

Paracrine factors released by γ-irradiated peripheral blood mononuclear cells inhibit neutrophil extracellular trap formation

Doctoral thesis at the Medical University of Vienna for obtaining the academic degree

Doctor of Philosophy

Submitted by

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> > Vienna, 10/2022

Declaration

The experiments conducted to accomplish this thesis have been carried out at the Department of Thoracic Surgery and the Department of Dermatology, at the Medical University of Vienna under the supervision of Hendrik Jan Ankersmit and Michael Mildner. The project was financed by the Aposcience AG and peer reviewed third party funding as described in the manuscript section "Funding". In vitro and ex vivo experiments were performed in cooperation with Michael Mildner, Anna Ondracek, Thomas Hofbauer and Karin Pfisterer. Peripheral blood mononuclear cell secretome was produced by the Austrian Red Cross Blood Transfusion Service for Upper Austria, Linz according to GMP-requirements. Interpretation of results, writing and experimental design leading to the publication underlying this thesis were accomplished under the supervision of Michael Mildner and Hendrik Jan Ankersmit.

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Abstract English

Neutrophils, representing the largest population of circulating granulocytes, are equipped with a great variety of highly efficient effector functions. Historically, neutrophils have been considered as critical players of the immune system by phagocytosing invading pathogens or eliminating those by degranulation. However, in the past decades, a rather new effector function has received increasing interest. The discovery of the neutrophils capability to form and extrude net-like structures, so called neutrophil extracellular traps (NETs), via a process called NETosis, has shaped various research fields. NETs have been continuously linked to a plethora of pathologies. Research increasingly demonstrates the tremendous implication of NETs in disease progression and patient prognosis in autoimmune disorders, cardiovascular diseases, wound healing and tissue regeneration, cancer as well as acute respiratory distress syndrome and Covid-19. NETosis is a highly versatile process, following various signalling cascades depending on the inducing stimulus, and is yet not fully understood. However, pharmacological inhibition of NET formation, to alleviate disease burden and complications is of critically high interest. Previous studies have attributed potent immunomodulatory, regenerative and tissue-protective effects to the secretome of y-irradiation-stressed peripheral blood mononuclear cells (PBMCsec). Nevertheless, the effect of PBMCsec on neutrophils has never been investigated. In this dissertation, it is demonstrated that PBMCsec prevents NET formation in in vitro experiments in calcium ionophore and phorbol 12 myristate 13 acetate (PMA) activated neutrophils. Furthermore, it was shown that the purified individual substance classes present in PBMCsec alone do not exert significantly reduced rates of NET formation. This indicates that an interplay of several substance classes or the whole secretome is required for this inhibitory effect. In depth analysis revealed that PBMCsec most likely inhibits NET formation via a dual mechanism. It was demonstrated that reactive oxygen species production is prevented upon PBMCsec-treatment, accompanied by the upregulation of anti-oxidative factors. Furthermore, one of the key enzymes required for NETosis, protein arginine deiminase 4 (PAD4), appears to be modulated by PBMCsec as a reduced enzymatic activity was observed. Together, these findings suggest a potent therapeutic potential of PBMCsec for a diverse set of NETs-associated diseases. These data lay the foundation for future diseasetargeted studies investigating the *in vivo* effect of PBMCsec.

Abstract German

Neutrophile, die die größte Population zirkulierender Granulozyten darstellen, sind mit einer Vielzahl hocheffizienter Effektorfunktionen ausgestattet. In der Vergangenheit wurden Neutrophile als entscheidende Akteure des Immunsystems betrachtet, indem sie eindringende Pathogene phagozytieren oder diese durch Degranulation eliminierten. In den letzten Jahrzehnten hat jedoch eine neue Effektorfunktion zunehmend an Interesse gewonnen. Die Entdeckung der Fähigkeit, dass Neutrophile netzartige Strukturen, sogenannte extrazelluläre Traps (NETs), über einen als NETose bezeichneten Prozess zu bilden und auszuschütten, hat verschiedenste Forschungsgebiete zunehmend geprägt. NETs wurden fortlaufend mit einer großen Anzahl von Pathologien in Verbindung gebracht. Neue Forschungserkentnisse zeigen zunehmend die enorme Bedeutung von NETs für den Krankheitsverlauf und die Patientenprognose bei Autoimmunerkrankungen, Herz-Kreislauf-Erkrankungen, Wundheilung und Geweberegeneration, Krebs sowie akutem Atemnotsyndrom und Covid-19. NETose ist ein sehr vielseitiger Prozess, der je nach auslösendem Stimulus verschiedenen Signalkaskaden folgt und noch nicht vollständig verstanden ist. Die pharmakologische Inhibierung der NET-Bildung, zur Linderung von Krankheitslast und Komplikationen, ist jedoch von äußerst hohem Interesse. Frühere Studien haben dem Sekretom von durch v-Strahlung gestressten mononukleären Zellen des peripheren Blutes (PBMCsec) starke immunmodulatorische, regenerative und gewebeschützende Wirkungen zugeschrieben. Dennoch wurde die Wirkung von PBMCsec auf Neutrophile bis dato noch nicht untersucht. In dieser Dissertation wird gezeigt, dass PBMCsec die NET-Bildung in in vitro Experimenten in Kalziumionophor- und Phorbol-12-Myristat-13-Acetat (PMA)-aktivierten Neutrophilen verhindert. Weiterhin konnte gezeigt werden, dass die in PBMCsec vorhandenen gereinigten Einzelsubstanzklassen allein keine signifikante Reduktion der NET-Bildung bewirken. Dies weist darauf hin, dass für diese Hemmwirkung ein Zusammenspiel mehrerer Substanzklassen bzw. des gesamten Sekretoms erforderlich ist. Eine detailiertere Analyse ergab, dass PBMCsec höchstwahrscheinlich die NET-Bildung über einen dualen Mechanismus hemmt. Es wurde gezeigt, dass die Produktion reaktiver Sauerstoffspezies bei PBMCsec-Behandlung verhindert wird, begleitet von der Hochregulierung antioxidativer Faktoren. Darüber hinaus scheint eines der für NETose erforderlichen Schlüsselenzyme, Protein-Arginin-Deiminase 4 (PAD4), durch PBMCsec moduliert zu werden, da eine verringerte enzymatische Aktivität beobachtet wurde. Zusammen deuten diese Ergebnisse auf ein starkes therapeutisches Potenzial von PBMCsec für eine Vielzahl von NET-assoziierten Erkrankungen hin. Diese Daten bilden die Grundlage für zukünftige krankheitsbezogene Studien zur Untersuchung der in vivo Wirkung von PBMCsec.

Publication arising from the thesis

Paracrine factors released by γ -irradiated peripheral blood mononuclear cells inhibit neutrophil extracellular trap formation

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Antioxidants (Basel), 2022 Aug 11; 11(8):1559. doi:10.3390/antiox11081559. PMID: 36009277

Abbreviations

AAA	abdominal aortic aneurism	MMP
AAV	ANCA associated vasculitis	MPO
AMI	acute myocardial infarction	MS
ANCA	anti-neutrophil cytoplasmic antibody	mTORC
APCs	antigen presenting cells	mtROS
ARDS	acute respiratory distress syndrome	MVO
BM	bone marrow	NE
C/EBP-c	CCAAT/enhancer-binding protein alpha	NET
CR	complement receptor 1	NETosis
CXCL12	chemokine stromal-derived factor 1	ΝFκB
CXCR	CXC chemokine receptor	NGAL
DAG	diacylglycerol	NOD
DAMPs	danger associated molecular patterns	PAD4
DC	dendritic cell	PBMC
DVT	deep vein thrombosis	PBMCse
ECM	extracellular matrix	pDC
ER	endoplasmic reticulum	PG
ESC	embryonic stem cell	PKC
EVs	extracellular vesicles	PMA
G-CSF	granulocyte-colony stimulating factor	PR3
Gfi1	growth factor independent-1	RA
GM-CSF	granulocyte-macrophage-colony stimulating factor	ROS
GMP	granulocyte-monocyte progenitor	RvD
GPCR	G protein-coupled receptor	RvT
HIF-1α	hypoxia inducible factor	SERCA
HMGB1	high mobility group protein B1	SLE
HO-1	heme oxygenase 1	SOCE
HSC	hematopoietic stem cell	T1D / T2
HSP	heat shock protein	TF
ICAM	intracellular adhesion molecule	TLR
IL	interleukin	TNF
iNOS	inducible nitric oxide synthase	VAMP
IP3	inositol-triphosphate	VASP
iPSC	induced pluripotent stem cell	VCAM
JAK	janus kinase	VEGF
MAC	macrophage	VLA4
МАРК	mitogen activated protein kinase	VWF

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MMP	matrix metalloproteinase	
MPO	myeloperoxidase	
MS	multiple sclerosis	
mTORC1	mammalian target of rapamycin	
mtROS	mitochondiral reactive oxygen species	
MVO	microvascular obstruction	
NE	neutrophil elastase	
NET	neutrophil extracellular traps	
NETosis	neutrophil extracellular trap formation	
NFκB	nuclear factor kappa B	
NGAL	Lipocalin 2	
NOD	nucleotide oligomerization domain	
PAD4	protein arginine deiminase 4	
PBMC	peripheral blood mononuclear cells	
PBMCsec	secretome of γ-irradiated PBMCs	
pDC	plasmacytoid dendritic cell	
PG	prostaglandin	
PKC	protein kinase C	
PMA	phorbol 12 myristate 13 acetate	
PR3	proteinase 3	
RA	rheumatoid arthritis	
ROS	reactive oxygen species	
RvD class D resolvin		
RvT	class T resolvin	
SERCA	sarco-endoplasmic reticulum calcium ATPase	
SLE	systemic lupus erythematosus	
SOCE	store operated calcium entry	
T1D / T2D	type 1 / type 2 diabetes	
TF	tissue factor	
TLR	toll like receptor	
TNF	tumour necrosis factor	
VAMP	vesicle-associated membrane protein	
VASP	vasodilator-stimulated phosphoprotein	
VCAM		
VEGF	vascular endothelial growth factor	
VLA4	integrin α4β1	
VWF	von Willebrand factor	

Acknowledgement

Conducting this thesis and all associated projects was only possible due to the support of Hendrik Jan Ankersmit who continuously contributed to my journey with new ideas and input.

I am deeply grateful to Michael Mildner for his invaluable advice on scientific as well as personal questions and for his immense knowledge and experience which he offered unceasingly throughout my time as PhD-student. He was the one of the main driving forces at times were I struggled.

My special thanks go to my colleagues in the Research Laboratory of the Department of Surgery and the Department for Dermatology. Most specifically, I would like to extend my sincere thanks to my PhD colleagues, Dragan, Daniel and Martin for a cherished time spent together in the lab as well as in social settings and for their unbroken support.

My deep gratitude goes to my family for paving my way and their continuous encouragement.

I would also like to express my deepest appreciation for my partner Thomas and thank for his tremendous understanding, unconditional support and aid on every step of the way.

