, ARDS, cystic fibrosis).⁶ NSPs have bactericidal and antifungal properties, and were first recognized as digestive enzymes that degrade ingested host pathogens.⁷ A web-like structure of DNA secreted by activated neutrophils (neutrophil extracellular trap, NET) was described as an additional antimicrobial mechanism.⁸ Following recruitment of PMNs, NSPs participate in the alteration of extracellular matrix components at inflammatory sites, and are key regulators of the local inflammatory response, being involved in the control of cellular signaling through the processing of chemokines and the modulation of the cytokine network.²

PR3 may be detected both in secondary granules and on membrane-bound secretory vesicles.⁹ Mature PR3 is dormant in the acidic environment of the azurophilic granules and is activated upon translocation to a neutral pH environment by a slow conformational change. PR3 is primarily regulated by α 1-protease inhibitor, serpin B1, and elafin, but not by secretory leukocyte protease inhibitor.¹⁰ Under pathological conditions, downregulation and/or inactivation of PR3 inhibitors result in an uncontrolled PR3 activity, which may further contribute to tissue damage and inflammatory cascades. More specifically, PR3 participates in the activation of IL-8, TNF- α and IL-1 β as well the degradation of fibronectin or laminin, and is involved in the maturation of cathelicidin hCAP-18 to generate LL-37, a potent antimicrobial peptide.¹¹ Also, PR3 favors endothelial cell apoptosis and takes part in platelet activation and regulation of granulopoiesis.¹²⁻¹⁴ Furthermore, PR3 is the main target antigen of anti-neutrophil cytoplasmic autoantibodies (ANCA) in granulomatosis with polyangiitis (GPA).¹⁵ Structural and functional properties of human PR3 (hPR3) were initially described as similar to those of human NE (hNE). hPR3 and hNE share an analogous catalytic mechanism¹⁶ and the elucidation of the crystal structure of hPR3 has confirmed that both enzymes display a common fold.^{17,18} Despite hPR3 and hNE preferentially accommodate small hydrophobic residues in the S1 pocket, their different substrate specificities have been exploited to develop synthetic inhibitors and FRET peptidyl substrates of human PR3.¹⁹⁻²²

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3 Although hPR3 is considered as a relevant therapeutic target, its precise contribution to
4 some diseases remains still debated. Accordingly, the appraisal of hPR3 activity within living
5 cells and extracellular milieu, using newly developed chemical biology tools, will obviously
6 assist to clarify the role of hPR3 during pathophysiological events. Foremost strategies of probe
7 design to profile enzymatic activity of serine and cysteine proteases, from cell to *in vivo* assays,
8 were recapitulated elsewhere.^{23–25} Activity-based probes (ABPs) consist of an electrophilic
9 warhead that binds stoichiometrically and covalently to the catalytic nucleophilic residue of the
10 active site, coupled to a linker or a targeting sequence, and a reporter group for visualization,
11 thus allowing to accurately pinpoint the protease activity. During the last decade, potent
12 chemical probes have been developed for assessment of NSP activities, including hPR3
13 activity.^{26–34} However, ABPs do not permit signal amplification, which can result in low-
14 intensity signals. By contrast, substrate-based probes (SBPs) contain a peptidyl (or pseudo-
15 peptidyl) backbone linked to a tag (usually a fluorophore) and a quencher group, allowing a
16 stronger reporter signal amplification. A detrimental counterpart is that, after cleavage, released
17 low molecular weight reporters could diffuse faster than an ABP-protease covalent complex
18 and thus lead to unfocused signals or images.^{35,36}

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31 A recent article reported a straightforward method for monitoring hNE and CatG
32 activity of purified neutrophils, but not hPR3 activity.³⁷ Alternatively, methods for monitoring
33 hPR3 are currently based on measurements of its enzymatic activity by peptidyl substrates.³⁸
34 Given that the extracellular activity of PR3 is basically associated to stimulation of PMNs and
35 release of NSPs stored in primary granules, an attractive manner to survey activation of
36 neutrophils is to design and synthesize a near-infrared fluorescence SBP. Hence, we conceive a
37 cell-impermeable near-infrared fluorescence (NIRF) SBP for monitoring active, secreted
38 hPR3.³⁹ We therefore designed a peptide hairpin loop-like structure that consists of a hPR3
39 recognition sequence and an electrostatic zipper, driving a close vicinity of the N- and C-
40 terminal FRET couple (fluorescent donor: sulfoCy5.5; non-fluorescent dark quencher: QSY21).
41 The recognition sequence is derived from both a hPR3 substrate (Abz-Val-Ala-Asp-Nva-Ala-
42 Asp-Tyr-Gln-EDDnp, with Nva: norvaline) and azapro-3, its corresponding azapeptide
43 inhibitor (Abz-Val-Ala-Asp-azaNva-Ala-Asp-Tyr-Gln-3-nitro-Tyr).⁴⁰ First, the probe, called
44 pro3-SBP, was characterized *in vitro*. Then, the ability of pro3-SBP to monitor secreted hPR3
45 and thus neutrophil activation was endorsed. Despite a scattered fluorescence signal, a pivotal
46 advantage of pro-SBP3, which emits in the near infrared, is to provide a sensitive detection
47 level of hPR3, allowing a laid-back and fast visualization of neutrophil activation.

Results & discussion

Design and characterization of the proteinase 3 NIRF probe

The hPR3 recognition domain was derived from both a peptidyl substrate and an azaNva-containing peptide inhibitor initially designed for targeting neutrophil hPR3⁴⁰ (see schematic representation: Figure 1a). The electrostatic zipper corresponded to a short double strand consisting of a complementary pair of a polyanionic (D-Glu)₅ segment and a (D-Arg)₅, a 5-mer polycation. The zipper was covalently linked to the N- and C-termini of hPR3 recognition peptide via (D-Ala)₂ spacers. The fluorescence donor-acceptor pair consisted of i/ a dark quencher (QSY21) conjugated to the peptide through an amide bond with the N-terminal amino group of D-Glu and ii/ a maleimido-functionalized fluorescent dye (sulfoCy5.5) which was linked to a C-terminal Cys via a thioether bond. The presence of the double strand polycationic/polyanionic zipper drives a tight vicinity between the fluorescent donor group and QSY21, the non-fluorescent acceptor quencher, leading to an efficient intramolecular FRET in the absence of proteolytic cleavage. Consequently, QSY21 returns from the excited state to the ground state via non-radiative decay pathways, without the emission of light.⁴¹ While cationic Arg-rich peptides as well poly-Arg 8 to 15-mers translocate through membranes and may act as potent CPPs^{42,43}, peptides containing less than six L/D -Arg residues are ineffective CPPs.⁴⁴ Accordingly, presence of a double strand electrostatic zipper consisting of only five D-Glu/D-Arg pairings ensure the inability of NIRF probe to cross membranes, avoiding any targeting of intracellular hPR3.⁴⁵ A similar strategy was successfully used to design a selective NIRF probe of secreted CatS.⁴⁶

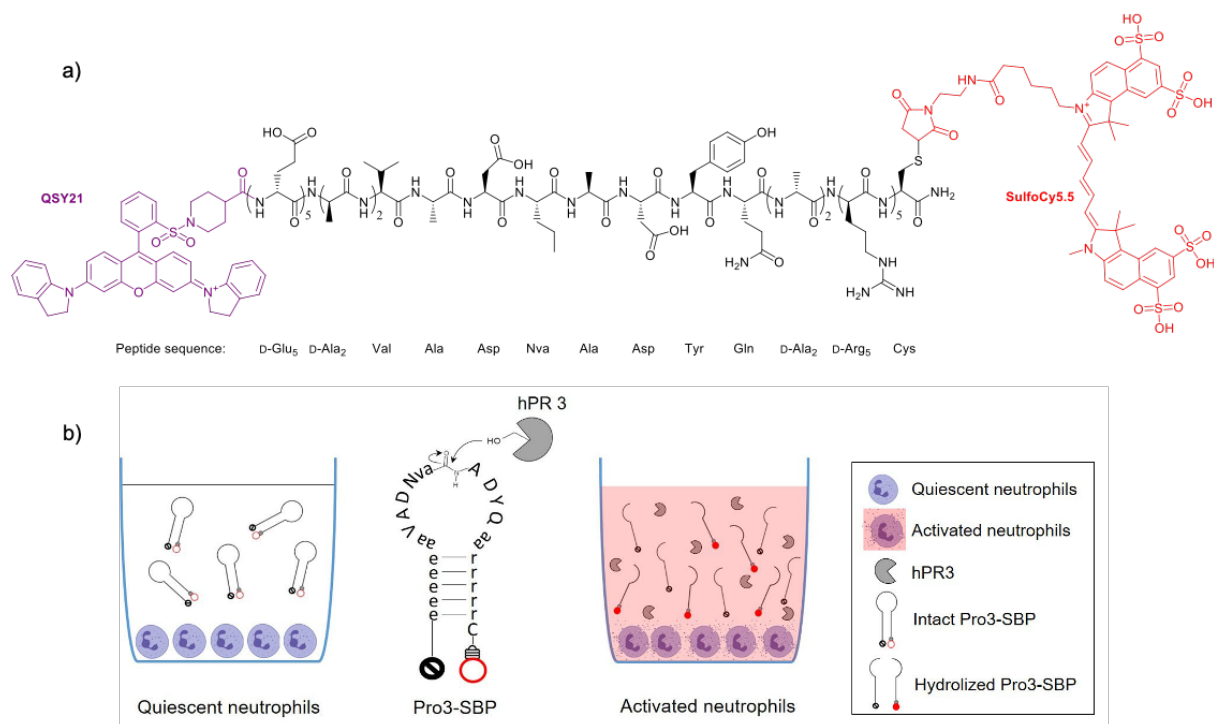


Figure 1: structure and design of the hPR3 substrate-based probe.