INTRODUCTION

1 Neutrophil granulocytes

Human blood not only provides living cells and tissues with oxygen and nutrients, contains platelets contributing to clotting in case of damaged vessels, but also delivers immune cells, including granulocytes, monocytes, T and B cells, to sites of infection.¹ Neutrophil granulocytes, also referred as polymorphonuclear leukocytes, contribute critically to the innate immune system as first line defenders, act as modulators of adaptive immune responses and additionally aid maintaining homeostasis.²⁻⁴ More than 10¹¹ neutrophils are produced per day within the bone marrow (BM), making them the most common leukocyte found in the circulation.⁵ The term polymorphonuclear leukocyte is derived from the uniquely characteristic segmented nucleus which precisely distinguishes neutrophils morphologically from other granulocytes as basophils or eosinophils.⁶ Besides different nuclear morphology, the varying granulocyte populations exert diverse cell type-specific functions. Basophils are mainly considered to take part in chronic allergy and allergy-induced inflammation, function as antigen presenting cells (APCs), and regulate immune cell memory and Th2 cell function.7-13 Eosinophils, the second least represented granulocyte population in the circulation, are critically involved in the control of parasitic infections but emerging evidence additionally suggests crucial contribution to bacterial and viral defence.¹⁴ Neutrophils were initially considered as solely phagocytic cells which aid in the resolution of inflammation and clearance of infections by engulfing and destroying pathogens, foreign material as well as dead cells and damaged tissue.⁴ However, increasing evidence accumulates indicating not only a proresolving involvement of neutrophils in various pathologies but also contribution to tissue damage in case of dysregulated effector functions.^{4,15-21}

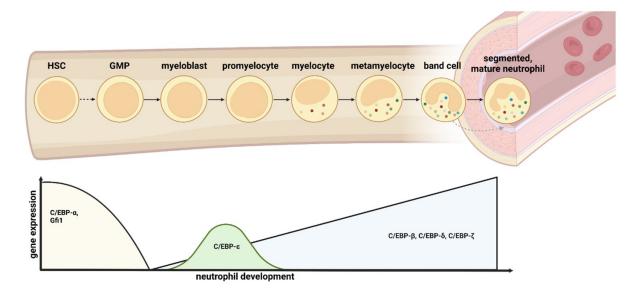
1.1 Neutrophil lineage committed granulopoiesis

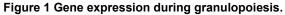
Neutrophils are produced at extensively high rates in the BM on a daily basis from hematopoietic stem cells. Upon terminally differentiation, they leave the BM and progress to patrol the circulation for signs of disturbed homeostasis and infection.²² In general, granulopoiesis describes the development of all cells belonging to the granulocyte-lineage with

the commonality of a high abundance of secretory granules. Specific neutrophil-lineage commitment is dependent on various factors including key transcription factors, inflammatory state, and environmental influence.²² Granulocyte development is initiated with the differentiation of hematopoietic stem cells into several intermediate multipotent progenitors which differentiate into granulocyte-monocyte progenitors (GMP)(Figure 1).^{5,22} Neutrophil lineage commitment is marked by further differentiation to myeloblasts followed by differentiation to promyelocytes, myelocytes, metamyelocytes and band cells. The final differentiation of band cells results in mature neutrophils.²³ To initiate commitment to myelopoiesis, the transcription factor PU.1 is of crucial importance. PU.1 expression increases throughout maturation from the promyelocyte stage.^{24,25} While GMP development is dependent on high expression levels of PU.1, continuous high expression favours monocyte-macrophage development. The transcription factor CCAAT/enhancer-binding protein alpha (C/EBP- α) antagonizes PU.1 function, thereby driving granulocyte differentiation.²⁶ Additionally, granulocyte-colony stimulating factor (G-CSF), a target of C/EBP-a, further promotes neutrophil-lineage commitment in GMPs.²⁷ However, neutrophil complexity and heterogeneity is reflected by the fact, that other cytokines including interleukin-6 (IL6) and granulocytemacrophage-colony stimulating factor (GM-CSF) can partially compensate in case of absence of G-CSF to ensure that at least some neutrophils are generated.^{28,29} Another essential transcription factor is growth factor independent-1 (Gfi1), which is upregulated in stem cells at the point of granulocyte-lineage commitment (Figure 1). High expression of Gfi1 represses monocyte-lineage favouring transcription factors and restricts stem cell proliferation promoting progressive differentiation.³⁰⁻³² C/EBP-α further contributes to granulocytic lineage decision at early stages of neutrophil development by inhibiting the cell-cycle regulator E2F1.^{33,34} Several transcription factors are required to ensure the further differentiation beyond promyelocytes. At the myelocyte stage, C/EBP- ε expression increases which is of vital importance for the transcription of granule proteins.^{25,35,36} While C/EBP- α expression gradually decreases after the myeloblast stage, C/EBP-ε peak expression levels overlap at the myelocytemetamyelocyte stage. Other C/EBP family members including C/EBP-β, C/EPB-γ, C/EBP-δ and C/EBP-ζ show continuously increasing expression levels from the metamyelocyte stage onwards.²⁵ During final maturation steps, neutrophil nuclei undergo several changes from a round shape to banded nuclei and eventually to a lobulated morphology. Progressive neutrophil differentiation is accompanied with changes in the expression of several factors involved in the retention or release of neutrophils from the BM. CXC chemokine receptor 4 (CXCR4) and integrin $\alpha 4\beta 1$ (VLA4) are downregulated while CXCR2 and Toll-like receptor 4 (TLR4) are upregulated.⁵ BM stromal cells express chemokine stromal-derived factor-1/SDF-1 (CXCL12) and vascular adhesion molecule 1 (VCAM1), ligands for CXCR4 and VLA4 respectively.⁵ It was shown that CXCR4-CXCL12 interaction represents a key mechanism

retaining neutrophils within the BM.³⁷ Only 1-2% of neutrophils enter the circulation under homeostatic conditions, highlighting the tight regulation of neutrophil liberation from the BM.³⁸ G-CSF stimulates neutrophil release by interfering with CXCR4-CXCL12 interactions and upregulation of additional CXCR2 ligands by the endothelium outside of the BM.³⁸⁻⁴⁰

Neutrophils were initially believed to be short-lived cells with a restricted life-span of only 1.5 and 8 hours in mice and humans, respectively.^{4,22} More recent studies revealed that under homeostatic conditions neutrophils appear to survive longer with an average circulatory lifespan of up to 12.5 hours in mice and 5.4 days in humans, however in this study all neutrophils, including those in the BM were assessed.⁴¹ Several studies reported a several fold extended lifespan upon neutrophil activation due to infection or inflammation resulting in a persisting tissue infiltration. ^{38,42-44} Aged circulatory neutrophils eventually return to the BM or alternatively die in peripheral tissues, both fates resulting in phagocytosis by dendritic cells (DCs) or macrophages (MAC).^{45,46} Phagocytosis of dying neutrophils was shown to function as feed-forward loop dampening neutrophil development in the bone marrow.⁴⁶ Both DCs and MAC produce IL23 upon neutrophil phagocytosis, thereby initiating IL17 production of T leukocytes, which in turn drives G-CSF dependent neutrophil precursor differentiation.⁴⁶⁻⁵⁰





Hematopoietic stem cells (HSC) differentiate upon G-CSF stimulus via several progenitor steps under the influence of C/EBP- α and Gfi1 into granulocyte-monocyte progenitors (GMP) within the bone marrow. C/EBP- ϵ expression peaks at myeloblast and promyeloblast stages. With progressing differentiation towards mature neutrophils C/EBP- β , C/EBP- δ and C/EBP- ζ expression increases and peaks when neutrophils reach a fully mature state and are released into the blood stream. This figure was created with BioRender.com.

1.2 Neutrophil effector functions

The decision of a neutrophil towards one of its effector functions, phagocytosis, degranulation or neutrophil extracellular trap (NET) formation, is a highly complex and yet poorly understood phenomenon.⁵¹ Within the circulation, neutrophils function as counterpart to tissue-resident macrophages by phagocytosing cell remnants and apoptotic cells.^{52,53} Certain cytokines, including the transcription factor GM-CSF, favour phagocytosis.⁵⁴⁻⁵⁶ Additionally, caspasemediated programmed cell death leads to the exposure of modified phosphatidylserine residues which facilitates the recognition of these cells for phagocytosis.^{57,58} Recognition of phosphatidylserines, accompanied by firm adhesion by tethering receptors, including PSGL1 for activated platelets and Tim4 for apoptotic cells, triggers phagocytosis.^{59,60} In contrast, the effect of neutrophil-platelet aggregates appears to be more complex and context dependent.⁵¹ The formation of these aggregates initially depends on the interaction of platelet P-selectin and neutrophil PSGL1 receptor.⁶¹⁻⁶⁵ Upon binding, both neutrophils and platelets upregulate factors further promoting strong and stable adhesion including integrins on neutrophils and fibrinogen on platelets.^{66,67} In dependence of overall environmental conditions, including metabolic, activation state and the interaction with the extracellular matrix, there are three possible outcomes of this interaction: 1) neutrophils dissociate from platelets thereby resolving the aggregate and may migrate towards sites of inflammation to exert effector functions ⁶⁸⁻⁷², 2) neutrophils phagocytose platelets in response to recognition of features of activated platelets as for example anionic phospholipids ^{66,72-74}, 3) the induction of NET formation.⁷⁵ In general, it appears as NET formation in vivo is triggered when inflammatory stimuli, regardless whether endogenous or microbial, exceeds a certain threshold of platelet activation.^{61,76,77} Furthermore, certain stimuli, including monosodium urate crystals cannot be phagocytosed and alternatively trigger NET formation.⁷⁸ Taken together, although not yet fully understood, pH of the microenvironment, metabolic and activation state of neutrophils, overall inflammatory state of immune cells and within tissues, distinct signalling cues as well as the size of the triggering stimuli contribute to the delicate fate decision of neutrophil effector functions.⁵¹ In contrast, degranulation is a tightly regulated receptor-coupled process requiring triggers distinctive of phagocytosis and NET formation.79

1.2.1 Degranulation

One key mechanism of neutrophil granulocytes is the rapid release of preformed granules, which are packed with toxic cargo including antimicrobial peptides and several proteases.^{80,81} There are at least three different types of granules (primary/azurophilic, secondary/specific,

tertiary/gelatinase granules) and some further subtypes as well as secretory vesicles have been suggested.⁸² It is believed that neutrophil granules contribute to the influence of both adaptive and innate immune responses.^{80,83} Neutrophil granules are obtained sequentially during differentiation and are named according to their timely appearance and major contents (Table 1). Primary granules are already formed at the promyelocyte stage and contain a large set of antimicrobial proteins and factors with antimicrobial activity, including serine proteases, myeloperoxidase (MPO), defensins and lysozyme.^{81,84} Granule proteases of primary granules were found to be activated prior to their incorporation into granules resulting in extensively potent and toxic contents that are readily available.⁸⁵ A high heterogeneity can be observed amongst primary granules as some are directed to cell surface trafficking, marked by expressing SIp1/JFC1 and Rab27, while others are prone to fuse with the phagosome lacking the before mentioned proteins.⁸⁶ Some granules were found to lack α-defensins while exhibiting all other characteristics of primary granules suggesting even greater granule heterogeneity.⁸⁷ Both, secondary and tertiary granules are formed from the myelocyte stage throughout band stage and share similar cargo and thus also functions.^{81,87} Approximately twothird of the peroxidase-negative granules contain lipocalin, MMP9 and lactoferrin and are suggested to be a hybrid form of granule subtype.⁸⁷ Furthermore, additional categorization of tertiary granules containing ficolin 1 as a ficolin 1-rich subtype was suggested. However, little is known about this potential subtype and requires more detailed research.^{88,89} During band and segmented stages of differentiation, secretory vesicles are formed by endocytosis.⁹⁰ In contrast to primary granules, which can also fuse with the phagosome, secondary as well as tertiary granules and secretory vesicles are restricted to releasing their contents by fusion with the plasma membrane and are additionally believed to aid in neutrophil adherence to previously activated endothelium.^{87,90} Mobilization and release of granules and their content is a tightly regulated process allowing selective degranulation upon specific receptor-coupling leading to distinct signalling events specific for each granule type.⁷⁹ Neutrophil degranulation is triggered by the ligation of Fcy receptors, Mac-1 or G protein-coupled receptors (GPCRs) by a diverse set of stimuli including chemokines, complement fragments, cytokines, bacterial products and endothelial adhesion molecules.⁹¹ Priming stimuli such as tumour necrosis factor (TNF), platelet-activating factor or GM-CSF induce markedly enhanced degranulation responses.⁹² Albeit different downstream signalling events are induced upon receptor ligation they all coincide to the induction of calcium flux and the activation of Rac2.93 Selective granule extrusion is further mediated by their individual requirements of certain calcium concentrations.^{87,93} Rac2, a member of the Rho GTPase subfamily, is required for actin skeleton remodelling thereby allowing granule mobilization.⁹⁴⁻⁹⁷ Rac2 deficiency in mice results in impaired primary granule release while secondary and tertiary granule degranulation was not affected, indicating the presence of additional factors influencing actin reorganization at least in mice.⁹⁶⁻⁹⁹ GPCR or Fcy receptor ligation results in the translocation of the Src-family kinase members Hck and Fgr to primary and secondary granules respectively.¹⁰⁰⁻¹⁰² Even though yet not well understood, the Src-family kinases appear to activate p38 mitogen activated protein kinase (MAPK) leading to actin rearrangement and facilitating the degranulation of all three granule types.¹⁰³ p38 MAPK is suggested to function in a similar manner to Rac2 as it is capable of inducing actin remodelling via heat shock protein 27 (HSP27).¹⁰⁴ An additional mode of action for selective degranulation is indicated by the ligation of toll like receptor 9 (TLR9) which activates NF-kB-mediated transcription of inducible nitric oxide synthase (iNOS).^{105,106} It was observed in macrophages that iNOS is capable of activating Src-kinases and could therefore be the signalling pathway by which TLR9 ligation results in primary granule degranulation.^{106,107} In addition to actin skeleton remodelling, an increased calcium concentration is an indispensable mechanism for functional degranulation. GPCR mediated calcium release is conducted via the activation of phospholipase C and subsequent inositol-triphosphate (IP₃) and diacylglycerol (DAG) production. IP₃ contributes to calcium liberation stored in the endoplasmic reticulum (ER) while DAG mediates protein kinase C (PKC) dependent store-operated channel activation.^{108,109} Additionally, other stimuli such as hypoxia or FcyRIIA-mediated phagocytosis further enhance degranulation via AKT signalling or Src-family kinase mediated IP₃ and DAG production.^{108,110} Activation of neutrophils via complement receptor 1 (CR1) and CR3 leads to the phosopholipase D dependent liberation of phosphatidic acid that can be converted into DAG, thus facilitating degranulation.⁸⁷ The final step of degranulation requires the fusion of the granule and target membrane. This mechanism is dependent on the binding of vesicle-associated membrane protein (VAMP also referred as SNARE) to cognate SNAREs (t-SNARE) found at the phagosome or plasma membrane.¹¹¹ The main SNAREs involved in neutrophil degranulation are syntaxin 6 and SNAP-23. Selective degranulation is further dependent on VAMP-SNARE interaction as VAMP-1 and VAMP-7 mediate the release of primary granules while VAMP-2 directs the release of tertiary granules.¹¹²⁻¹¹⁴ The fusion of granules with phagosomes is exclusively observed for primary granules and seems to be mediated by Rab GTPases, specifically Rab5a.¹⁰¹ Contrary, another Rab GTPase, Rab27a appears to be inflicted in degranulation of all three granule subtypes.⁸⁷ Released neutrophil granule contents such as MPO or elastase not only contribute to antimicrobial host defence but also function in an autocrine and paracrine manner modulating the mode of neutrophil cell death, life-span and function.¹¹⁵⁻¹¹⁷ MPO is capable of regulating Mac-1 expression and activation of MAPK/ERK as well as PI3K/Akt pathways which contribute to the prolongation of neutrophil life-span by preserving anti-apoptotic proteins. This feedforward loop may amplify neutrophil persistence and thus the inflammatory response.^{115,118,119} Contrasting, a pro-resolving regulatory role was attributed to lactoferrin by its ability to stimulate

IL10 secretion from macrophages and selectively inhibiting activation and migration of neutrophils.^{120,121}

Nomenclature	Primary granules	Secondary granules	Tertiary granules	Secretory vesicles
Alternative nomenclature	Azurophilic granules	Specific granules	Gelatinase granules	
ldentifying markers	Azurocidin, Myeloperoxidase (MPO), CD63	Lipocalin 2 (NGAL), CD66b	Gelatinase B (MMP9), CD11b	Albumin, CD45, Mac-1, CD13
Antimicrobial proteins	Defensins, Cap57, Lysozyme	Lactoferrin, Pentraxin 3, Lysozyme, Haptoglobin, Gp91 ^{phox} , Gp22 ^{phox}	Cathelicidin (CAP- 18), Lysozyme, Gp91 ^{phox} , Gp22 ^{phox}	Gp91 ^{phox} , Gp22 ^{phox}
Proteases	Neutrophil elastase, Cathepsin G, Proteinase 3	uPA, MMP8	MMP25, MMP8	MMP25, Proteinase 3
Adhesion molecules		Mac-1, CD66, CD67	Mac-1, CD67	Mac-1, CD67
Receptors	CD63	uPAR, Laminin- receptor, thrombospondin receptor	Ficolin-1	Complement receptor 1, CD14, FCγR, C1q-receptor, formylpeptide receptor
Trafficking and docking	VAMP-1, VAMP- 7, Rab5/Rab27a	VAMP-7	VAMP-2, VAMP-7	VAMP-7, Rab3d

Table 1 Nomenclature and key molecules found within neutrophil granules.

The distinct distribution of individual molecules allows for the delicate differentiation between the three individual granule subtypes as well as the secretory vessels. In addition to the frequently used identifying markers, these granules and vesicles can further be distinguished by the presence or absence of certain antimicrobial proteins, functional proteases and receptors.

1.2.2 Phagocytosis

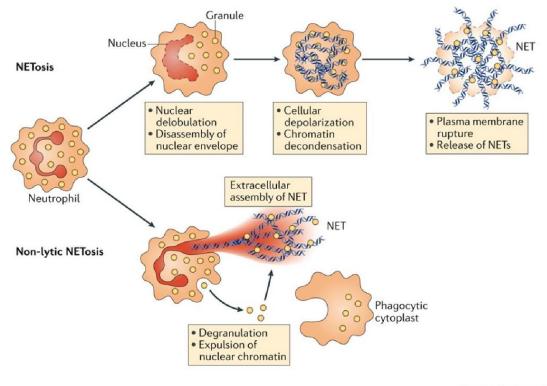
Phagocytosis by neutrophils is less well described compared to other effector functions and the majority of the molecular understanding of phagocytosis is derived from studies based on macrophages.¹²² However, together with macrophages neutrophils belong to professional phagocytes capable of internalizing opsonized bacteria, invading pathogens or dead cells.^{51,122} So far, two general mechanisms of phagocytosis have been identified including the trigger mechanism and the zipper mechanism.^{123,124} Distinct signalling by a limited number of pathogens including for example *Salmonella* or *Shigella*, initiates the trigger mechanism resulting in the formation of plasma membrane protrusions surrounding the engulfed target.^{123,125} On the other hand, the zipper mechanism involves a large number of surface receptors that sequentially bind to target ligands thereby wrapping the target particles of a

great variety of pathogens.¹²⁵ The main receptor relevant for neutrophil phagocytosis is the complement receptor 3, a Fcy class receptor recognizing IgG. Additionally, even though not considered as phagocytic receptors, NOD receptors and TLRs may increase phagocytosis upon activation.¹²³ Phagocytosis is initiated by lateral clustering of complement receptor 3 (CR3 or Mac-1) and regulated by a balance between complement C5a receptor (C5aR or CD88) and CR3.^{110,126,127} Pharmacological inhibition or genetic deletion of C5aR as well as reduced expression of CR3 results in defective phagocytosis in both human and mouse neutrophils with subsequent defective intracellular killing of bacteria.^{127,128} It appears as neutrophils are equipped with an intracellular self-regulating mechanism as certain granule components, including neutrophil elastase (NE), cathepsin G and proteinase 3, are capable of cleaving C5aR.^{106,129} The influence of granule contents is further complexed by its contextdependent involvement. While C5a induces the release of NE and thereby the reduction of C5aR, TLR9 ligation mediates the additional release of proteinase 3, both leading to compromised phagocytosis.¹⁰⁶ Under normal conditions persuading phagocytosis, receptor activation induces lipid remodelling within the cell membrane accompanied by actin cytoskeleton rearrangement. This membrane reorganization is required for proper pathogen/particle internalization within the phagosome.¹³⁰ The initial composition of a fully formed phagosome is not automatically antimicrobial. Phagosome maturation describes the process by which the fully formed, yet not fully functional phagosome acquires the contents essential for intracellular killing.¹²² To fully cover the events during neutrophil phagocytosis, it is noteworthy that this process shares several parallels with the endocytic pathway.¹³¹ One major difference between macrophage and neutrophil phagocytosis is characterized by the fact that conventional endosomes as well as lysosomes are absent in neutrophils and not readily available.¹²² The antimicrobial effect observed in neutrophil phagosomes is derived by their fusion with preformed granules and secretory vesicles.¹³² Similar to other effector functions, phagocytosis impacts neutrophil life-span. Usually, neutrophil apoptosis is accelerated upon phagocytosis of opsonized bacteria in a reactive oxygen species (ROS)dependent manner that results in caspase 8 activation.^{43,133,134} Overall, it is still not fully understood how neutrophils decide whether to phagocytose or to form NETs.¹³⁵

1.2.3 Neutrophil Extracellular Trap formation (NETosis)

Neutrophil extracellular trap (NET) formation was initially described as programmed cell death distinct from necrosis and apoptosis as potent host defence mechanism to trap and kill invading pathogens including a diverse set of bacteria, fungi and viruses.¹³⁶⁻¹⁴¹ Besides neutrophils, other leukocytes such as macrophages, basophils, eosinophils and mast cells were described

to release extracellular DNA resembling extracellular traps.¹⁴²⁻¹⁴⁵ However, increasing evidence accumulates that NET formation, also called NETosis, does not inevitably result in cell death as vital NETosis may also occur without cell lysis.^{139,146-149} Furthermore, NETosis is frequently considered as a double-edged sword of innate immunity as neutrophils are capable of forming NETs also in sterile inflammation thereby eliciting damage to host cells.^{147,150,151} While the presence or increased occurrence of NETs has been associated to various pathologies, thereby demonstrating the pathophysiological relevance of NET formation, the exact molecular and cellular mechanisms underlying this process are not yet fully understood and are still continuously being discovered.¹⁵² NETs mainly consist of condensed chromatin and DNA, forming web-like structures with a pore size of approximately 200 nm.¹⁵³ Nuclear proteins such as histones, a diverse set of granule proteins including NE and MPO, as well as cytosolic proteins and actin are attached to these structures.¹⁵⁴⁻¹⁵⁶



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Figure 2 NET formation pathways (NETosis).

Two major NET formation pathways have been suggested. The classic and more common lytic form, usually termed as NETosis or NET formation is characterized by striking intracellular re- and disassembly of the nucleus and the cytoskeleton, followed by chromatin decondensation and membrane rupture eventually leading to the release of NETs and remnants of the cellular body. During non-lytic NET formation neutrophils only partially degranulate and extrude NET components for extracellular NET assembly. Neutrophils performing this form of NET formation remain capable of phagocytosis. Adapted from Papayannopoulos et al. 2018.¹⁵⁷

1.2.3.1 Neutrophil activation and NETosis initiation

The ultimate prerequisite for NETosis is the preceding activation of neutrophils, as resting neutrophils do not form NETs in non-inflammatory conditions.¹⁴⁸ Neutrophil activation involves a broad spectrum of surface receptors.^{148,158-160} Genetically modified mice representing with a deficiency in Toll-like receptor 2 (TLR2) or complement component 3, or with generally impaired interleukin-1 (IL1)-receptor/TLR signalling were unable to form NETs upon stimulation with Staphylococcus aureus, indicating an involvement in NETosis signalling in terms of specific bacterial stimuli.¹⁴⁸ Furthermore, ligands of tumour necrosis factor (TNF), Fc and G-protein-coupled receptors (GPCRs) were observed to initiate NETs formation.¹⁵⁸⁻¹⁶⁰ In addition to this diverse set of surface receptors, neutrophils express surface proteins such as CD18, a β 2 integrin, which enables them to form NETs upon encountering activated platelets, bacteria or viruses, which further highlights the miscellaneous activating possibilities of neutrophils.^{76,160-162} Besides plasma membrane surface receptors, neutrophils additionally express nucleotide oligomerization domain (NOD)-like receptors which have been shown to induce NETosis upon activation.¹⁵² Additional routes of neutrophil activation circumventing surface receptor ligation have been described for ROS or bacterial toxins, such as nigericin and ionomycin.^{117,163,164} Activation of neutrophils leading to NETosis signalling initiation subsequently leads to increased intracellular calcium concentrations due to the liberation of calcium storages of the endoplasmic reticulum (ER).¹⁶⁵⁻¹⁶⁸ Ligation of Fcy receptors, GPCRs, complement receptors or TLR4 contribute to calcium release from the ER which is usually followed by additional calcium influx upon opening of plasma membrane channels.¹⁶⁵⁻¹⁶⁸ Calcium chelation was observed to impair NET formation in stimuli-dependent manner as well as in dependence of extracellular or intracellular calcium chelation. Extracellular calcium chelation prevents phorbol myristate acetate (PMA)-, IL8-, calcium ionophore- (ionomycin and nicericin) and *Pseudomonas aeruginosa*-stimulated NET formation.^{158,163,169} While intracellular calcium chelation also inhibits IL8, calcium ionophore- and PMA-induced NET release, it does not affect NET formation upon Candida albicans or group B Streptococcus stimulation.^{163,170} Even though the exact cellular processes dependent on increased calcium concentrations during NETosis signalling are not yet entirely understood, increased calcium availability is widely accepted to be a key requirement for fully functional NET formation and release.¹⁵²

1.2.3.2 Morphodynamics

A multitude of morphological changes of neutrophils have been well described in the context of migration to inflamed or infected tissue where high flexibility is inevitable for proper transmigration and extravasation.⁴ Several changes during NETosis initiation have been observed, however it is not yet clear whether the observed morphodynamics are required for functional NET formation.¹⁵² It was observed that upon activation neutrophils spread prior to shedding of plasma membrane microvesicles followed by rounding up.^{171,172} It is suggested that the increased cell spreading is accompanied by the activation or upregulation of extracellular matrix (ECM) surface receptors, which further promotes NETosis.^{76,160,161,173,174} Nevertheless, it was shown that at least upon PMA stimulation, neutrophils enter NETosis despite being seeded on substrate surfaces lacking integrin ligands.¹⁷¹ Plasma membrane microvesicles, shed by activated neutrophils, are annexin V positive and contain cytosolic granules.¹⁷² It was suggested that these microvesicles may serve as messengers contributing to systemic effects, including for example the promotion of thrombosis.¹⁷⁵ Furthermore, it is hypothesized that the presence of annexin V subjects the microvesicles to phagocytosis by macrophages without inducing further inflammatory responses.¹⁷⁶ Despite the lack of complete understanding of the exact role and involvement of the neutrophil-derived microvesicles, it was shown that they contribute to the limitation of bacterial growth, activation of platelets, downregulation of macrophage activation and stimulate endothelial cells to secrete cytokines such as IL6 and IL8.177-180