(a) Structure of the NIRF probe (pro3-SBP). The PR3 recognition domain (VADXADYQ, with X=norvaline, Nva) was derived from a selective azaNva-containing peptide inhibitor of human neutrophil PR3, hPR3. The electrostatic zipper corresponds to a complementary (D-Glu)₅/(D-Arg)₅ pair. The PR3 recognition peptide is flanked in both N- and C-positions by a (D-Ala)₂ spacer. The FRET donor-acceptor pair consists of a dark quencher (QSY21) and a maleimido-functionalized fluorophore (sulfoCy5.5) which is linked to a C-terminal Cys. (b) Cartoon representation. General scheme for monitoring neutrophil activation by pro3-SBP.

Pro3-SBP was synthesized as detailed extensively in the experimental part. The characteristics of the purified probe (RP-HPLC profile, ESI-MS spectrum and UV-vis spectrum) were disclosed in the supplementary file: Figure S1. Stability of the intramolecularly quenched fluorescent pro3-SBP was further examined. No fluorescence signal corresponding to a self-dequenching ($\lambda_{\text{ex}}=684\text{ nm}$; $\lambda_{\text{em}}=710\text{ nm}$) was observed after one hour of incubation, supporting that the intramolecular FRET between sulfoCy5.5 and QSY21 was robust. This was confirmed by preincubation of pro3-SBP in the presence of a high ionic strength (1 M NaCl) where no dequenching was observed. Alternatively, addition of hPR3 led to a straight fluorescence release, indicating that the cleavage within the recognition sequence triggered an immediate dissociation of the salt bridges between (D-Glu)₅ and (D-Arg)₅ strands, directing the dequenching of the sulfoCy5.5 donor/QSY21 acceptor pair, as previously described for a molecular SBP conjugate of similar assembly, but carrying a different donor/acceptor pair.⁴⁶ Moreover, cleavage of increasing amounts of pro3-SBP (0-8 μM) conducted a linear dose-

dependent level of emitted fluorescence ($R^2 = 0.98$), demonstrating that intermolecular fluorescence quenching did not occur.

Specificity of the hPR3 substrate-based probe

As indicated before, incubation of pro3-SBP with hPR3 conducted to a NIRF release. As expected, the cleavage site was located at the Nva-Ala bond (MALDI mass spectrometry analysis) (Figure S2). The second-order rate constant, which was determined under pseudo-first order conditions, corresponded to $k_{cat}/K_m = 440,000 \pm 5,500 \text{ M}^{-1} \cdot \text{s}^{-1}$ (with Michaelis constant $K_m = 1.33 \mu\text{M}$), supporting a worthy specificity (Table I).

Table I: Kinetic parameters (second-order rate constant) for the hydrolysis of pro3-SBP by human proteases.

Proteases	$k_{cat} / K_m (\text{M}^{-1} \cdot \text{s}^{-1})$
hPR3	$440,000 \pm 5,500$
CatG	n.h.
hNE	$33,000 \pm 3,000$
CatS	n.h.
CatL	$<1,000$
CatB	n.h.
CatH	n.h.
CatD	n.h.
MMP-8	n.h.
MMP-9	n.h.
MMP-12	n.h.

Determination of the second-order rate constants were carried out under pseudo first-order conditions as detailed in the experimental section. Kinetic data were reported as means \pm S.D ($n=3$, three independent experiments). n.h., no hydrolysis.

Also, a similar specificity constant was maintained, compared to the parent FRET substrate (Abz-Val-Ala-Asp-Nva-Ala-Asp-Tyr-Gln-EDDnp)⁴⁰, sustaining that substitution of the Abz/EDDnp pair by a bulky hydrophobic donor/quencher (sulfoCy5.5/QSY21) pair did not affect the k_{cat}/K_m value, which mostly relied on the peptidyl hPR3 recognition pattern. Pro3-SBP was also cleaved, but to a lesser extent, by hNE ($k_{cat}/K_m = 33,000 \pm 3,000 \text{ M}^{-1} \cdot \text{s}^{-1}$). Analysis of the hydrolysis products by mass spectrometry demonstrated the presence of two cleavage sites for hNE, located at the Nva-Ala bond and its neighboring Ala-Asp bond (respective cleavage ratio: $\sim 55\%$ vs $\sim 45\%$) (Figure S3). Also, cathepsin L hydrolyzed slightly pro3-SBP (estimated $k_{cat}/K_m < 1,000 \text{ M}^{-1} \cdot \text{s}^{-1}$). On the other hand, pro3-SBP was not susceptible to

hydrolysis by CatG. Similarly, biologically relevant aspartic CatD, cysteine cathepsins B, H and S, MMP-8 and MMP-9, which are matrix metalloproteinases produced by neutrophils, as well as MMP-12 (macrophage elastase) did not hydrolyze pro3-SBP.

We evaluated the sensitivity (i.e. detection limit) of pro3-SBP. In the presence of a constant amount of pro3-SBP (0.5 μM), fluorescence release was observed with subnanomolar concentrations of hPR3 (IVIS Caliper Lumina XR imaging system) (Figure 2a). Alternatively, in the presence of a constant concentration of hPR3 (5 nM), we detected a fluorescence threshold with submicromolar concentration (0.25 μM) of pro3-SBP (Figure 2b).

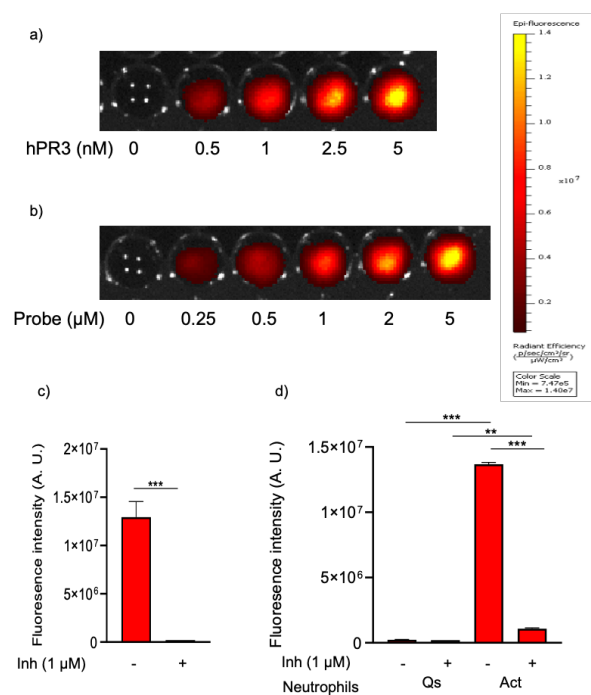


Figure 2: Detection threshold and selectivity of pro3-SBP.

(a) Increasing amounts of hPR3 (0-5 nM) were incubated in the presence of pro3-SBP (0.5 μM) (0.05 M HEPES buffer, pH 7.4, 0.05% NP40, 150 mM NaCl). After 5 min, fluorescence release was measured ($\lambda_{\text{ex}}=640$ nm, $\lambda_{\text{em}}=675$ nm; 5 s exposure; IVIS Caliper Lumina XR imaging system). (b) PR3 (5 nM) was incubated in the presence of increasing concentrations of pro3-SBP (0-5 μM), before recording the fluorescence release ($t=25$ min). (c) Control experiments were conducted by pre-incubating for 30 min PR3 (5 nM) in the presence of a selective irreversible hPR3 inhibitor (Inh: Biotinyl-Val-Tyr-Asp-Nva-P(O-C₆H₄-4-Cl)₂, 1 μM). After addition of pro3-SBP (5 μM), fluorescence release was monitored (IVIS Caliper Lumina XR imaging system, $t=10$ min). Data were expressed as mean \pm SD fluorescence intensity (A.U., arbitrary units). Statistical analyses were performed using Mann-Whitney U test ($n=3$, ***: $P<0.0005$). (d) Following incubation of neutrophils with ionomycin, as detailed in the experimental section, conditioned media of activated neutrophils (Act) were withdrawn and incubated with pro3-SBP (5 μM). Fluorescence release was further examined (IVIS Caliper Lumina XR imaging system, $t=30$ min). The same experiments were carried out in the presence of unstimulated neutrophils (quiescent, Qs). Results were expressed as mean \pm SD fluorescence intensity (A.U., arbitrary units). Statistical analyses were performed using Kruskal-Wallis test ($n=3$, **: $P<0.005$, ***: $P<0.0005$). Signal selectivity was checked in the presence of the hPR3 inhibitor (Inh).