1.2.3.3 Involvement of kinase signalling pathways

Elevated levels of calcium as well as cytokine engagement has been shown to be involved in the activation of several kinases and cell cycle regulators during NETosis.^{158,181-183} The protein kinase C (PKC), specifically the isoforms PKC α , PKC β 1 and PKC ζ , are activated by calcium and phorbol esters in a phospholipid-dependent manner. PKC serves as critical cell cycle regulator and mediates ionomycin-, PMA-, IL8-, *C. albicans*-, group B *Streptococcus*- and platelet-activating factor-induced NETosis.^{158,163,181,184} Stimulation of neutrophils with PMA additionally revealed the involvement of cyclin-dependent kinase 6, regulating G₁/S phase transition during the cell cycle, as well as the Raf-MEK-ERK MAP kinase pathway.¹⁸² Contrasting, monosodium urate crystal- as well as *S. aureus*-induced NET release is mediated by the SYK-PI3K-mTorc2 pathway.¹⁷⁰ Furthermore, activating mutations in the non-receptor tyrosine kinase Janus kinase 2 (JAK2) were observed to enhance neutrophils tendency to release NETs.¹⁸³ Taken together, this highlights the delicate stimulus-dependent heterogeneity of neutrophil downstream signalling.¹⁵²

1.2.3.4 The role of reactive oxygen species

Reactive oxygen species (ROS) are dually implicated in NETosis, as they are a direct product of NET formation but also serve as activating stimuli for resting neutrophils.^{117,163} In neutrophils ROS are either derived from mitochondria or NADPH oxidase.^{117,163} An increase in ROS production is observed rapidly within 20 minutes after neutrophil stimulation with calcium ionophores, C. albicans, group B Streptococcus, PMA or S. aureus. Similar to calcium and kinase signalling, the production of ROS of different origin is highly stimulus dependent.^{117,152,163} Furthermore, it was proposed that certain stimuli, including PMA or S. aureus seem to strictly depend on NADPH oxidase derived ROS, as neutrophils from patients suffering from chronic granulomatous disease lack functional NADPH oxidase and fail to form NETs upon stimulation.¹⁸⁵ The critical role of NADPH oxidase is further highlighted as inhibition of NADPH oxidase was shown to indirectly decrease histone citrullination by mediating protein arginine deiminase 4 (PAD4) activity, thereby compromising NET formation.^{186,187} NADPH oxidase activation is mainly PKC-mediated but other kinase pathways, such as the c-Raf-MEK-ERK cascade as well as Akt, were shown to promote the assembly of functional NADPH oxidase.¹⁸¹ The mode of action activating NADPH oxidase during NETosis appears to be highly stimulus-specific, since its induction in neutrophils stimulated with parasites was shown to be c-Raf-MEK-ERK signalling-dependent but PKC-independent.¹⁸⁸ It is noteworthy that both, PKC and the c-Raf-MEK-ERK cascade stimulate Mcl-1 expression, which is the main protein in neutrophils exerting an anti-apoptotic effect, which may explain the association with suppressed apoptosis in activated neutrophils dedicated to NETosis.^{181,189}

Contrasting, *Leishmania donovani, Paracoccidioides brasiliensis, C. albicans*, calcium and potassium ionophores, soluble immune complexes, monosodium urate crystals as well as *S. aureus* do not depend on NADPH oxidase, thereby indicating an involvement of mitochondrial ROS (mtROS).^{163,170,173,190-192} Stimulation of NADPH oxidase-deficient neutrophils lead to robust NETosis induction relying on mtROS, which is believed to be produced due to increased calcium concentrations. This mode of action is commonly referred as NADPH-independent NETosis.¹⁸⁹ However, it was shown that mtROS activates NADPH oxidase, thereby further promoting NETosis.¹⁹³ Based on these findings, it was suggested to omit the term "NADPH oxidase-dependent NETosis".¹⁸⁹

1.2.3.5 The involvement of protein arginine deiminase 4 in DNA decondensation

One of the main features distinguishing NETosis from other forms of cell death is marked by chromatin decondensation, which is not seen in necrosis, pyroptosis or apoptosis, where chromatin remains unchanged or becomes condensed.^{194,195} During NETosis, chromatin heterogeneity is lost and decondensation is mediated by posttranslational histone modifications. In contrast to marginally described histone acetylation, citrullination of histones has been well described and is considered to drive NET formation.^{164,196} Furthermore, histone modification by serine protease mediated cleavage is also considered as critical step during NET formation.¹⁹⁷ The PAD family comprises five isoforms, which are expressed in mammals and catalyse citrullination.^{198,199} PAD4 is primarily expressed in granulocytes and is the only subtype that holds a nuclear localization signal.^{200,201} However, precise localization has been observed inconsistently as cytosolic activity was reported, yet visualization of PAD4 showed a nuclear localization.^{172,201,202} It was shown that PAD4 activation occurs in a calcium-dependent manner. Interestingly, in vitro PAD4 enzyme activity required higher calcium concentrations as were observed intracellularly in activated neutrophils, indicating that additional or alternative pathways are contributing to PAD4 activation.²⁰³⁻²⁰⁵ Furthermore, it was suggested that ROS may also contribute to PAD4 activation, however, direct evidence has not been shown yet.²⁰⁶ PAD4 specifically citrullinates histones H3, H4, H2A and the linker histone H1 at different arginines.²⁰⁷⁻²⁰⁹ Neutrophils obtained from PAD4 knockout mice were shown to lack citrullinated histones and fail to undergo NETosis upon a plethora of stimuli.^{187,210-212} In addition to neutrophil activation in an inflammatory milieu, PAD4 was also described to be required during sterile inflammation, including cancer and deep vein thrombosis.^{212,213} Despite questionable specificity, pharmacological inhibition of PADs was described to reduce NET extrusion in primary human and mouse neutrophils. ²¹⁴⁻²¹⁹ However, contradictory reports describe NETosis lacking citrullinated histones upon PMA stimulation and some studies showed no inhibitory effect of pharmacological PAD inhibitors upon bacterial-, PMA- or calcium ionophore-induced NETosis.^{163,184} Additionally, NETosis was observed in PAD4 knockout mice exposed to *C. albicans*.²²⁰ It is suggested that other PAD family members, specifically PAD2 contribute to protein citrullination, thereby substituting potential PAD4 deficiency.²²¹⁻²²³ Even though delayed, chromatin decondensation was observed in PAD4-deficient neutrophil-like HL60 cells.¹⁷² These findings propose that PAD4 enzymatic activity may be critical during a specific phase of decondensation, such as initiation, but not exclusively required for fully functional chromatin decondensation.¹⁵² Nevertheless, it is widely accepted that PAD4 mediated histone citrullination is a prerequisite for end-stage NETosis.^{163,224,225}

1.2.3.6 Cellular adaptations required for the release of NET content

Nuclear as well as cytoplasmic rearrangement during NETosis resembles morphological changes observed during migration and cell division. It involves lamin remodelling and pore formation in the nuclear envelope before it eventually ruptures.^{117,171,226-228} Activation of neutrophils induces local discontinuities of the lamin networks which are believed, even though not yet fully proven, to be mediated by either PAD4 and/or PKC-induced phosphorylation.^{171,227,228} The complete sequence of actions occurring during NETosis are not fully uncovered yet and several controversial observations have been described. These include that nuclear envelope vesiculation occurs during suicidal and vital NETosis.117,139 During classical suicidal NETosis, this vesiculation is believed to mediate the disintegration of the nuclear envelope which subsequently facilitates the release of chromatin into the cytosol.^{117,229} Furthermore, it is hypothesized that the nuclear envelope is vesiculated but not ruptured during vital NETosis. Electron microscopy of stimulated neutrophils suggests that these vesicles bud off of the outer nuclear membrane prior to exocytosis at the plasma membrane.¹³⁹ However, these findings have fuelled controversial discussions as it remains to elucidate how these vesicles are formed without rupturing the inner nuclear membrane yet forming with only a single-membrane.^{152,172} Contrasting, high-resolution, live-cell imaging revealed that the nuclear membrane breaks at various sites instead of being disintegrated by vesiculation.152,172

To ensure proper release of nuclear contents into the extracellular space, a great variety of cytoskeletal rearrangements has to be performed including the disassembly of microtubules, actin filaments as well as vimentin intermediate filaments.^{171,172,182,230} It was shown that the pharmacological stabilization of the actin cytoskeleton, by actin filament polymerization, resulted in compromised NET formation.^{171,172} Contrary, NETosis was also observed to be impaired if drugs inducing actin filament depolymerisation were applied early after neutrophil stimulation, thus suggesting a specifically and temporally requirement of actin destabilization during later stages of NETosis.^{152,171,172}

The molecular mechanisms underlying the extrusion of NET contents into the extracellular space remain mostly unknown. However, it has been suggested that plasma membrane rupture follows a multistep process with progressively increasing membrane permeability.¹⁷² Despite contradictory findings, it was recently proposed that the eventual extracellular DNA release takes place as passive phenomenon due to chromatin-swelling mediated mechanical rupture, rather than being actively mediate as it appeared to be independent of MPO, glycolysis, metabolism and ATP.¹⁷¹

1.2.3.7 Suicidal versus non-lytic NETosis

The tremendous diversity of neutrophil effector functions is further expanded by their capability to undergo a non-lytic form of NETosis retaining other effector functions and not altering their viability. Initially, this phenomenon was described as vital NETosis observed in neutrophils in presence of LPS stimulated platelets.¹⁴⁷ Due to the discrepancy in describing a form of cell death using the term "vital", the Cell Death Nomenclature Committee recommended omitting this specific terminology.²³¹ The extrusion of nuclear content from viable neutrophils was observed in vivo in skin infected with gram-positive bacteria. This process was induced by the activation of the complement system and interaction with TLR2. In this context, nuclear-free neutrophils retained functional chemotactic and phagocytic ability.¹⁴⁸ Furthermore, release of mitochondrial DNA was observed in GM-CSF primed neutrophils upon LPS stimulation in a NADPH oxidase-dependent mechanism.^{145,232} Suicidal, or lytic NETosis and non-lytic NET extrusion can be distinguished by three fundamental differences. Firstly, certain stimuli, including microbial-specific molecular patterns, TLR2 and complement activation, seem to trigger non-lytic NET extrusion in a drastically more rapid fashion.^{139,147,148} Secondly, non-lytic NET extrusion is associated with retaining other neutrophil effector functions such as degranulation and phagocytosis.^{148,173} And lastly, molecular mechanisms required for non-lytic NET release differ as this phenomenon relies on vesicular trafficking of nuclear content to the plasma membrane where it is released eventually.¹³⁹

1.3 Neutrophils and NETs in disease

Neutrophil effector functions, specifically NET formation has been linked to protection in various pathologies as means of host defence. NETs were shown to immobilize pathogens and prevent pathogen spreading as well as penetration into the bloodstream.^{137,148} However, deficiency, mutation or impaired function of several factors required for functional NET extrusion, including for example MPO and NADPH oxidase, were associated with increased disease burden and poor prognosis in various pathologies.^{185,233} Furthermore, patients suffering from defects of neutrophil functions, as seen in chronic granulomatous disease, Chédiak-Higashi disease, leukocyte adhesion deficiencies or the Papillon-Lefévre syndrome present with high incidences of recurrent infections, chronic non-resolving inflammation and hyper-inflammation, mainly due to the incompetency of proper pathogen clearance.²³⁴⁻²³⁸ Accumulating evidence has linked dysregulated and excessive NET occurrence with heavily promoting host damage in infection as well as sterile inflammation and several pathologies including thrombosis, pulmonary diseases, rheumatoid arthritis, systemic lupus

erythematosus, sepsis, inflammatory skin diseases and chronic wounds, diabetes and associated complications, Covid-19 as well as heart failure to name only a few (Figure 3).^{157,189,221,239-241}

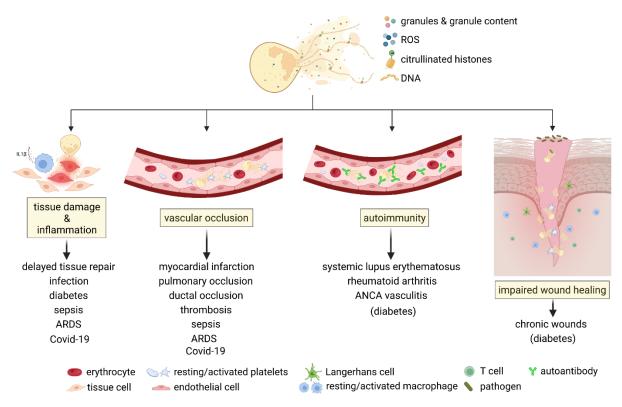


Figure 3 Implications of NETs in pathologies.

NETs and NET-contents are potent drivers of various diseases and pathologic conditions. Direct contact of NETs to tissue induces tissue damage and inflammation. Vascular occlusion is favoured by NETs and NET-platelet aggregates. Continuous exposure to NETs-contents may induce the production of autoantibodies resulting in autoimmunity. Prolonged and/or excessive infiltration and persistence of neutrophils and NETs within wounds is associated with delayed or impaired wound healing. Diseases named in this figure are not necessarily caused by NETs but strongly correlate with NET formation. This figure was created with BioRender.com

1.3.1 Cardiovascular diseases

1.3.1.1 Atherosclerosis

Atherosclerosis onset is marked by endothelial cell damage resulting in the deposition of lipids followed by plaque formation.²⁴² Endothelial cell damage may be neutrophil-induced or a result of hyperlipidemia, which was also shown to promote neutrophilia thereby indicating a feedforward loop with detrimental consequences.^{243,244} NETs were detected in human plaques and in superficial erosions in close proximity of apoptotic endothelial cell clusters.^{242,245} Murine models of atherosclerosis revealed a sterile inflammation-induced production of cytokines triggering neutrophils to form NETs.²⁴⁶ The considerable influence of NETs in atherosclerosis was further proven as atherosclerotic plaques were approximately 3-fold smaller in ApoE/NE/proteinase 3 deficient mice compared to controls and DNase I treatment of control

animals showed a comparable plaque size reduction.²⁴⁶ Inhibition of NET formation by using a PAD4 inhibitor, chlor-amidine, yielded a significant reduction in atherlosclerotic lesion size, delayed thrombosis in carotid artery, decreased neutrophil recruitment and NET formation.^{247,248}

1.3.1.2 Vascular occlusion and thrombosis

NETs formed within the circulation provide a scaffold promoting deep vein thrombosis (DVT).²⁴⁹ It was shown that DVT treatment with PAD4 inhibitors and DNase I prevented thrombosis in mouse models.^{212,250} Release of von Willebrand factor (VWF) and P-selectin from the endothelium recruit neutrophils and initiate NETs formation.^{249,251} P-selectin dependent accumulation of neutrophils further recruits platelets, which are then stimulated to produce thromboxane A2 that subsequently triggers endothelial cells to upregulate intercellular adhesion molecule 1 (ICAM1) surface expression, thereby ensuring firm adhesion of neutrophils.²⁵² Additionally, platelet-derived high mobility group protein B1 (HMGB1) as well as integrins and ROS promote NET formation.^{251,253} Besides mechanical occlusion, Factor XIIa is recruited by NETs and contributes to coagulation and the mobilization of VWF, Factor XIIa and P-selectin containing Weibel-Palade bodies from endothelial cells.^{253,254} Binding of NET derived histones to fibrin and VWF leads to additional recruitment of red blood cells and platelets.^{249,250} Formation of neutrophil-platelet aggregates themselves also trigger a robust pro-coagulant response by enhancing the intravascular tissue factor activity.^{250,255} Endogenous anticoagulants such as thrombomodulin may be degraded or modified by neutrophil proteases, specifically NE, or inactivated by oxidases.^{256,257} Activated platelets are major contributors to thrombosis and are highly responsive to histone mediated activation via TLR2 and TLR4.258,259

1.3.1.3 Acute Myocardial infarction

Acute myocardial infarction (AMI) represents one of the most common cardiac emergencies and usually results of ischemic heart diseases such as coronary artery stenosis, thrombosis or the rupture of a coronary atherosclerotic plaque.^{260,261} AMI is accompanied by cardiac wound healing, necrosis, inflammation and increased leukocyte influx into the infarcted zone.²⁶¹ High numbers of neutrophils infiltrate the infarcted area within hours post AMI where they interact with danger associated molecular patterns (DAMPs), released by necrotic and apoptotic myocytes, which in turn induces an inflammatory response.^{262,263} Tissue damage, reduced resolution of inflammation and poor patient prognosis is associated with excessive neutrophil infiltration or delayed regression due to accumulation of inflammatory mediators.^{19-21,264} In addition to the preceding involvement of NETs in atherothrombosis generation and accumulation in coronary thrombi, NETs continue to interact with platelets during AMI and were shown to further promote platelet activation and express functional tissue factor.²⁶⁵ Specifically histone H3 and H4 were found to induce thrombin generation in a platelet-dependent manner, thereby promoting thrombogenesis.^{265,266} Both, DNA-histone complexes as well as double stranded DNA, which represent key components of NETs, were observed at increased levels at the culprit site and were found to be correlated with the coronary thrombus NET burden. Furthermore, the latter showed a positive correlation with the infarct size, which highlights the detrimental involvement of NETs in the infarcted myocardium. In terms of NET clearance, DNase activity was observed to negatively correlate with the area at risk, infarct size and NET burden.²⁶⁴

1.3.1.4 Anti-neutrophil cytoplasmic antibody associated vasculitis

Vasculitis describes a diverse group of conditions marked by inflammation of blood vessels leading to organ ischaemia and damage. Due to the enormous heterogeneity observed in vasculitis no consensus on exact nomenclature and characterization has been found yet. However, it can be classified broadly according to the size of the vessel involved (e.g. large: aorta; small: intraparenchymal arteries, arterioles, capillaries, venules) and the presence or absence of autoantibodies.²⁶⁷ Anti-neutrophil cytoplasmic antibody (ANCA) associated vasculitis (AAV) describes an exceptional form of vasculitis characterized by tissue damage, endothelial injury and inflammation of blood vessels featuring the loss of tolerance towards the neutrophil granule proteins proteinase 3 (PR3) or MPO.²⁶⁸ AAV is associated with severe organ damage or life-threatening complications and is typically characterized by microvascular endothelial inflammation leading to progressive tissue injury, fibrosis and ultimately to loss of function.²⁶⁸ Several hypotheses have been established which factors could cause AAVs. On the one hand, it appears as genetic predispositions may be involved to a certain extent. However, it was also suggested that increased exposure of neutrophil antigens to DCs may induce loss of tolerance.²⁶⁹⁻²⁷¹ Immunological loss of tolerance of T cells and B cells to neutrophil granule proteins results in the development of ANCAs which not only target PR3 or MPO, but also contribute to neutrophil activation. Neutrophil activation by ANCAs leads to cytoskeletal alterations, generation of ROS and an upregulation of adhesion molecules, resulting in neutrophil accumulation at already stressed tissue sites, where they further promote tissue injury and additionally release autoantigens by degranulation of NET formation.²⁷²⁻²⁷⁵ Released autoantigens are processed by antigen presenting cells with

subsequent antigen recognition by T cells, which eventually results in a vicious cycle of continuous inflammation, ANCA production and neutrophil activation.²⁷⁶ Furthermore, AAVs are associated with an increased risk of cardiovascular events including myocardial infarction, stroke or cardiovascular death and additional dysregulation of either the immune or coagulation system reinforces the risk of venous thromboembolism.²⁷⁷⁻²⁷⁹

1.3.2 Autoimmunity

NETs, specifically certain components such as MPO, DNA, citrullinated proteins and proteinase 3, were proposed to promote autoimmune diseases by providing a source of self-antigens. This phenomenon was first discovered in kidney biopsies from patients presenting with ANCA-associated vasculitis and was further corroborated by studies evaluating NETs deposition in patients suffering from rheumatoid arthritis (RA) and systemic lupus erythematosus (SLE).¹⁵⁷

1.3.2.1 Rheumatoid arthritis

Antibodies targeting citrullinated proteins were detected in two thirds of RA patients.²⁸⁰ RA is a chronic inflammatory disease primarily affecting the joints, however in severe manifestations it is associated with neutropenia, splenomegaly and excessive systemic inflammation.^{280,281} It was shown that neutrophils derived from RA patients are highly prone to spontaneously form NETs as well as upon stimuli, including antibodies targeting citrullinated proteins, TNF-α, IL-17 and immune complexes.²⁸²⁻²⁸⁴ This observation is due to a pre-activation of neutrophils by the rheumatoid factor which drives neutrophils to excessively produce ROS, form NETs and degranulate, thereby further extruding self-antigens.²⁸⁵ Targeting PAD4 in animal models of RA using a pan-PAD inhibitor showed alleviation of disease burden, marked by reduced erosive changes and inflammation.²⁸⁶ Furthermore, a therapeutic scheme using monoclonal antibodies against citrullinated histones showed similar results as PAD4 inhibition in a collagen-induced RA model.²⁸⁷ However, a NETs-dependent mechanism driving RA severity has been challenged by the observation that the excessive citrullination of proteins, present in the synovial fluid of human arthritic joints, resembles citrullination seen in T cell-derived poreforming toxin treated neutrophils.²⁸⁸

1.3.2.2 Systemic lupus erythematosus

SLE is an autoimmune disease characterized by large numbers of autoantibodies, which are mainly directed against nuclear antigens, the generation of immune complexes and abnormal myeloid as well as lymphoid cells.^{222,289} Patients suffering from SLE show reduced NETs clearance, which is thought to result from impaired DNA degradation, due to DNase inhibitors and antibodies binding to NETs.²⁸⁹ NETs are believed to worsen SLE severity by stimulating plasmacytoid dendritic cells (pDCs) to abundantly produce type I interferons via TLR9 and TLR7 signalling.²⁹⁰⁻²⁹² In addition to pDCs, NETs activate macrophages in a feedforward loop resulting in further neutrophil stimulation to produce NETs.²⁹³ It is suggested that the prolonged persistence of NETs as well as individual NETs components lead to complement activation, which in turn further promotes disease activity.²⁹⁴ Studies in a murine SLE model revealed that inhibition of NETs alleviated SLE pathogenesis by additionally reducing organ damage and reduction of vascular injury.^{295,296} NETs targeted therapy may also suppress macrophage activation, thereby limiting immune complex formation, which contributes to the suppression of the adaptive immune response in SLE.²⁹¹

1.3.2.3 Diabetes

Despite highly contrasting causative events, neutrophil effector functions and response to infection are exceedingly dysregulated in both, Type 1 Diabetes (T1D) and Type 2 Diabetes (T2D).²⁹⁷ While T1D represents an autoimmune disorder marked by T cell orchestrated pancreatic ß cell loss accompanied by insulin deficiency, T2D is an acquired disease characterized by defective insulin production in combination with insulin resistance in the liver and skeletal muscle.^{298,299} Patients with diabetes are reportedly at high risk of infection at the urinary tract, respiratory system and especially the skin.^{300,301} Hyperglycaemia and advanced glycation end products are critical factors altering neutrophil function.^{302,303} It was observed that neutrophil phagocytosis as well as chemotaxis were reduced in diabetic patients.³⁰⁴ Other effector functions such as pro-inflammatory cytokine release, ROS production and NET formation, were found to be increased in diabetic patients.^{305,306} Increased NET burden is associated with complications including retinopathy, attendant cardiovascular diseases and impaired resolution of infection, specifically in diabetic foot ulcera.³⁰⁷ Elevated levels of glucose concentrations promote increased rates of NET formation as well as amplified ROS production.^{305,308} This triggers a feedforward loop, since ROS serves as potent trigger of NET formation itself. Furthermore, it was found that NETs persist in patients with diabetes up to one year despite normalisation of their blood glucose levels.³⁰⁹ Potent proteases such as NE

showed higher enzymatic activity in diabetic patients, further exacerbating inflammation.³¹⁰ Furthermore, both, human and murine neutrophils isolated from diabetic patients or experimentally induced diabetes were already primed to undergo NETosis.³⁰⁵ Several therapeutic approaches targeting PAD4 or ROS showed promising results specifically in terms of improved wound healing.^{305,311}