The fluorescence release correlated markedly with hPR3 enzymatic activity, since there was a full impairment of signal in the presence of Biotinyl-Val-Tyr-Asp-Nva-P(O-C₆H₄-4-Cl)₂ (Figure 2c), a potent peptidyl di(chlorophenyl)-phosphonate ester inhibitor, previously used to

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3 selectively inhibit extracellular activity of hPR3 in sputa from cystic fibrosis patients.³¹
4 Following treatment of purified neutrophils by a calcium ionophore, PMN conditioned medium
5 was collected and pro3-SBP was added. An extensive NIRF signal, which was abolished in the
6 presence of Biotinyl-Val-Tyr-Asp-Nva-P(O-C₆H₄-4-Cl)₂, was monitored (Figure 2d).
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8 Conversely to ionomycin-activated neutrophils, hydrolysis of pro3-SBP was undetectable in
9 supernatants issued from untreated PMNs. Present results indicate that the specific fluorescence
10 release from cleavage of pro3-SBP corresponded to extracellular hPR3 activity, which could
11 be entirely abrogated by addition of the hPR3 inhibitor.³¹ Data also support that pro3-SBP could
12 be a valuable tool for monitoring the secretion of mature active hPR3 by activated neutrophils,
13 since no NIRF signal was detected in the presence of quiescent neutrophils.
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22 **Imaging of secreted hPR3 by confocal microscopy**

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24 Both ionomycin-treated and untreated neutrophils were cultured for 4 h. After removal
25 of non-adherent PMNs, pro3-SBP (10 μ M) was immediately added to the culture medium and
26 imaging was performed by confocal fluorescence microscopy. In a preliminary way, a cell
27 viability assay was performed since incubation of neutrophils with ionomycin results in an
28 increase of intracellular Ca²⁺ levels and a subsequent decrease of cell viability.^{47,48} Accordingly,
29 we observed a decrease of cell viability following addition of ionomycin compared to untreated
30 neutrophils. However, the addition of pro3-SBP did not induce any significant modification of
31 the viability of ionomycin-activated neutrophils as well quiescent cells (Figure S4), which
32 supports that pro3-SBP did not exhibit deleterious cellular outcomes under the present
33 experimental conditions. Confocal microscopy analysis showed that NIR fluorescence release
34 was rapidly detected for activated neutrophils, but not for quiescent cells (Figure 3a-b). The
35 emitted fluorescence was found exclusively in the extracellular medium, but not intracellularly
36 within living neutrophils, as supported by bright-field microscopy, thus validating our probe
37 design strategy. In parallel, control trials, cell media of both quiescent, and activated PMNs
38 were withdrawn and hPR3 activity was measured by transmitted light microscopy. Addition of
39 an irreversible phosphonate hPR3 inhibitor²⁶ fully abrogated the fluorescence release (Figure
40 3c), confirming that pro3-SBP is a potent and selective SBP for hPR3 imaging following
41 neutrophil activation.
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55 Present data confirmed that the peptide hairpin loop-structure of pro3-SBP is highly
56 stable in culture media. Indeed, no fluorescence signal, associated to a spontaneous self-
57 dequenching, was detected for quiescent neutrophils in the absence of hPR3 release, supporting
58 the selective character of pro3-SBP. Moreover, data endorsed that pro3-SBP is a potent probe,
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aiming to image secreted/extracellular hPR3 activity following activation of quiescent neutrophils and degranulation by calcium ionophore. Nevertheless, a limitation for the use of pro3-SBP is that the released reporter group bearing Cy5.5 led to an unfocused signal, which is due to its diffusion into the extracellular medium, consistently with its substrate-derived backbone.³⁵ Nevertheless, a fine NIRF signal-noise ratio should be noticed, as demonstrated by the lack of fluorescence background of uncleaved pro3-SBP. Also, the worthy turnover rate constant value of pro3-SBP ($k_{\text{cat}}/K_m = 440,000 \pm 5,500 \text{ M}^{-1} \cdot \text{s}^{-1}$) thereby makes possible to detect *in situ* nanomolar concentrations of extracellular active hPR3.

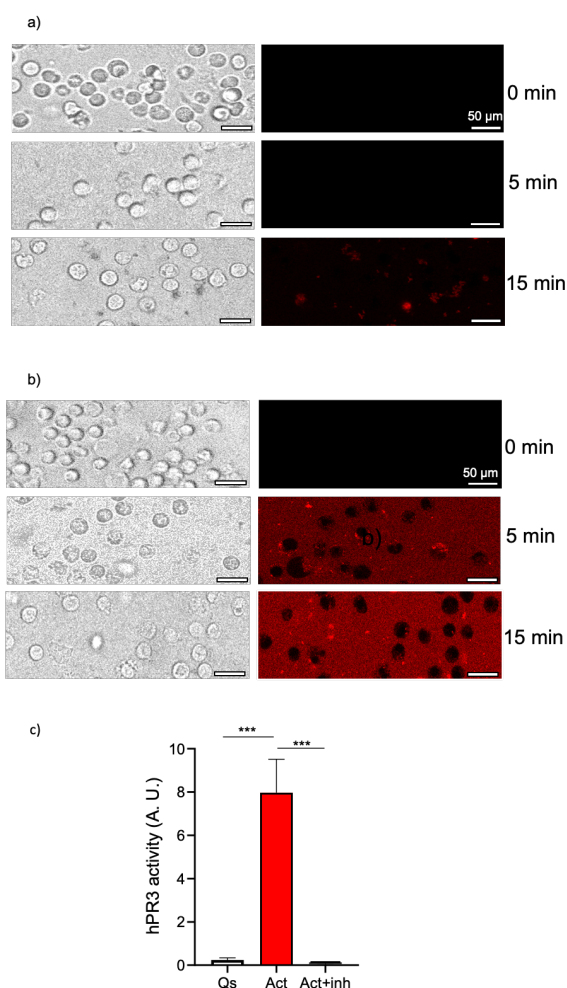


Figure 3: Probing neutrophil activation by confocal microscopy.

(a) Ionomycin-treated neutrophils were seeded in Well Black ibiTreat microplates (Ibidi) as described in “material & methods” (500,000 neutrophils/well). After removal of non-adherent cells, pro3-SBP (10 μM) was immediately added to the culture medium. Imaging was performed by monitoring the NIRF signal release (confocal fluorescence microscopy, Leica SP8 confocal microscope, laser 675 nm, red) at 0, 5, and 15 min (right panel), using white light as control (left panel). All snapshots were obtained by using objective: 60 \times and scale bar: 50 μm . (b) The selectivity of the NIRF signal was confirmed by carrying out the same experiments in the presence of quiescent neutrophils. (c) hPR3 activity in cell-free supernatants of quiescent (Qs) and activated (Act) neutrophils. Alternatively, media (corresponding to 500,000 neutrophils/well) were withdrawn, and hPR3 activity was measured using ABZ-Val-Ala-Asp-Nva-Ala-Asp-Tyr-Gln-EDDnp (20 μM) as substrate ($\lambda_{\text{ex}}=320 \text{ nm}$, $\lambda_{\text{em}}=420 \text{ nm}$). Biotinyl-Val-Tyr-Asp-Nva-P(O-C₆H₄-4-Cl)₂ (1 μM), a selective hPR3 inhibitor was used as control

(Act+inh). Data were expressed as arbitrary units (A.U). Results were analyzed using Kruskal-Wallis test (n=3, *P*-value ***; *P*<0.0005).

In conclusion, current results support that pro3-SBP is a specific NIRF substrate-based probe for monitoring activity of secreted hPR3. In addition to selective and powerful activity-based probes designed by several colleagues to specifically target and visualize hPR3 activity, pro3-SBP may represent an alternative and convenient molecular tool for a casual monitoring of neutrophil activation.

Experimental procedures

Peptide chemistry:

General information: All reagents and solvents were used without further purification. Protected amino acids were purchased from Gyros Protein Technology (Uppsala, Sweden). DIEA (di-*isopropyl*-ethylamine), HPLC gradient-grade MeCN and TFA were purchased from Carlo Erba (Val-de-Reuil, France). Rink amide ChemMatrix resin was purchased from Biotage (Uppsala, Sweden). Sulfo-cyanine5.5 NHS ester was obtained from Lumiprobe GmbH (Hannover, Germany). QSY21 maleimide was purchased from AAT Bioquest (Sunnyvale, CA, USA). Peptide synthesis-grade DMF was purchased from VWR (Fontenay-sous-Bois, France). Ultrapure water was obtained using a Milli-Q water system from Millipore (Molsheim, France). Polypropylene syringes fitted with polypropylene frits were purchased from Torviq (Niles, MI, USA) and were equipped with PTFE (5 polytetrafluoroethylene) stopcock from Biotage.

Peptide characterization and purification: HPLC analyses were carried out on a Chromaster system equipped with a 5160 pump, a 5430 diode array detector and a 5260 auto sampler. Semi-preparative purifications were carried out on a LaChromElite system equipped with a Hitachi L-2130 pump, a Hitachi L-2455 diode array detector and a Hitachi L-2200 auto sampler. Chromolith High Resolution RP-18e (150 Å, 10 × 4.6 mm, 3 mL/min flow rate) columns were used for analysis, and Nucleosil C18 (300 Å, 5 μm, 250 × 10 mm, 3 mL/min flow rate) for purification. As mobile phases, mixtures of 0.1% TFA in H₂O (A) and 0.1% TFA in MeCN (B) were used. ESI-MS analyses were carried out on an Agilent 1260 Infinity HPLC system, coupled with an Agilent 6120 mass spectrometer (ESI + mode), and fitted with an Aeris Widespore XB-C18 2 (3.6 μm, 150 × 2.1 mm, 0.5 mL/min flow rate, 60 °C) column. The reported *m/z* values correspond to the average mass. As mobile phases, mixtures of 0.1% formic acid in H₂O (A') and 0.1% formic acid in MeCN (B') were used. Gradient: 3% B' for 1 min,