1.3.3 Wound healing

Wound healing is a tightly controlled process marked by four phases: haemostasis, inflammation, proliferation and subsequent remodelling. Dysregulation of either of these phases may result in delayed or impaired wound healing.³¹² One of the first cell types recruited to the wound site are neutrophils.¹⁵ NETs have increasingly gained attention in terms of delayed wound healing.^{305,313,314} While neutrophil granule contents are partly required for functional wound healing, excessive granule and NETs components are linked to impaired wound healing.^{315,316} Chronic wounds show elevated levels of proteases and particularly NE is capable of degrading proteins critical for structural maintenance including collagen, fibronectin and proteoglycans, leading to the disruption of cell connections.^{316,317} MPO contributes to increased NET integrity and stability by crosslinking individual NETs, leading to inflammation and local tissue damage.^{315,318} Other NET components such as histones are capable of integrating into phospholipid membrane bilayers, thereby compromising membrane permeability and integrity, resulting in increased calcium influx and ultimately cell death.^{258,319} Other factors such as NET-derived MMP9 take part in ECM and intracellular matrix degradation, which potentially affects proper reepithelialisation.^{4,320-322} In addition to direct effects, NET aggregates in the vasculature promote vascular occlusion via thrombus formation, leading to insufficient blood supply to the wounded area, which in turn is accompanied by delayed clearance of dead tissue as well as ischemia and ultimately impaired wound healing.^{157,323,324} NETs may also dampen the formation of new capillaries via NEmediated cleavage of wound healing associated growth factors, including platelet-derived growth factor and vascular endothelial growth factor.³²⁵ The deleterious effect of NETs during wound healing is particularly noticeable in diabetic wounds. Hyperglycaemia itself sufficiently primes neutrophils to rapidly form NETs.³⁰⁵ Counteracting the detrimental effect of NETs during wound healing by genetic PAD4 deletion was observed to improved wound healing in diabetic mice significantly.³²⁶ However, hyperglycaemia is not a diabetes specific prerequisite as normoglycaemic, aseptic wound healing as similarly improved upon PAD4 knockdown.³⁰⁵

1.3.4 Cancer

The role of neutrophils and neutrophil effector functions has been described to be rather controversial and highly dependent on the type of cancer as well as disease progression stage and composition of the microenvironment. Neutrophils were reported to exert anti- as well as pro-cancer effects, as they are extremely versatile in adapting to various microenvironmental signals.^{2,327} Several studies described the anti-cancer effect of neutrophils to be dependent on effector molecules, such as nitric oxide or NE, which either delay tumour growth and induce killing or are capable of directly killing cancer cells.^{328,329} Furthermore, neutrophils were shown to indirectly limit tumour progression at early stages by stimulating macrophages, which in turn trigger the release of T cell derived IFNy.³³⁰ The high plasticity of neutrophils was highlighted by a study in early-stage human lung cancer, where immature neutrophils differentiated into a hybrid state, expressing an antigen-presenting-cell phenotype and cross-presented tumour antigens to T cells, thereby inducing an anti-tumour effect.³³¹ In contrast, accumulating evidence suggests that neutrophils may also contribute to tumour incidence, formation and growth, due to maintaining a chronic inflammatory state, genome instability, promoting angiogenesis and inducing DNA damage.^{327,332-335} Neutrophils were also shown to amplify DNA damage and persuading cancer initiation in chemically induced tumorigenesis such as cigarette smoke.³³⁶ Furthermore, neutrophil-derived ROS were shown to favour oxidative DNA damage mutations leading to intestinal cancer growth.³³⁷ Neutrophils may exert proangiogenic effects, as the release of their vascular endothelial growth factor (VEGF)-reservoir stimulates neo-angiogenesis, thus promoting tumour growth.³³⁸ Additionally, neutrophils may indirectly promote angiogenesis by releasing MMP9 which degrades the extracellular matrix, thereby forcing mesenchymal cells to secrete VEGF.³³⁹ Interactions between circulating tumour cells and neutrophils were described to promote cycle progression as well as the metastatic capacity.³⁴⁰ Studies showed that neutrophil depletion was accompanied by a reduction of complete inhibition of metastasis.341

1.3.5 Acute respiratory distress syndrome and Covid-19

Acute respiratory distress syndrome (ARDS) represents a life-threatening respiratory pathology with increasing incidence and mortality rates.^{342,343} Its severity was recalled upon the recent outbreak of Covid-19 as 29-42% of Covid-19 patients developed ARDS and 15-52% of these patients died.^{344,345} ARDS patients typically developh extensive pulmonary oedema, diffuse bilateral pulmonary infiltrates and hypoxemia. Lungs of ARDS patients are marked by increased vascular permeability, leading to protein-rich fluid leaking into the alveolar space

and tremendous accumulation of activated immune cells, which compromises gas exchange and results in respiratory failure.^{346,347} One hallmark of ARDS is the excessive infiltration of neutrophils in the inflamed lung.³⁴⁸ Circulating neutrophils follow a chemokine gradient, released by damaged pulmonary tissue. Neutrophil recruitment and activation is not limited to either exogenous or endogenous inflammatory stimuli, nor to infectious or sterile inflammatory stimuli, thereby facilitating high levels of activation and promoting a potent neutrophil-driven immune response resulting in damaged tissue and lung dysfunction.³⁴⁹⁻³⁵¹ Accumulating NETs were observed to obstruct small airways and damage alveolar capillaries.³⁵² The role of neutrophils and specifically NETs during Covid-19 is highly complex and was shown to indirectly promote disease pathogenesis.²⁴⁰ Patients with Covid-19 show endothelial inflammation, which in turn promotes coagulation as well as thrombus formation.^{353,354} NETs associated biomarkers are elevated in the circulation during early infection response and are associated with disease severity and thrombosis.³⁵⁵ NETs-mediated thrombosis during Covid-19 was proposed to be complement dependent. Covid-19 activates the complement cascade leading to thrombin mediated tissue factor (TF) expression in neutrophils, which results in TF carrying NETs with pro-coagulatory properties.³⁵⁶ Furthermore, it was observed that live as well as heat-inactivated SARS-CoV-2 virus is capable of inducing NETs formation, most likely via binding to angiotensin converting enzyme 2 receptor on neutrophils.^{357,358} Additionally, NETs were found to increase virus infectivity as NE can cleave S protein thereby facilitating virus entry into the cell.³⁵⁹

1.4 Pharmacological targeting of NETs

Indirect immunomodulatory inhibition of NET formation was observed in several studies using acetylsalicylic acid (Aspirin), which is an anti-thrombotic, non-steroidal drug with antiinflammatory effects.^{360,361} It inhibits thromboxane A2, thereby counteracting platelet aggregation and activation.³⁶² Upon activation, platelets show increased surface expression of P-selectin and secrete several mediators, such as platelet factor 4, CCL5 and HMBG-1, thus facilitating neutrophil binding, subsequent activation and NET formation.^{147,251,363} Targeting platelets or blocking platelet-neutrophil interactions may serve as indirect anti-NET therapeutic approach. This was shown to be efficient in an endotoxin-triggered acute lung injury model, where pre-treatment with aspirin lead to reduced intravascular NET formation and decreased lung injury.^{364,365} Another mechanism to indirectly prevent NET formation is the therapeutic administration of thrombomodulin, which is capable of limiting procoaculant responses and prevents NET formation in neutrophils co-cultured with activated platelets.³⁶⁶ Cyclosporine A, an immunosuppressive drug, directly inhibits NET formation by inhibiting the calcineurin pathway.³⁶⁷ It was shown that a combinatory therapy of cyclosporine A together with ascomycin yielded potent inhibition of NET formation *in vitro*.¹⁵⁸ However, the benefit of strong immunosuppressant drugs targeting NETs is questionable as they are accompanied by potentially severe adverse effects and predisposing patients to recurring and more severe infections.³⁶⁷

Direct and irreversible inhibition of PAD4 activity, and thus reduced or inhibited NET formation, is mediated by chlor-amidine.³⁶⁸ Systemic therapeutic administration in a murine model of SLE protected animals from NET-induced vascular damage, kidney injury as well as endothelial dysfunction.²⁹⁶

Prostaglandins (PGs) are endogenously synthesized enzymes that were attributed to exert both anti- as well as pro-inflammatory effects.³⁶⁹ PGE2 was shown to counteract PMA-induced NET formation by modulating EP2 and EP4.³⁷⁰ Defective intracellular bacterial killing was observed in a PGE2 overproducing mouse model, while EP2 and EP4 antagonist treatment could restore NET formation, thus suggesting that blocking PGE2-EP2 or PGE2-EP4 signalling axis restores NET formation.³⁷¹

Certain complement components were shown to interact with neutrophils and trigger NET formation resulting in a vicious cycle as NET products can in turn activate the complement cascade.³⁷² A peptide inhibitor of complement C1 was observed to reduce MPO activity and therapeutic complement inhibition is already successfully employed in paroxysmal nocturnal hemoglobuinuria, a rare disease characterized by a variety of symptoms including haemolytic anemia, thrombosis, and renal insufficiency.^{373,374}

The commonly used antidiabetic drug metformin induces AMP-activated protein kinase activity, which in turn prevents mitochondrial ROS production via mammalian target of rapamycin (mTORC1) inhibition.³⁷⁵ *In vitro* studies revealed that metformin reduces overall NETs concentrations and particularly cell free DNA, proteinase 3, histones and elastase and additionally prevents the activation of NOX in neutrophils by preventing PKC-βII membrane translocation.^{376,377}

DNase treatment is widely used in *in vitro* experiments and despite not preventing NET formation per se, it was observed to potently alleviate disease burden, by reducing NET-mediated tissue damage in different animal models.^{305,378,379}

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2 Cell-based therapy in regenerative medicine

The importance of regenerative medicine has increasingly gained attraction due to several studies providing promising results, meeting the aim of repairing or replacing damaged cells or organs with potentially regaining normal function.³⁸⁰ Bone marrow transplants used to treat immune system or blood disorders are the most established forms of stem cell therapy.³⁸¹ Stem cell-based therapies comprise any therapeutic intervention for a disease that involves any type of human stem cells, including induced pluripotent stem cells (iPSCs), adult stem cells and embryonic stem cells (ESCs).³⁸² However, despite promising results, stem cell therapy is accompanied by several limitations. One of the biggest concerns is reflected by ethical conflicts using ESCs but also iPSCs, arising from the fact that processing of human embryos is required as well as the unlimited differentiation capacity potentially leading to tumour formation, respectively.³⁸⁰ The efficacy of stem cell therapy was further denounced by initial clinical trials in the setting of AMI, revealing only minor clinical improvements upon autologous progenitor cell treatment.³⁸³ Several studies yielded contradictory results, reporting on the one hand reduced risk of mortality as well as reduced re-hospitalization in patients suffering from heart failure upon autologous cell therapy, while other studies revealed no beneficial effect associated with stem cell therapy in AMI.^{384,385} Mesenchymal stromal cells (MSCs), also referred as mesenchymal stem cells, have been of high interest in terms of therapeutic application due to their immune regulatory function and high safety profile.³⁸⁶ It was shown that MSCs do not exert immunosuppressive functions at baseline, but require microenvironmental cues, including inflammation, to adopt an immunosuppressive phenotype.³⁸⁶ Despite initially promising results, 28% of registered clinical trials have been completed and results were reported for only 2% of the completed clinical trials.³⁸⁶ Immunoselection of mesenchymal precursor cells (MPCs) for STRO-1 expression, which is associated with high co-expression of STRO-3, was shown to be superior to unspecific adherence-isolated MSCs in terms of paracrine activity and thus potential cardiovascular therapeutic approaches.³⁸⁷ Pluripotent, self-renewing MPCs with a high capacity for differentiation and proliferation were demonstrated to exert beneficial effects in animal models of MI by improving left ventricular (LV) function accompanied by reducing infarct expansion upon targeted delivery. Furthermore, already low concentrations of directly intra-myocardially injected MPCs showed therapeutic effects and resulted in alterations of MMP tissue inhibitor and MMP levels as well as collagen dynamics.³⁸⁸ The beneficial effect of immunoselected STRO-3 positive MPCs was further demonstrated in a large animal model of monoarthritis, as MPC-treated animals showed reduced lymphocytic and monocytic infiltration of synovial tissues, decreased cartilage erosions and overall decreased clinical signs of joint pain and swelling as well as decreased lameness.³⁸⁹ Further investigation of the cardioprotective effects of STRO-3 positive MPCs in a rat model of MI revealed that not only the injection of MPCs but also the treatment with MPC-conditioned media decreased fibrosis and myocyte apoptosis, preserved LV dimensions and function and promoted neovascularization.³⁹⁰

The initial theory that transplanted stem cells would incorporate at injured sites and transdifferentiate into the required cell types thereby exerting tissue regenerative effects was further tremendously challenged by studies reporting that only a minor percentage, 1-5% of injected stem cells, engraft at the desired tissue site.³⁹¹⁻³⁹⁵ It was shown that the majority of injected cells was sequestered in the lymph nodes, bone marrow, lung, liver and spleen already a few hours post infusion.^{393,396} These findings raised the question how the beneficial effects observed in stem cell therapy were obtained. Gnecchi et al. shifted the theory from direct stem cell induced immunomodulation towards a secretome-assisted effect, which does not solely rely on direct cell-cell interactions, but is supported by paracrine factors released from infused cells.^{397,398} Conditioned media from mesenchymal stem cells, exposed to hypoxic stress, was shown to significantly decrease hypoxia-induced cell death in ventricular cardiomyocytes of rats.^{397,399} Shortly thereafter, it was shown that the stem cell secretome deviates marginally from supernatants derived from peripheral blood mononuclear cells (PBMCs).⁴⁰⁰ The controversy around stem cell therapy was further expanded when the immunomodulatory effect was proposed to be derived from the secretome of apoptotic stem cells and may even be similar to the secretome of any apoptotic cell type.^{399,401} Apoptosis, representing a tightly regulated active process, is critically involved in the maintenance of homeostasis and constant self-renewal. It is known to drive strong immunomodulatory effects by directly secreting immunocompetent mediators or indirectly by activating dendritic cells and phagocytes.³⁹⁹ A systemic decrease of inflammatory concentrations as well as a release of anti-inflammatory cytokines is observed upon infusion of apoptotic cells.^{402,403} The broad spectrum of signals released by apoptotic cells serves to alarm their surrounding environment of dysregulated homeostasis or danger to promote immune responses, survival and stimulate cell proliferation.399

Using the secretome of stem cells instead of direct cell transfusion allows to circumvent stem cell therapy associated risk factors such as immune rejection and teratoma formation.⁴⁰⁴

2.1 Stressed PBMCs and their secretome

2.1.1 Cardiology

Ankersmit et al. demonstrated that an LPS-induced pro-inflammatory phenotype in PBMCs and monocytes was reduced by treatment with apoptotic γ-irradiated PBMCs.⁴⁰⁵ The injection of irradiated PBMC suspensions in an *in vivo* rodent model of AMI resulted in reduced infarct size and improved functional parameters including end-systolic and end-diastolic diameters as well as ejection fraction.⁴⁰⁵ Lichtenauer et al. sought to further elucidate these findings and investigating the pleiotropic effect of apoptotic PBMCs using the same rat model of AMI.⁴⁰⁶ Irradiated PBMCs were injected either intravenously or intramyocardially upon onset of ischemia. Similar to previously reported results, treatment with irradiated PBMC cell suspensions yielded significantly reduced infarction sizes accompanied with improved remodelling post AMI. Furthermore, an increased homing of cells associated with a regenerative phenotype, including c-kit, IGF-1, FGF-2 and FLK-1 positive cells as well as macrophages, was observed. Ventricular remodelling appeared to be reduced due to a change in the ratio of collagenous and elastic fibres, favouring reduced scar formation.⁴⁰⁶

Interestingly, already in the first manuscript by Ankersmit et al, it was shown that the application of conditioned medium of irradiated PBMCs (PBMCsec) increased the production of VEGF and MMP9 in human primary keratinocytes and fibroblasts. This finding attributes anit-inflammatory and pro-angiogenic properties to PBMCsec for the first time and suggests a promising treatment option for AMI.⁴⁰⁵

Furthermore, this observation led to the development of a cell-derived yet cell-free therapeutic approach, in which the sole effects of paracrine factors, released by irradiated PBMCs (PBMCsec or APOSEC), were evaluated in an experimental AMI model in rats as well as in a porcine closed chest reperfused AMI model.⁴⁰⁷ In both AMI models, a single intravenous dose of PBMCsec yielded a significant reduction of myocardial scar tissue. Furthermore, an improved cardiac output, higher ejection fraction and decreased infarct size was also observed in the porcine AMI model. *In vitro* experiments with primary human cardiomyocytes showed that treatment with PBMCsec prevented starvation-induced cell death. PBMCsec activated pro-survival signalling cascades, including c-Jun, CREB, Erk1/2 and AKT, while simultaneously increasing anti-apoptotic factors such as BAG1 and Bcl-2.⁴⁰⁷ To further unravel the mode of action of PBMCsec during AMI, the effect of PBMCsec on platelets and microvascular obstruction (MVO) was assessed in a porcine closed chest reperfused AMI model.⁴⁰⁸ The animals were treated with a single intravenous dose of PBMCsec 40 minutes post occlusion. Magnetic resonance imaging revealed significantly reduced areas of MVO in

PBMCsec treated animals compared to the control group. Furthermore, improved ventricle contraction capacity as well as ST segment resolution accompanied by reduced ventricular arrhythmias were observed in the PBMCsec treatment group. Platelet activation, representing a critical contributor to MVO, was observed to be significantly reduced, characterized by decreased levels of sCD40L, TSP-1, sP-selectin and PF-4. Additional in vitro experiments with isolated human and porcine platelets further corroborated the in vivo data, since despite strong activating stimuli, platelet aggregation was prevented by PBMCsec treatment. This mechanism was attributed to PBMCsec-induced upregulation of vasodilator-stimulated phosphoprotein (VASP).⁴⁰⁸ Moreover, Pavo et al. showed an increased homing of CD31⁺ cells at the border zone of infarcted area in a porcine closed chest AMI model, suggesting a PBMCsec-induced enhancement of angiogenesis, accompanied with increased accumulation of endogenous cardiac stem cells.⁴⁰⁹ Gene expression patterns in infarcted, border zone and remote myocardium revealed strong downregulation of lipid metabolism-, inflammation- and apoptosis-associated genes such as caspase-1, stromal derived factor 2-like protein 1, TNFα, arachidonate 15-lipoxygenase and claudin 3. In parallel, PBMCsec induced the upregulation of angiogenic growth factors such as insulin-like growth factor, as well as Kruppel-like factor which is known to function as a regulator of homeostasis, vascular tone and exert antiatherogenic effects.⁴⁰⁹ Attempting to unravel the underlying transcriptional changes and pharmacodynamics induced by PBMCsec treatment, transcriptomic analysis of the perfused heart, transition zone and non-perfused heart, as well as liver and spleen, revealed highly organ-specific alterations.⁴¹⁰ Overall, a uniform trend towards the downregulation of proinflammatory mediators was observed. PBMCsec induced strong expression of genes crucial for cardiomyocyte function in the infarcted area.⁴¹⁰ The immunomodulatory effects of PBMCsec were further addressed by Hötzenecker et al. in an experimental autoimmune myocarditis mouse model mimicking critical aspects of human inflammatory dilated cardiomyopathy.⁴¹¹ Intraperitoneal injection of PBMCsec lead to almost complete inhibition of myocarditis characterized by sparse lymphocytic infiltration and absence of apoptotic or necrotic cardiomyocytes. Circulating levels of autoantibodies were partially reduced by PBMCsec treatment. Despite no statistical significance, a trend of reduced amounts of pro-inflammatory cytokines was observed in the PBMCsec group. One substantial finding of this study was that PBMCsec induces caspase-8-dependent apoptosis in CD4⁺ T cells both in vitro and in vivo and lymphocytes derived from treated animals did not show a proliferative response upon stimulation.411

2.1.2 Neurology

Based on the early findings that PBMCsec ameliorates ischemia-induced tissue damage in AMI, Altmann et al. assessed the impact of rat- and human-PBMCsec treatment on ischemic lesion volumes in a rat middle cerebral artery occlusion model.^{405-408,412} The right hemisphere was occluded and animals received either a single intraperitoneal dose of PBMCsec 40 minutes post-ischemia induction or an additional dose 24 hours post-surgery. Both treatment groups, allogenic PBMCsec and xenogenic PBMCsec, showed comparably reduced infarction volumes and improved neuroscores. Furthermore, cytoprotection mediating signalling cascades, such as Akt, c-Jun, Erk1/2 and CREB were activated in Schwann cells and astrocytes treated with PBMCsec in vitro. Administration of PBMCsec to cultured astrocytes and neurons lead to a dose dependent activation of CREB phosphorylation and increased length of newly formed neuronal sprouts.⁴¹² In a second attempt, the effect of PBMCsec on spinal cord injury was investigated.⁴¹³ Neurological damage and deterioration is frequently induced upon spinal cord injury by an inflammatory response, ischemia and increased oxidative stress.⁴¹³ Administration of PBMCsec in a rat spinal cord injury model revealed substantially improved functional recovery and motor function. 28 days after trauma the PBMCsec-treated animals showed significantly smaller cavity formation marked by reduced areas of white matter. Furthermore, acute axonal injury was reduced and an increased vascular density was observed in the spinal cord. A reduction of oxidative stress was characterized by reduced expression levels of inducible nitric oxide synthase.⁴¹³

2.1.3 Wound healing

Additionally, the beneficial effect of the secretome of non-stressed PBMCs was assessed in a full thickness wound mouse model.⁴¹⁴ Cell culture supernatants of freshly isolated PBMCs, cultured for 24 hours were applied to 6 mm punch biopsy wounds daily for three days and resulted in significantly elevated wound closure. Furthermore, treatment of primary human keratinocytes and fibroblasts induced migration but not proliferation *in vitro*. In line with previous findings, a tremendously enhanced infiltration of CD31⁺ cells was observed, suggesting increased angiogenesis.^{409,414} Application of the secretome of non-stressed PBMCs on endothelial cells further induced tube-formation and proliferation in an *in vitro* matrigel-based assay.⁴¹⁴ Hacker et al. further investigated the effect of the secretome of non-stressed PBMCs and PBMCsec in a full-thickness burn injury model in pigs.⁴¹⁵ Comparing the secretome of non-stressed and irradiated PBMCs during wound closure of burn wounds revealed a superiorly beneficial effect of PBMCsec in terms of wound healing. PBMCsec

treated wounds showed increased epidermal thickness, more advanced epidermal differentiation as well as increased numbers of rete ridges. Again, an enormously increased accumulation of CD31⁺ cells was observed in PBMCsec treated wounds.⁴¹⁵ Furthermore, the beneficial effects of PBMCsec on wound healing were further assessed in a rodent epigastric flap model.⁴¹⁶ A single dose of PBMCsec in combination with fibrin sealant was administered intraoperatively immediately prior to wound closure and led to significantly reduced tissue necrosis rate compared to control or fibrin sealant only groups. While no difference was observed in the amount of lymphatic vessels, PBMCsec treatment resulted in a significant increase in von Willebrand Factor positive blood vessels 7 days postoperative. It was suggested that the beneficial effect of PBMCsec is exerted by reducing post-ischemic inflammation additionally to increased re-vascularization.⁴¹⁶ One critical factor during wound healing is a sufficient antimicrobial defence response. Robust antimicrobial activity against certain gram-negative and gram-positive bacteria, all known to be critically involved in the pathogenesis of diabetic foot ulcers, was observed.⁴¹⁷ The secretome of non-stressed PBMCs already displayed modest antimicrobial activity towards the gram-negative bacteria Pseudomonas aeruginosa and Escherichia coli as well as gram-positive bacteria as for example Staphylococcus aureus, yet PBMCsec exhibited significantly stronger effects on these strains and inhibited their growth significantly.⁴¹⁷

2.1.4 Components of the secretome

PBMCsec consists of a plethora of components including proteins, lipids, DNA, extracellular vesicles (EVs), which were attributed to exert regenerative effects.⁴¹⁸ Wagner et al. sought to dissect the molecular composition of EVs found in the secretome of non-stressed as well as irradiation-stressed PBMCs and their impact on wound healing.⁴¹⁸ Irradiation of PBMCs strikingly changed the composition, number and size of EVs and in contrast to non-irradiated PBMC secretome-derived EVs their molecular components exhibited strong association with regenerative processes. Irradiated PBMCs released higher rates of EVs containing a broad set of native as well as oxidized bioactive phospholipids. *In vitro* aortic ring sprouting assays as well as reporter gene assays revealed a reduced potency of individual subfractions of PBMCsec, including EVs, proteins and lipids, compared to the whole PBMCsec. Furthermore, PBMCsec was first applied in the context of diabetic wound healing in a full-thickness skin wound mouse model using LepR^{db/db} diabetic mice. Topical administration of PBMCsec accelerated wound closure and a significantly reduced wound area was observed 25 days post wounding. Contrasting previous findings, enhanced angiogenesis marked by increased homing of CD31⁺ cells was not observed.⁴¹⁸ Furthermore, it was observed that irradiation not

only changes the PBMC-derived EVs of the secretome but also induces changes in both miRNA and mRNA profiles. ⁴¹⁹ A time-dependent alteration in miRNA and mRNA expression was reported. Comparing miRNA and mRNA expression data revealed a negative correlation between mRNA-miRNA and lead to the identification of a significantly downregulated transcription factor, the hepatic leukaemia factor, which together with a plethora of other regulated gene sets is critically involved in the modulation of irradiation-responsive pathways including endocytosis, MAPK signalling, cytokine-cytokine interactions and apoptosis.⁴¹⁹ More in depth analysis of the secretome further revealed that irradiation induces a tremendous alteration in gene expression patterns in PBMCs coding for secreted proteins.⁴²⁰ Gene ontology assessment showed strong correlation of the altered genes with biological processes such as wound healing, leukocyte trafficking regulation and angiogenesis. Comparing PBMCsec with the secretome of non-stressed PBMCs further revealed drastically elevated levels of bioactive lipids, including triglycerides, cholesterol esters, free fatty acids, cholesterol, cholesterol sulfate and phospholipids. Particularly phospholipids appear to be highly susceptible to irradiation-induced changes, specifically those with intact but oxidized sn-2 chains.420