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3 then 3 to 50% B' over 15 min. MS spectrum of purified Pro3-SBP was obtained by LC-MS
4 (liquid chromatography coupled to mass spectrometry) analysis using this system.
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6 **Solid-phase peptide synthesis:** Automated Fmoc-based solid phase peptide syntheses
7 (SPPS) was carried out on a Prelude synthesizer from Protein technologies. Standard side-chain
8 protecting groups were used: Arg(Pbf), Asp(OtBu), Cys(Trt), Gln(Trt) and Tyr(tBu). Synthesis
9 was performed on a 25 μmol -per-reactor scale. Protected L-amino acids (250 μmol , 10 equiv.)
10 were coupled without pre-activation using HCTU (2-(6-chloro-1*H*-benzotriazole-1-yl)-1,1,3,3-
11 tetramethylammonium hexafluorophosphate, 90 mg, 238 μmol , 9.5 equiv.) and DIEA (87 μL , 500
12 μmol , 20 equiv.) in NMP (3 mL) for 30 min. A modified protocol was used for D-amino acids
13 building blocks: protected D-amino acids (87.5 μmol , 3.5 equiv.) were coupled using HATU
14 (1-[*bis*(dimethylamino)methylene]-1*H*-1,2,3-triazolo[4,5-*b*]pyridinium 3-oxide
15 hexafluorophosphate, 32 mg, 85 μmol , 3.4 equiv.) and DIEA (30 μL , 175 μmol , 7 equiv.) in
16 NMP (3 mL) for 2 h. After each coupling, capping of potential unreacted amine groups was
17 achieved by treatment with acetic anhydride (143 μL , 1.51 mmol, 60 equiv.), DIEA (68 μL ,
18 0.39 mmol, 15.5 equiv.) and HOBt (*N*-hydroxybenzotriazole, 6 mg, 0.044 mmol, 1.8 equiv.) in
19 NMP (3 mL) for 7 min. Fmoc groups were removed by three successive treatments with 20%
20 piperidine in NMP (3 mL) for 3 min. Peptidyl resin was extensively washed with DCM
21 (dichloromethane) then dried under an air flow, and the crude peptide was deprotected and
22 cleaved from the resin through a treatment with TFA/H₂O/*i*Pr₃SiH/phenol, 88:5:2:5 for 2 h. The
23 peptide was precipitated by dilution into an ice-cold diethyl ether/petroleum ether 1:1 mixture,
24 recovered by centrifugation, and washed twice with diethyl ether and dried under reduced
25 pressure.
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41 **Synthesis and physico-chemical characterization of pro3-SBP:** The synthesis of the
42 peptide was performed as described above to give crude peptide H-
43 eeeeeaaVADXADYQaarrrrrC-NH₂, which was used without further purifications
44 (X=norvaline, L-amino acids are in uppercase and D-amino acids in lowercase). The crude
45 peptide (2.65 mg, 0.785 μmol , calculated from a molecular weight (MW) = 3377 g/mol taking
46 into account six trifluoroacetate counteranions) was dissolved in 500 μL of 1 M HEPES buffer
47 pH 7.5, deoxygenated through several successive vacuum (\sim 15 mbar) / argon cycles in order
48 to avoid cysteine oxidation. Then, a solution of sulfo-cyanine5.5 maleimide (0.65 mg, 0.877
49 μmol , 1.12 equiv.) in 100 μL deoxygenated DMF was added under argon and the resulting
50 mixture was further deoxygenated. After 13 h stirring at RT, a second portion of the maleimide
51 reagent was added (1.12 equiv. in 100 μL deoxygenated DMF), and the reaction was stirred for
52 30 min. At this stage, HPLC analysis showed total consumption of the starting peptide material.
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3 A solution of QSY21 *N*-hydroxy succinimide ester (1.28 mg, 1.570 μmol , 2.0 equiv.) in 100
4 μL DMF was then added, and the reaction was stirred for 1 h, at RT. A second portion (1.76
5 equiv. in 88 μL DMF) was then added and the reaction stirred for 18 h. A third portion was
6 added (2.24 equiv. in 112 μL DMF) and the reaction stirred for an additional 6 h. Reaction
7 mixture was analyzed by HPLC to assess the completion of the reaction. The reaction was
8 quenched by dilution into 5 mL of a $\text{H}_2\text{O}/\text{MeCN}/\text{TFA}$ 79.5:19.5:1 mixture then purified by
9 semi-preparative HPLC and lyophilized to yield Pro3-SBP as a deep blue powder (37 nmol,
10 determined by UV titration at 675 nm, $\epsilon = 315\ 100\ \text{M}^{-1}\cdot\text{cm}^{-1}$, 4.7% yield). Theoretical molecular
11 absorption coefficient of pro3-SBP at 675 nm was calculated by addition of the reported one
12 for sulfo-Cy5.5 at this wavelength (λ_{max} , $\epsilon = 235\ 000\ \text{M}^{-1}\cdot\text{cm}^{-1}$)^a to the one determined for
13 QSY21 ($\epsilon^{675} = 80\ 100\ \text{M}^{-1}\cdot\text{cm}^{-1}$) from its absorption spectrum,^b based on its reported $\epsilon = 89\ 000$
14 $\text{M}^{-1}\cdot\text{cm}^{-1}$ at $\lambda_{\text{max}} = 660\ \text{nm}$. Note that the conjugate showed limited solubility and tendency to
15 precipitate under HPLC purification conditions, which explains the moderate (non-optimized)
16 isolated yield. However, pro3-SBP is perfectly soluble and stable in a DMSO
17 (dimethylsulfoxide) stock solution and does not show tendency to aggregate or precipitate
18 under our *in vitro* or *in cellulo* tests conditions.

19 HPLC: semi-preparative purification gradient: 20-80% B over 30 min, retention time = 19.8
20 min. Analytical gradient: 20-80% B over 5 min, retention time: 3.76 min.

21 ESI-MS (m/z): $[\text{M}+\text{H}]^{3+}$ calcd. for $\text{C}_{195}\text{H}_{262}\text{N}_{47}\text{O}_{58}\text{S}_6$: 1461.6, found: 1461.4; $[\text{M}+2\text{H}]^{4+}$ calcd.
22 for $\text{C}_{195}\text{H}_{263}\text{N}_{47}\text{O}_{58}\text{S}_6$: 1096.5, found: 1096.2. UV: $\lambda_{\text{max}} = 675.0\ \text{nm}$ in $\text{H}_2\text{O}/\text{MeCN}/\text{TFA}$
23 70/30/0.1.

41 Enzymes, inhibitors, reagents:

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43 Abz-Val-Ala-Asp-Nva-Ala-Asp-Tyr-Gln-EDDnp was synthesized according to
44 Epinette et al. (2012).⁴⁰ Synthesis of the peptidyl di(chlorophenyl)-phosphonate ester, Biotinyl-
45 Val-Tyr-Asp-Nva-P(O-C₆H₄-4-Cl)₂ was detailed elsewhere.²⁶ Human PR3 (hPR3) was
46 produced as reported earlier.⁴⁹ Human neutrophil elastase (hNE) was provided by BioCentrum
47 (Krakow, Poland). Human CatG was from ICN Pharmaceuticals (Costa Mesa, CA, USA).
48 Human CatS was produced as previously described.⁵⁰ Human CatD was provided by Merck
49 KGaA (Darmstadt, Germany). Human cathepsins B, L, and H and human matrix
50 metalloproteinases (MMP-8, MMP-9, MMP-12) were purchased from Calbiochem (VWR
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59 ^a <https://www.lumiprobe.com/p/sulfo-cy55-nhs-ester>, last retrieved : 12/05/2021

60 ^b <https://www.aatbio.com/absorbance-uv-visible-spectrum-graph-viewer/>, last retrieved : 12/05/2021

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3 International, Pessac, France). Brij35, EDTA, DTT and NP40 were obtained from Sigma-
4 Aldrich (Saint Quentin Fallavier, France). Activity buffers were 0.05 M HEPES buffer, pH 7.4,
5 0.05% NP40, 150 mM NaCl for hPR3, hNE, and CatG, 0.1 M sodium citrate buffer, pH 4.0 for
6 aspartic CatD, 0.1 M sodium phosphate buffer, pH 7.4, 2 mM DTT, 1 mM EDTA and 0.01%
7 Brij35 for CatS, 0.1 M sodium acetate buffer, pH 5.5, containing 2 mM DTT, 1 mM EDTA and
8 0.01% Brij 35 for cathepsins B, H, and L, and 0.05 M Tris buffer, pH 7.5, containing 0.05%
9 Brij35, 150 mM NaCl, and 10 mM CaCl₂ for MMP-8, MMP-9, and MMP-12, respectively.

16 17 **Preparation of human neutrophils:**

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19 Blood samples were collected from healthy volunteers by the "Etablissement Français
20 du Sang Centre-Pays de Loire" (EFS Tours, France). Specimens were delivered by EFS
21 (University of Tours/EFS agreement) in accordance with ethical rules (approval number
22 #CPDL-PLER-2019 188). Neutrophils were isolated from EDTA-whole blood by negative
23 magnetic selection using the "EasySep Direct Human Neutrophil Isolation Kit" (Stemcell
24 Technologies, Cambridge, UK).

27 28 29 30 **Characterization of the NIRF probe (pro3-SBP):**

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32 **Kinetics parameters of pro3-SBP:** hPR3 (10 nM) was incubated for 5 min at 37 °C in
33 the assay buffer prior addition of pro3-SBP (1 μM final concentration). Determination of the
34 second-order rate constants (k_{cat}/K_m) were carried out under pseudo first-order conditions. The
35 enzymatic activity was followed by monitoring the hydrolysis of pro3-SBP (λ_{ex} : 684 nm; λ_{em} :
36 710 nm) (Cary Eclipse spectrofluorimeter, Agilent Technologies, Les Ulis, France). The half-
37 time of the reaction ($t_{1/2}$) was obtained graphically by plotting $[P] = f(t)$, with $k_{obs} = \ln 2/t_{1/2}$.
38 The value of k_{cat}/K_m ($M^{-1}.s^{-1}$) was deduced from the equation $k_{cat}/K_m = k_{obs}/[E]$ (GraphPad Prism
39 6, San Diego, CA, USA). Kinetic data were reported as means \pm S.D (three independent
40 experiments). The maximal velocity of the reaction (V_{max}) and the Michaelis constant (K_m)
41 were fitted from initial velocity measurements (GraphPad Prism 6), by varying pro3-SBP
42 concentrations (0.5-8 μM) for a constant hPR3 concentration (5 nM) (n=3). The same
43 experience was repeated in the presence of hHNE.

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52 **Selectivity of pro3-SBP:** Pro3-SBP selectivity was checked by incubating proteases
53 (10 nM) 5 min at 37°C in their respective activity buffer prior addition of pro3-SBP (1 μM).
54 Enzymatic activity was recorded by monitoring the fluorescence release (λ_{ex} : 684 nm; λ_{em} : 710
55 nm) (Cary Eclipse spectrofluorimeter). On the other hand, pro3-SBP (1 μM) was incubated
56 with hPR3 for 30 min at 37 °C in its assay buffer (final volume: 100 μL). The reaction was
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3 stopped by addition of ethanol (500 μ L). After centrifugation (6,000 rpm/10 min, 4 $^{\circ}$ C), the
4 precipitate was removed and the native peptide and/or its proteolytic fragments was evaporated
5 to dryness and re-dissolved in water containing 0.1% trifluoroacetic acid. Proteolysis products
6 were separated by reverse-phase chromatography (Brownlee Aquapore RP-300, C8 Phase
7 Cartridge Column, PerkinElmer), using a 41-min linear (0–90%) gradient of acetonitrile in
8 water (0.1% TFA) at a flow rate of 0.5 mL/min (P1000XR, SpectraSystem), and analyzed by
9 the ChromQuest chromatography Workstation (ThermoFinnigan, Les Ulis, France). The same
10 experiment was conducted for hNE, CatG and CatS (10 nM) in their respective assay buffer.
11 Identification of cleavage sites were performed by analysis of the hydrolysis products by
12 MALDI-TOF mass spectrometry (MALDI-TOF/TOF UltraFleXtreme, Bruker Daltonics).
13 QSY21-eeeeeeaaVADX-OH (X=norvaline) (m/z): monoisotopic $[M]^+$ calculated for
14 $C_{89}H_{109}N_{14}O_{28}S$: 1853.73, found: 1853.72. QSY21-eeeeeeaaVADXA-OH (X=norvaline) (m/z ,
15 monoisotopic): $[M]^+$ calculated for $C_{92}H_{114}N_{15}O_{29}S$: 1924.76, found: 1924.75. H-
16 ADYQaarrrrrC(sulfoCy5.5)-NH₂ (m/z , monoisotopic): $[M]^+$ calculated for $C_{106}H_{154}N_{33}O_{31}S_5$:
17 2545.01, found: 2545.00. H-DYQaarrrrrC(sulfoCy5.5)-NH₂ (m/z , monoisotopic): $[M]^+$
18 calculated for $C_{103}H_{149}N_{32}O_{30}S_5$: 2473.97, found: 2473.95.

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31 **Physicochemical properties of pro3-SBP:** The stability of pro3-SBP (1 μ M) in an
32 aqueous buffer was endorsed by incubation for 0-60 min at 37 $^{\circ}$ C in its assay buffer (final
33 volume: 100 μ L) as reported elsewhere.⁴⁶ Alternatively, after addition of ethanol (500 μ L),
34 centrifugation (6,000 rpm/10 min, 4 $^{\circ}$ C) and evaporation to dryness, integrity of pro3-SBP was
35 checked by RP-HPLC (Brownlee Aquapore RP-300) using conditions reported above
36 (ChromQuest chromatography Workstation). Moreover, the possible hazard of an
37 intermolecular fluorescence quenching associated to pro3-SBP hydrolysis and the subsequent
38 release of intramolecular sulfoCy5.5/QSY21 couple was evaluated by incubation of hPR3 (5
39 nM) with increasing amounts of the probe (0-8 μ M) at 37 $^{\circ}$ C in the assay buffer (λ_{ex} : 684 nm;
40 λ_{em} : 710 nm).

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48 **Detection threshold of pro3-SBP:** Increasing amounts of hPR3 (0-5 nM) were
49 incubated in the presence of pro3-SBP (0.5 μ M) in 0.05 M HEPES buffer, pH 7.4, 0.05% NP40,
50 150 mM NaCl (black 96-well microtiter plates, Nunc, ThermoFisher Scientific, Villebon-sur-
51 Yvette, France). After 5 min, fluorescence release was measured using IVIS Caliper Lumina
52 XR imaging system (10 s exposure, Caliper LifeSciences, Villepinte, France) (λ_{ex} : 640 nm; λ_{em} :
53 675 nm). Data were presented on color scale overlaid on a grayscale photograph (microplate
54 wells). They were quantified as total radiant efficiency (photons (p) per second (s) per cm² per
55 steradian (sr) [photons/s/cm²/sr]) within a circular region of interest (1×10^3 pixels) using the
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3 Living Image 4.3.1 software (Caliper Life Sciences).⁴⁶ Similarly, hPR3 (5 nM) was incubated
4 in the presence of increasing concentrations of pro3-SBP (0-5 μ M), before recording the
5 fluorescence release (t = 0-25 min). Assay was repeated by pre-incubating hPR3 (5 nM) in the
6 presence of a hPR3 inhibitor, Biotinyl-Val-Tyr-Asp-Nva-P(O-C₆H₄-4-Cl)₂ (10 μ M)²⁶, prior
7 addition of pro3-SBP (5 μ M) and recording of the fluorescence release 10 min latter. On the
8 other hand, purified neutrophils (5x10⁶ neutrophils) were activated by ionomycin, a calcium
9 ionophore (1 μ M, Sigma-Aldrich), for 20 min at 37°C in phosphate-buffered saline/1 mM
10 CaCl₂/1 mM MgCl₂ (final volume: 5 mL). Then, 500,000 neutrophils/well were seeded in 96-
11 well plates at 37°C (5% CO₂) in Gibco free-fetal calf serum RPMI-1640 medium (Fisher
12 Scientific, Illkrich, France) for 24 h. Cell supernatants were collected and the fluorescence
13 release was monitored (IVIS Caliper Lumina XR imaging system, t = 0-30 min), following
14 addition of pro3-SBP (5 μ M). Controls were carried out in the presence of quiescent PMNs,
15 and selectivity of NIRF signal was checked by addition of a hPR3 inhibitor, Biotinyl-Val-Tyr-
16 Asp-Nva-P(O-C₆H₄-4-Cl)₂ (1 μ M).
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29 **Activation of human neutrophils: imaging by confocal microscopy:**

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31 Following their isolation, purified neutrophils (5x10⁶ neutrophils) were immediately
32 stimulated with phosphate-buffered saline/1 mM CaCl₂/1 mM MgCl₂, containing 250 nM
33 ionomycin for 20 min at 37°C (final volume: 5 mL). Then, cells (500,000 neutrophils/well)
34 were seeded in 96-well Well Black ibiTreat microplates (Ibidi, CliniSciences, Nanterre, France)
35 at 37°C (5% CO₂) in Gibco FCS-free RPMI-1640 medium for 4 h. After removal of non-
36 adherent cells, pro3-SBP (10 μ M) was immediately added to the culture medium and imaging
37 performed by confocal fluorescence microscopy (Leica SP8 Confocal Microscope, Leica
38 Microsystemes, Nanterre, France) (0-15 min.). Data were acquired using White Light Laser
39 (Cys5.5; wavelength 675 nm) and transmitted light detector (diode 405; wavelength 405nm)
40 with constant microscope settings (laser power, detector gain, and amplification offset). Assays
41 were repeated with non-activated (quiescent) neutrophils. Alternatively, a control experiment
42 was conducted; cell media of both quiescent and activated neutrophils were withdrawn and
43 hPR3 activity was monitored using Abz-Val-Ala-Asp-Nva-Ala-Asp-Tyr-Gln- EDDnp (20 μ M)
44 as substrate in 0.05 M HEPES, 0.75 M NaCl, and 0.05% NP40, pH 7.4
45 (spectromicrofluorimeter SpectraMax Gemini, Molecular Devices, Saint Grégoire, France;
46 λ_{ex} =320 nm, λ_{em} =420 nm). Selectivity of substrate was checked by adding Biotinyl-Val-Tyr-
47 Asp-Nva-P(O-C₆H₄-4-Cl)₂ (1 μ M). Experiments were performed in triplicate.
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Cell viability assay:

CellTiter 96® AQueous One Solution Cell Proliferation Assay (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium, MTS) kit (Promega, Charbonnière-Les-Bains, France) was used to evaluate cytotoxicity of pro3-SBP, according to the manufacturer's instructions. Briefly, both ionomycin-treated and untreated neutrophils (500,000 cells/well) were cultured for 4 h in 96-well plates containing 100 μ L of medium as described above. At various time (0, 5, and 15 min), pro3-SBP (10 μ M) was incubated with culture medium at 37°C (5% CO₂). The MTS reagent was added (20 μ L/well) and cells were incubated under the same conditions for 2 h. The absorbance was measured at 490 nm (Microplate Reader VersaMAX, Molecular Devices, St Grégoire, France). All the experiments were repeated three times. MTS assay results were normalized using as reference the proliferation rate of quiescent neutrophils in the absence of pro3-SBP (control value defined as 100%). Assays were performed in triplicate.

Statistical analysis:

Data are expressed as means \pm standard deviation (SD). Statistical significance between the different values was analyzed by nonparametric Mann-Whitney U test and group comparisons were performed with nonparametric Kruskal-Wallis test. Statistical analysis was performed using GraphPad Prism v8.4.2 (San Diego, CA, USA). Differences at a *P*-value < 0.05 were considered significant (**: *P* < 0.005; ***: *P* < 0.0005).

Supporting Information:

Figure S1. Characterization of purified pro3-SBP (HPLC trace, UV-vis and ESI-MS spectra)

Figure S2. Cleavage of pro3-SBP by hPR3.

Figure S3. Cleavage of pro3-SBP by hNE.

Figure S4. Cell viability assay.

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Competing interests: The authors declare that they have no competing interests.

Abbreviations: Abz, *o*-aminobenzoic acid; AMC, 7-amino-4-methyl coumarin; ARDS, acute respiratory distress syndrome ; Brij35, polyethylene glycol lauryl ether; CatB, cathepsin B; CatD, cathepsin D; CatG, cathepsin G; CatH, cathepsin H; CatL, cathepsin L; CatS, cathepsin S; CPP, cell-penetrating peptide; COPD, chronic obstructive pulmonary disease ; DMF, dimethylformamide; DTT; dithiothreitol; EDDnp, *N*-(2,4-dinitrophenyl)ethylenediamine; FCS, fetal calf serum; FRET, fluorescence resonance energy transfer; HEPES, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid; hNE, human neutrophil elastase; Fmoc, fluorenylmethyloxycarbonyl; IL-8, interleukin 8; IL-1 β , interleukin-1 beta; MALDI-TOF, matrix assisted laser desorption ionization time-of-flight; MeCN, acetonitrile ; MMP-8, matrix metalloproteinase-8; MMP-9, matrix metalloproteinase-9, MMP-12, matrix metalloproteinase-12; MTS, 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium; NHS, *N*-hydroxysuccinimide; NIRF, near infrared fluorescence; NP40, nonyl phenoxyethoxyethanol; NSP, neutrophil serine protease; NMP, *N*-methyl-2-pyrrolidone; Pbf, pentamethyldihydrobenzofuran-5-sulfonyl; PMN, polymorphonuclear neutrophil; hNE, human neutrophil elastase; hPR3, human proteinase 3; RP-HPLC, reversed phase high performance liquid chromatography; SPPS, solid phase peptide synthesis; TFA, trifluoroacetic acid; TNF- α , tumor necrosis factor alpha ; Trt, trityl.

References

- (1) Pham, C. T. N. (2006) Neutrophil serine proteases: specific regulators of inflammation. *Nat. Rev. Immunol.* 6, 541–550.
- (2) Korkmaz, B., Horwitz, M. S., Jenne, D. E., and Gauthier, F. (2010) Neutrophil elastase, proteinase 3, and cathepsin G as therapeutic targets in human diseases. *Pharmacol. Rev.* 62, 726–759.
- (3) Perera, N. C., Schilling, O., Kittel, H., Back, W., Kremmer, E., and Jenne, D. E. (2012) NSP4, an elastase-related protease in human neutrophils with arginine specificity. *Proc Natl. Acad. Sci. U. S. A.* 109, 6229–6234.
- (4) AhYoung, A. P., Lin, S. J., Gerhardy, S., van Lookeren Campagne, M., and Kirchhofer, D. (2019) An ancient mechanism of arginine-specific substrate cleavage: What’s “up” with NSP4? *Biochimie* 166, 19–26.
- (5) Korkmaz, B., Caughey, G. H., Chapple, I., Gauthier, F., Hirschfeld, J., Jenne, D. E., Kettritz, R., Lalmanach, G., Lamort, A.-S., Lauritzen, C., et al. (2018) Therapeutic targeting of cathepsin C: from pathophysiology to treatment. *Pharmacol. Ther.* 190, 202–236.
- (6) Owen, C. A. (2008) Roles for proteinases in the pathogenesis of chronic obstructive pulmonary disease. *Int. J. Chron. Obstruct. Pulmon. Dis.* 3, 253–268.
- (7) Korkmaz, B., Moreau, T., and Gauthier, F. (2008) Neutrophil elastase, proteinase 3 and cathepsin G: physicochemical properties, activity and physiopathological functions. *Biochimie* 90, 227–242.
- (8) Brinkmann, V., Reichard, U., Goosmann, C., Fauler, B., Uhlemann, Y., Weiss, D.S., Weinrauch, Y. and Zychlinsky, A. (2004) Neutrophil extracellular traps kill bacteria. *Science* 303, 1532–1535.
- (9) Witko-Sarsat, V., Cramer, E.M., Hieblot, C., Guichard, J., Nusbaum, P., Lopez, S., Lesavre, P. and Halbwachs-Mecarelli, L. (1999) Presence of proteinase 3 in secretory vesicles: evidence of a novel, highly mobilizable intracellular pool distinct from azurophil granules. *Blood* 94, 2487–2496.
- (10) Weiss, S. A. I., Rehm, S. R. T., Perera, N. C., Biniossek, M. L., Schilling, O., and Jenne, D. E. (2021) Origin and Expansion of the Serine Protease Repertoire in the Myelomonocyte Lineage. *Int. J. Mol. Sci.* 22.
- (11) Sørensen, O. E., Follin, P., Johnsen, A. H., Calafat, J., Tjabringa, G. S., Hiemstra, P. S., and Borregaard, N. (2001) Human cathelicidin, hCAP-18, is processed to the antimicrobial peptide LL-37 by extracellular cleavage with proteinase 3. *Blood* 97, 3951–3959.
- (12) Renesto, P., Halbwachs-Mecarelli, L., Nusbaum, P., Lesavre, P., and Chignard, M. (1994) Proteinase 3. A neutrophil proteinase with activity on platelets. *J. Immunol.* 152, 4612–4617.

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2
3 (13) Sköld, S., Rosberg, B., Gullberg, U., and Olofsson, T. (1999) A secreted proform of
4 neutrophil proteinase 3 regulates the proliferation of granulopoietic progenitor cells. *Blood*
5 93, 849–856.
6
7 (14) Yang, M., Li, K., Lam, A. C., Yuen, P. M., Fok, T. F., Chesterman, C. N., and Chong, B.
8 H. (2001) Platelet-derived growth factor enhances granulopoiesis via bone marrow
9 stromal cells. *Int. J. Hematol.* 73, 327–334.
10
11 (15) Lüdemann, J., Utecht, B., and Gross, W. L. (1990) Anti-neutrophil cytoplasm antibodies
12 in Wegener’s granulomatosis recognize an elastinolytic enzyme. *J. Exp. Med.* 171, 357–
13 362.
14
15 (16) Hedstrom, L. (2002) Serine protease mechanism and specificity. *Chem. Rev.* 102, 4501–
16 4524.
17
18 (17) Fujinaga, M., Chernaia, M. M., Halenbeck, R., Koths, K., and James, M. N. (1996) The
19 crystal structure of PR3, a neutrophil serine proteinase antigen of Wegener’s
20 granulomatosis antibodies. *J. Mol. Biol.* 261, 267–278.
21
22 (18) Bode, W., Wei, A. Z., Huber, R., Meyer, E., Travis, J., and Neumann, S. (1986) X-ray
23 crystal structure of the complex of human leukocyte elastase (PMN elastase) and the third
24 domain of the turkey ovomucoid inhibitor. *The EMBO J.* 5, 2453–2458.
25
26 (19) Korkmaz, B., Attucci, S., Moreau, T., Godat, E., Juliano, L., and Gauthier, F. (2004)
27 Design and use of highly specific substrates of neutrophil elastase and proteinase 3. *Am.*
28 *J. Respir. Cell. Mol. Biol.* 30, 801–807.
29
30 (20) Korkmaz, B., Kellenberger, C., Viaud-Massuard, M.-C., and Gauthier, F. (2013) Selective
31 inhibitors of human neutrophil proteinase 3. *Curr. Pharm. Des.* 19, 966–976.
32
33 (21) Korkmaz, B., Lesner, A., Guarino, C., Wysocka, M., Kellenberger, C., Watier, H., Specks,
34 U., Gauthier, F. and Jenne, D.E. (2016) Inhibitors and Antibody Fragments as Potential
35 Anti-Inflammatory Therapeutics Targeting Neutrophil Proteinase 3 in Human Disease.
36 *Pharmacol. Rev.* 68, 603–630.
37
38 (22) Popow-Stellmaszyk, J., Wysocka, M., Lesner, A., Korkmaz, B., and Rolka, K. (2013) A
39 new proteinase 3 substrate with improved selectivity over human neutrophil elastase.
40 *Anal. Biochem.* 442, 75–82.
41
42 (23) Edgington, L. E., Verdoes, M., and Bogoyo, M. (2011) Functional imaging of proteases:
43 recent advances in the design and application of substrate-based and activity-based probes.
44 *Curr. Opin. Chem. Biol.* 15, 798–805.
45
46 (24) Sanman, L. E., and Bogoyo, M. (2014) Activity-based profiling of proteases. *Annu. Rev.*
47 *Biochem.* 83, 249–273.
48
49 (25) Vizovišek, M., Fonović, M., and Turk, B. (2018) Cysteine cathepsins in extracellular
50 matrix remodeling: Extracellular matrix degradation and beyond. *Matrix Biol.* 75-76,
51 141-159.
52
53
54
55
56
57
58
59
60

- 1
2
3 (26) Guarino, C., Gruba, N., Grzywa, R., Dyguda-Kazimierowicz, E., Hamon, Y., Łęgowska,
4 M., Skoreński, M., Dallet-Choisy, S., Marchand-Adam, S., Kellenberger, C., et al. (2018)
5 Exploiting the S4-S5 Specificity of Human Neutrophil Proteinase 3 to Improve the
6 Potency of Peptidyl Di(chlorophenyl)-phosphonate Ester Inhibitors: A Kinetic and
7 Molecular Modeling Analysis. *J. Med. Chem.* 61, 1858–1870.
8
9
10 (27) Kasperkiewicz, P., Poreba, M., Snipas, S. J., Parker, H., Winterbourn, C. C., Salvesen, G.
11 S., and Drag, M. (2014) Design of ultrasensitive probes for human neutrophil elastase
12 through hybrid combinatorial substrate library profiling. *Proc. Natl. Acad. Sci. U. S. A.*
13 111, 2518–2523.
14
15 (28) Kasperkiewicz, P., Poreba, M., Snipas, S. J., Lin, S. J., Kirchhofer, D., Salvesen, G. S.,
16 and Drag, M. (2015) Design of a Selective Substrate and Activity Based Probe for Human
17 Neutrophil Serine Protease 4. *PLoS One* 10, e0132818.
18
19 (29) Kasperkiewicz, P., Altman, Y., D'Angelo, M., Salvesen, G. S., and Drag, M. (2017)
20 Toolbox of Fluorescent Probes for Parallel Imaging Reveals Uneven Location of Serine
21 Proteases in Neutrophils. *J. Am. Chem. Soc.* 139, 10115–10125.
22
23 (30) Kasperkiewicz, P., Poreba, M., Groborz, K., and Drag, M. (2017) Emerging challenges in
24 the design of selective substrates, inhibitors and activity-based probes for
25 indistinguishable proteases. *FEBS J.* 284, 1518–1539.
26
27 (31) Guarino, C., Legowska, M., Epinette, C., Kellenberger, C., Dallet-Choisy, S., Sieńczyk,
28 M., Gabant, G., Cadene, M., Zoidakis, J., Vlahou, A., et al. (2014) New selective peptidyl
29 di(chlorophenyl) phosphonate esters for visualizing and blocking neutrophil proteinase 3
30 in human diseases. *J. Biol. Chem.* 289, 31777–31791.
31
32 (32) Anderson, B.M., Poole, D.P., Aurelio, L., Ng, G.Z., Fleischmann, M., Kasperkiewicz, P.,
33 Morissette, C., Drag, M., van Driel, I.R., Schmidt, B.L., et al. (2019) Application of a
34 chemical probe to detect neutrophil elastase activation during inflammatory bowel
35 disease. *Sci. Rep.* 9, 13295.
36
37 (33) Poreba, M., Groborz, K.M., Rut, W., Pore, M., Snipas, S.J., Vizovisek, M., Turk, B.,
38 Kuhn, P., Drag, M. and Salvesen, G.S. (2020) Multiplexed Probing of Proteolytic
39 Enzymes Using Mass Cytometry-Compatible Activity-Based Probes. *J. Am. Chem. Soc.*
40 142, 16704–16715.
41
42 (34) Burster, T., Gärtner, F., Knippschild, U., and Zhanapiya, A. (2021) Activity-Based Probes
43 to Utilize the Proteolytic Activity of Cathepsin G in Biological Samples. *Front. Chem.* 9,
44 628295.
45
46 (35) Vizovišek, M., Vidmar, R., Drag, M., Fonović, M., Salvesen, G. S., and Turk, B. (2018)
47 Protease Specificity: Towards In Vivo Imaging Applications and Biomarker Discovery.
48 *Trends Biochem. Sci.* 43, 829–844.
49
50 (36) Kasperkiewicz, P. (2021) Peptidyl Activity-Based Probes for Imaging Serine Proteases.
51 *Front. Chem.* 9, 639410.
52
53 (37) Frey, D. L., Guerra, M., Mall, M. A., and Schultz, C. (2021) Monitoring Neutrophil
54 Elastase and Cathepsin G Activity in Human Sputum Samples. *J. Vis. Exp.* (171).
55
56
57
58
59

- 1
2
3 (38) Korkmaz, B., Attucci, S., Juliano, M. A., Kalupov, T., Jourdan, M.-L., Juliano, L., and
4 Gauthier, F. (2008) Measuring elastase, proteinase 3 and cathepsin G activities at the
5 surface of human neutrophils with fluorescence resonance energy transfer substrates. *Nat.*
6 *Protoc.* 3, 991–1000.
7
8
9 (39) Jiang, T., Olson, E. S., Nguyen, Q. T., Roy, M., Jennings, P. A., and Tsien, R. Y. (2004)
10 Tumor imaging by means of proteolytic activation of cell-penetrating peptides. *Proc. Natl.*
11 *Acad. Sci. U. S. A.* 101, 17867–17872.
12
13 (40) Epinette, C., Croix, C., Jaquillard, L., Marchand-Adam, S., Kellenberger, C., Lalmanach,
14 G., Cadene, M., Viaud-Massuard, M.-C., Gauthier, F. and Korkmaz, B. (2012) A selective
15 reversible azapeptide inhibitor of human neutrophil proteinase 3 derived from a high
16 affinity FRET substrate. *Biochem. Pharmacol.* 83, 788–796.
17
18 (41) Chin, J., and Kim, H.-J. (2018) Near-infrared fluorescent probes for peptidases.
19 *Coord. Chem. Rev.* 354, 169–181.
20
21 (42) Futaki, S., Suzuki, T., Ohashi, W., Yagami, T., Tanaka, S., Ueda, K., and Sugiura, Y.
22 (2001) Arginine-rich peptides. An abundant source of membrane-permeable peptides
23 having potential as carriers for intracellular protein delivery. *J. Biol. Chem.* 276, 5836–
24 5840.
25
26 (43) Bechara, C., and Sagan, S. (2013) Cell-penetrating peptides: 20 years later, where do we
27 stand? *FEBS Lett.* 587, 1693–1702.
28
29 (44) Mitchell, D. J., Kim, D. T., Steinman, L., Fathman, C. G., and Rothbard, J. B. (2000)
30 Polyarginine enters cells more efficiently than other polycationic homopolymers. *J. Pept.*
31 *Res.* 56, 318–325.
32
33 (45) Walrant, A., Cardon, S., Burlina, F., and Sagan, S. (2017) Membrane Crossing and
34 Membranotropic Activity of Cell-Penetrating Peptides: Dangerous Liaisons? *Acc. Chem.*
35 *Res.* 50, 2968–2975.
36
37 (46) Wartenberg, M., Saidi, A., Galibert, M., Joulin-Giet, A., Burlaud-Gaillard, J., Lecaille, F.,
38 Scott, C.J., Aucagne, V., Delmas, A.F. and Lalmanach, G. (2019) Imaging of extracellular
39 cathepsin S activity by a selective near infrared fluorescence substrate-based probe.
40 *Biochimie* 166, 84–93.
41
42 (47) Fukuda, T., Ackerman, S. J., Reed, C. E., Peters, M. S., Dunnette, S. L., and Gleich, G. J.
43 (1985) Calcium ionophore A23187 calcium-dependent cytolytic degranulation in human
44 eosinophils. *J. Immunol.* 135, 1349–1356.
45
46 (48) Hoppenbrouwers, T., Autar, A.S.A., Sultan, A.R., Abraham, T.E., van Cappellen,
47 W.A., Houtsmuller, A.B., van Wamel, W.J.B., van Beusekom, H.M.M., van Neck,
48 J.W. and de Maat, M.P.M. (2017) In vitro induction of NETosis: Comprehensive live
49 imaging comparison and systematic review. *PLoS ONE* 12, e0176472.
50
51 (49) Jégot, G., Derache, C., Castella, S., Lahouassa, H., Pitois, E., Jourdan, M.L., Remold-
52 O'Donnell, E., Kellenberger, C., Gauthier, F. and Korkmaz, B. (2011) A substrate-based
53 approach to convert SerpinB1 into a specific inhibitor of proteinase 3, the Wegener's
54 granulomatosis autoantigen. *FASEB J.* 25, 3019–3031.
55
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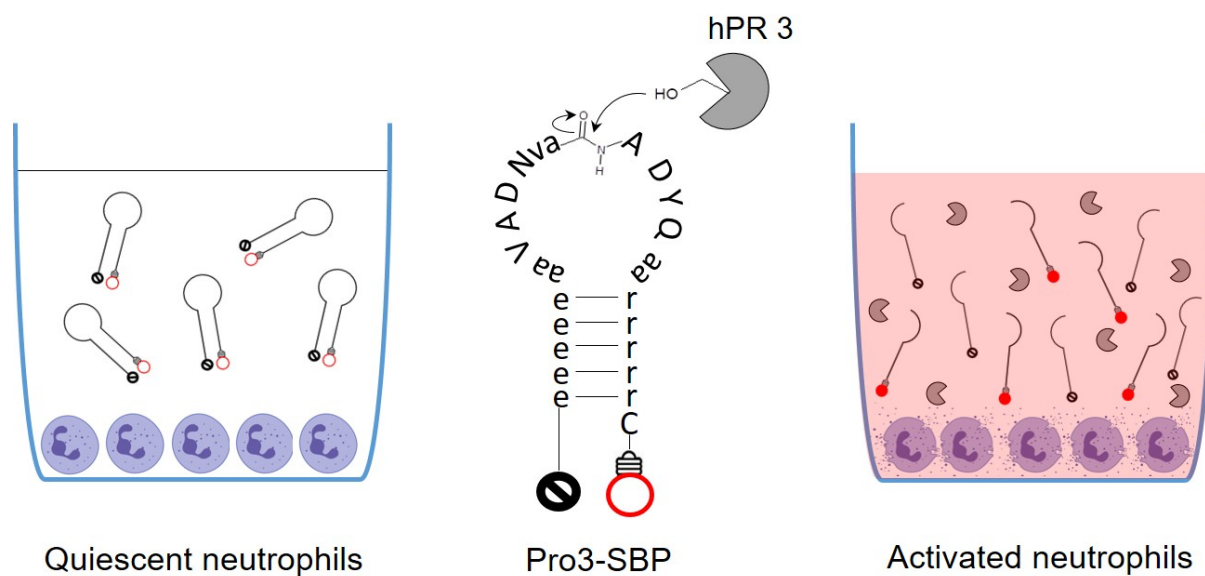
- (50) Sage, J., Leblanc-Noblesse, E., Nizard, C., Sasaki, T., Schnebert, S., Perrier, E., Kurfurst, R., Brömme, D., Lalmanach, G. and Lecaille, F. (2012) Cleavage of nidogen-1 by cathepsin S impairs its binding to basement membrane partners. *PloS One* 7, e43494.

Footnotes:

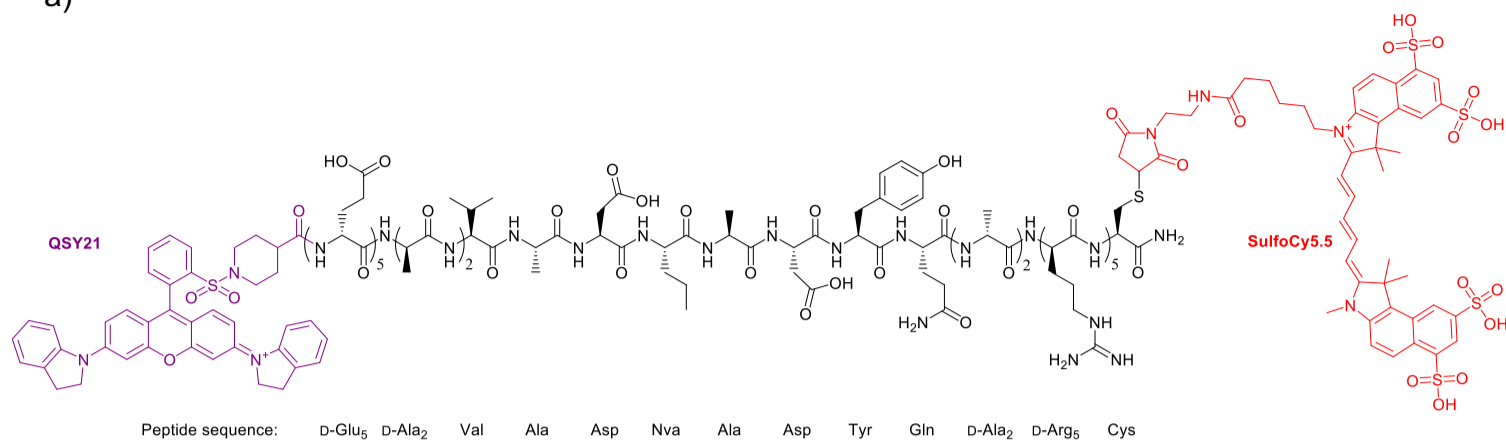
^a <https://www.lumiprobe.com/p/sulfo-cy55-nhs-ester>, last retrieved: 12/05/2021

^b <https://www.aatbio.com/absorbance-uv-visible-spectrum-graph-viewer/>, last retrieved: 12/05/2021

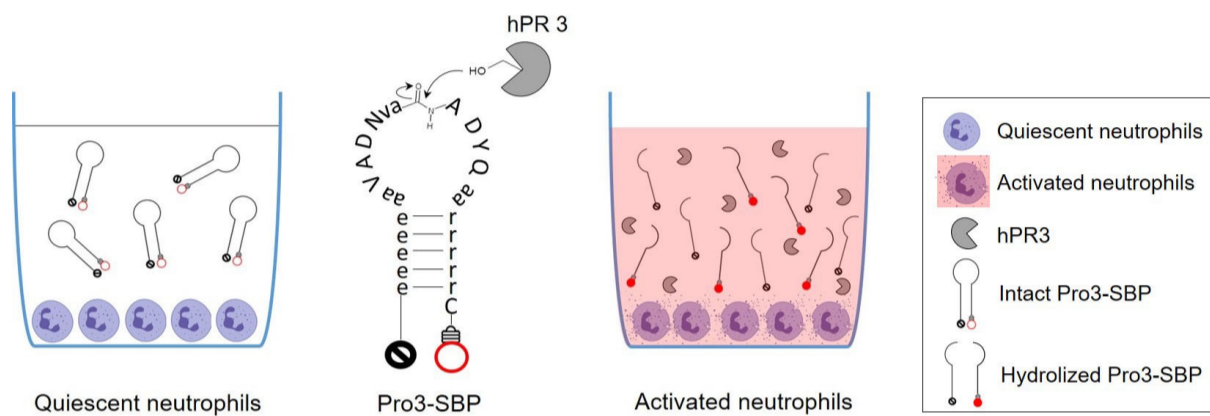
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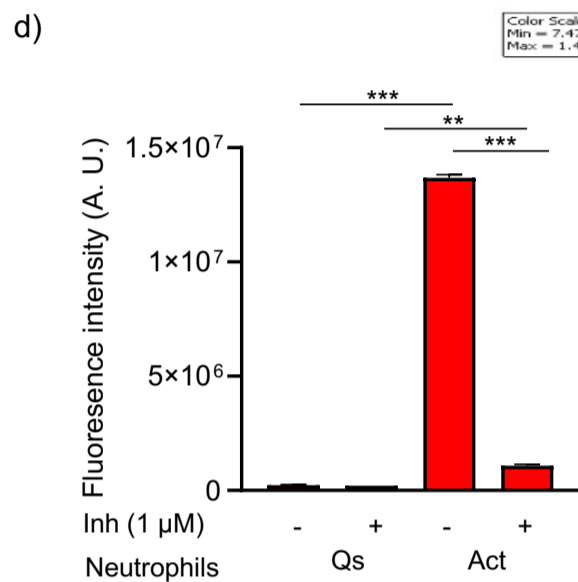
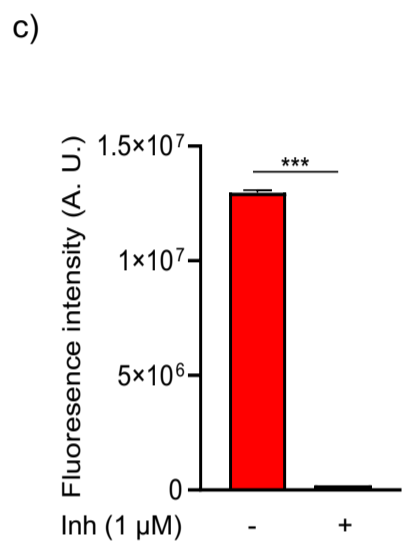
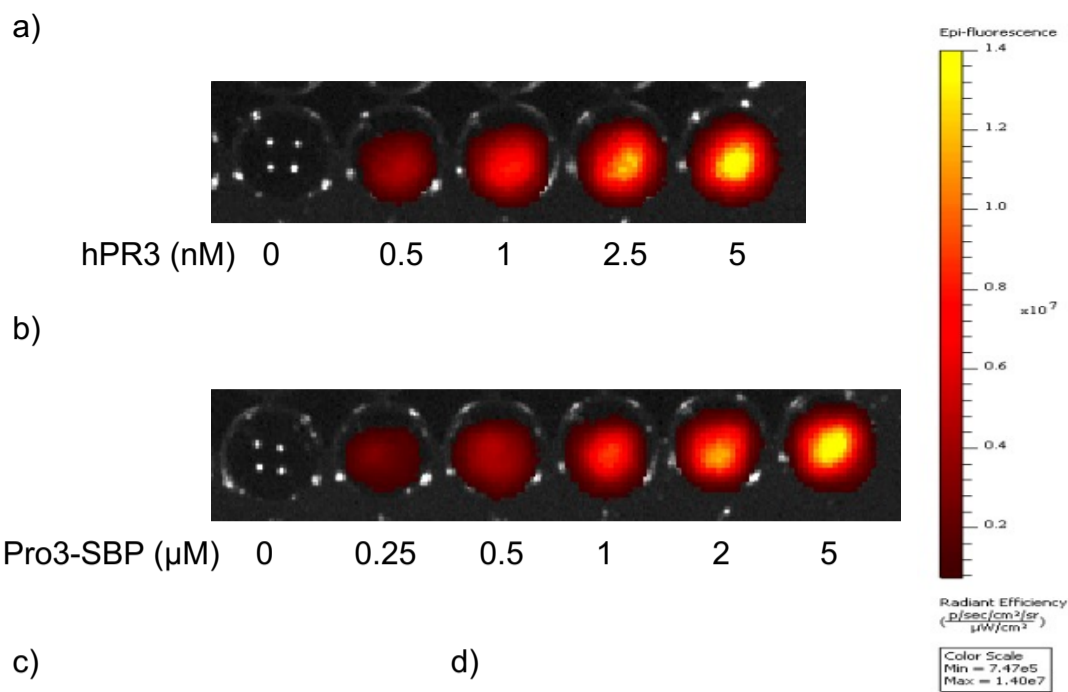


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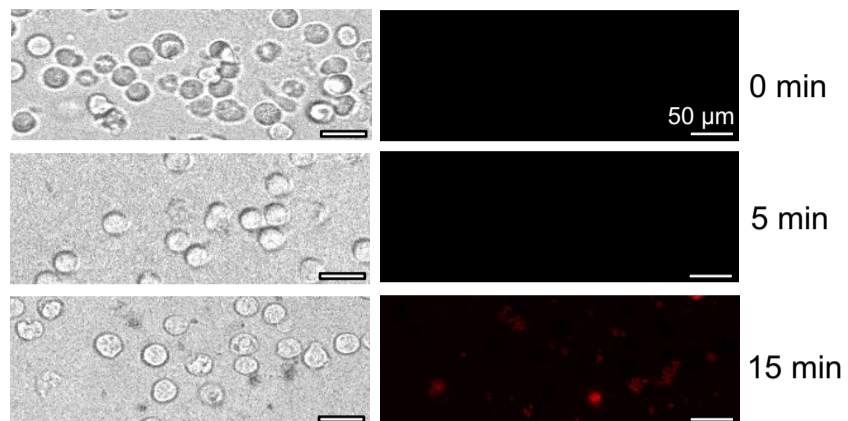
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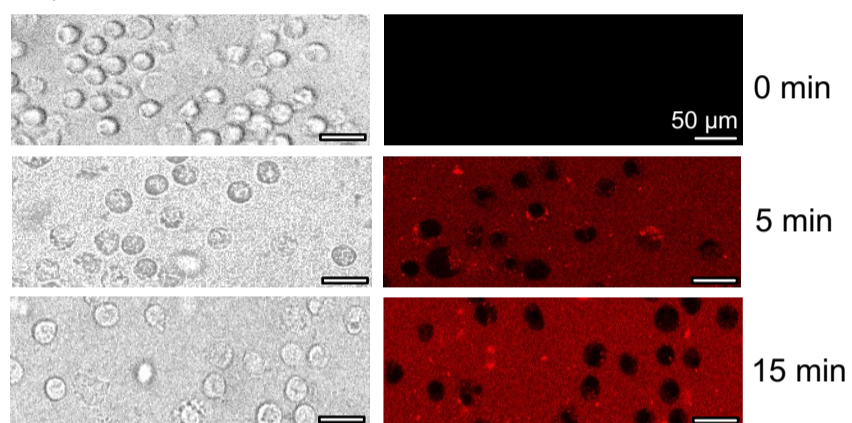


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