2.1.5 Irradiation induced apoptosis and necroptosis

PBMCsec represents a rather complex compound with several different cell types contributing to the resulting secretome. Simader et al. investigated whether certain subpopulations, including B cells, natural killer cells, CD4⁺ and CD8⁺ T cells as well as monocytes, account for the tissue-regenerative effect.⁴²¹ Individual isolated and purified cell populations were exposed to high-dose γ-irradiation, and subsequently their secretomes analysed using transcriptomics. It was demonstrated, that each cell type responds to γ-irradiation by a distinct cytokine production profile, death receptor signalling as well as different pro-angiogenic pathways.⁴²¹ Furthermore, it was observed that the beneficial effect by PBMCsec requires interactions of the individual subsets. Moreover, Simader et al. revealed that γ-irradiated PBMCs not only undergo apoptosis, but also necroptosis. In depth analysis showed that inhibition of apoptosis abrogated the previously observed effect. TNF receptor superfamily member 1B was identified as the key molecule of necroptosis in PBMCs upon γ-irradiation.⁴²¹

2.1.6 Immunomodulatory effects

In addition to improved wound healing, PBMCsec was found to reduce the inflammatory response and cellular infiltration in dendritic cell (DC)-mediated skin inflammation in mice.⁴²² PBMCsec prevented the differentiation of monocyte-derived DCs which was characterized by reduced expression of DC markers including MHC class II, CD11c and CD1a. DC maturation, lipopolysaccharide-induced cytokine secretion, DC-mediated immune cell proliferation as well as antigen uptake was significantly reduced upon treatment with PBMCsec. The presence of PBMCsec during monocyte-derived DC differentiation resulted in an impaired capability to prime naïve CD4⁺ T cells into both T_{H1} and T_{H2} cells. In situ analysis of skin further revealed a modified DC phenotype. The key discovery was that these modifications appear to be based on PBMCsec-derived lipid-mediated immunomodulatory changes.⁴²² In another study, the immunomodulatory effect of PBMCsec on mast cell and basophil activation in the context of IgE-mediated hypersensitivity was investigated.⁴²³ Experimentally induced mast cell degranulation in mouse ears was reduced by topical administration of PBMCsec. Several genes involved in Fc-receptor signalling as well as immune cell degranulation were significantly downregulated in murine mast cells. PBMCsec treatment of activated primary human dermal mast cells robustly inhibited α-IgE- and compound 48/80-induced mediator release in vitro. In addition to the suppression of mast cell degranulation, allergen driven activation of basophils derived from allergic donors was attenuated by PBMCsec treatment in vitro. Transcriptomic analysis of PBMCsec-treated basophils revealed a similar pattern including strong downregulation of gene sets relevant for Fc-receptor signalling and immune cell degranulation. In this study, lipids were found to be the major contributors of the observed immunomodulatory effects.423

Taken together, PBMCsec has been attributed a diverse set of mode of actions positively influencing various pathologies such as AMI, wound healing and inflammatory responses (Figure 4).

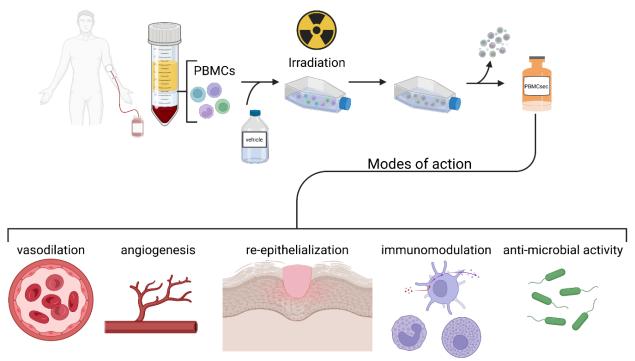


Figure 4 Modes of action of PBMCsec.

During the cultivation of irradiated PBMCs a plethora of paracrine factors is secreted. These factors have been attributed several modes of action with overall cytoprotective effects. The so far investigated modes of action include vasodilation, angiogenesis, re-epithelialization, immunomodulation of various cell types as well as anti-microbial activity. This figure was created with BioRender.com

2.1.7 Clinical trial

The first in-human application of PBMCsec was performed in the MARSYAS I trial (therein referred as APOSEC), a randomized, single-centre, placebo-controlled, double-blinded phase 1 trial. Healthy male volunteers received two 4 mm punch biopsy wounds on their upper arm and were treated with both placebo and autologous PBMCsec in NuGel for 7 consecutive days. The participants were randomly assigned into either a high dose or low dose group. The primary interest was focused on investigating the tolerability of PBMCsec followed by the impact on wound closure and re-epithelization. Recorded adverse events were all characterized as mild and unlikely to be related to the treatment. The main limitation of the study was the short intervention time which restricted the assessment of wound closure which revealed no significant differences.⁴²⁴

3 Aims of this thesis

Dysregulated NET formation is known to critically contribute to host damage in infectious conditions and (sterile) inflammation. The progression of several pathologies is closely associated with increased NET formation.¹⁵⁷

Previous studies analysing the effect of PBMCsec in a diverse set of pathological conditions already provides deeper insights into potential mode of actions. Particularly the immunomodulatory effect is of high interest.^{418,420,422,423}

However, the effect of PBMCsec on neutrophils in the context of NET formation has not been investigated so far.

The main aim of this thesis was to evaluate to what extent NET formation is influenced by PBMCsec treatment.

Furthermore, we aimed to unravel potential deviations in the potency of the PBMCsec subfractions to influence NET formation.

Finally, we sought to identify the mode of action by which PBMCsec influences NET formation.

RESULTS

1. Prologue

It was previously shown that PBMCsec exerts various anti-inflammatory, cytoprotective, immunomodulatory and pro-angiogenic effects in a diverse set of pathologies.³⁹⁹ However, the influence of PBMCsec on neutrophils, particularly the formation of NETs, has not been elucidated yet, despite prominent involvement of neutrophils in the vast majority of investigated pathologies.¹⁵⁷

The main objective of this thesis was to evaluate whether PBMCsec harbours any inhibitory capacity towards NET formation. Furthermore, we sought to investigate if certain substance classes present in PBMCsec are responsible for these effects and pursued to analyse whether molecular mechanisms critical for NET formation may be impaired.

1.1 Paper



Article



The Effect of Paracrine Factors Released by Irradiated Peripheral Blood Mononuclear Cells on Neutrophil Extracellular Trap Formation

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Abstract: Neutrophil extracellular trap (NET)-formation represents an important defence mechanism

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Citation: Klas, K.; Ondracek, A.S.; Hofbauer, T.M.; Mangold, A.; Pfisterer, K.; Laggner, M.; Copic, D.; Direder, M.; Bormann, D.; Ankersmit, H.J.; et al. The Effect of Paracrine Factors Released by Irradiated Peripheral Blood Mononuclear Cells on Neutrophil Extracellular Trap Formation. Antioxidants 2022, 11, 1559. https://doi.org/10.3390/ antiox11081559

Academic Editors: Mercedes Camacho Pérez de Madrid and Josep Julve

Received: 29 July 2022 Accepted: 10 August 2022 Published: 11 August 2022

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Copyright © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). for the rapid clearance of infections. However, exaggerated NET formation has been shown to negatively affect tissue-regeneration after injury. As our previous studies revealed the strong tissueprotective and regenerative properties of the secretome of stressed peripheral blood mononuclear cells (PBMCsec), we here investigated the influence of PBMCsec on the formation of NETs. The effect of PBMCsec on NET formation was assessed ex vivo in ionomycin stimulated neutrophils derived from healthy donors using flow cytometry, image stream analysis, and quantification of released extracellular DNA. The effect of PBMCsec on molecular mechanisms involved in NET formation, including Ca-flux, protein kinase C activity, reactive oxygen species production, and protein arginine deiminase 4 activity, were analysed. Our results showed that PBMCsec significantly inhibited NET formation. Investigation of the different biological substance classes found in PBMCsec revealed only a partial reduction in NET formation, suggesting a synergistic effect. Mechanistically, PBMCsec treatment did not interfere with calcium signalling and PKC-activation, but exerted antioxidant activity, as evidenced by reduced levels of reactive oxygen species and upregulation of heme oxygenase 1 and hypoxia inducible-factor 1 in PBMCsec-treated neutrophils. In addition, PBMCsec strongly inhibited the activation of protein arginine deiminase 4 (PAD4), ultimately leading to the inhibition of NET formation. As therapeutics antagonizing excessive NET formation are not currently available, our study provides a promising novel treatment option for a variety of conditions resulting from exaggerated NET formation.

Keywords: neutrophil; neutrophil extracellular traps (NETs); PAD4; ROS; secretome; peripheral blood mononuclear cell secretome

1. Introduction

Neutrophil granulocytes represent the main population of circulating leukocytes in the blood [1]. They exert a plethora of functions critical for maintaining immune homeostasis, and their contribution to immune regulatory mechanisms is of vital importance during infectious conditions [2]. Being amongst the first cell populations to be recruited to a site of infection, they use a broad machinery of defence mechanisms, including the production of reactive oxygen species (ROS), excretion of cytotoxic granules, phagocytosis of pathogens, and the formation of neutrophil extracellular traps (NETs), to fight invading

Antioxidants 2022, 11, 1559. https://doi.org/10.3390/antiox11081559

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pathogens [3]. In addition to these functions, NET formation represents another potent defence mechanism for the elimination of pathogens [4]. Neutrophils are equipped with a vast array of surface receptors, including Toll-like and NOD-like receptors, G-protein coupled receptors, cytokine receptors, as well as Fc and complement receptors, rendering them highly responsive to a multitude of stimuli [5]. Some of these stimuli, such as ROS or bacterial toxins, are potent inducers of NETs and operate independently of the classical neutrophil activation pathways via surface receptors [6-8]. After induction of NET formation, intracellular Ca2+ levels increase due to influx and its release from the endoplasmic reticulum, promoting protein kinase C (PKC) activation and phosphorylation of Gp91^{phox} [9]. Activation of PKC, in turn, leads to the assembly of functional NADPH oxidase, which generates reactive oxygen species (ROS) [10]. Furthermore, increased Ca2+ levels activate protein arginase deiminase 4 (PAD4) [11], which promotes chromatin decondensation by converting arginine residues of core histones H3 and H4 into citrulline [8]. In addition, ROS also lead to the gradual disassembly of the nuclear membrane, followed by the dispersion of chromatin throughout the cytoplasm, where it is decorated with granular and cytoplasmic contents [12]. Ultimately, chromatin, DNA, granular, and cytoplasmic contents are released into the extracellular space as NETs [6,13].

Neutrophil functions, specifically the extrusion of NETs, are considered beneficial during infection [14]. However, dysregulated or extensive NET formation may result in undesirable tissue damage [15,16] and is linked to many inflammatory disorders, including sepsis, asthma, lupus, rheumatologic diseases, as well as diabetes [17]. Additionally, neutrophils receive increasing interest in cancer research as potential drivers of metastasis [18]. Furthermore, NETs are discussed as potential inducers of endothelial tissue damage, leading to various forms of vasculitis [19]. The accumulation of neutrophils, as well as the entailing activation and NET formation, at the culprit site lesion during acute coronary syndrome or acute myocardial infarction, is associated with poor disease prognosis and an increased long-term mortality rate [20–22]. In addition to systemic disorders, excessive NET formation is also associated with locally impaired or prolonged tissue regeneration, due to increased neutrophil-derived ROS in the microenvironment of the injury [16,23–26].

Recent advances in cell-derived, yet cell-free medicinal products have increasingly gained attention in regenerative medicine [27,28]. Although initial research on cell-free therapeutic agents focused on secretomes derived from stem cells, we could demonstrate that the secretome of peripheral blood mononuclear cells (PBMC) exhibits comparable regenerative effects [29-35]. The potency of the PBMC-derived secretome (PBMCsec) was further increased by exposing PBMCs to 60 Gy γ -irradiation, which induces apoptosis and necroptosis, resulting in the release of a plethora of pro-regenerative paracrine factors [29]. Lichtenauer et al. showed strong regenerative potential of PBMCsec in rodent and porcine models of acute myocardial infarction [35]. These pioneering findings laid the foundation for further studies, which identified a broad spectrum of therapeutic implications for PBMCsec in a vast variety of pathologic conditions, including chronic heart failure after myocardial infarction [36], cerebral ischemia [35], burn injury [35], diabetic wound healing [30], and acute spinal cord injury [35]. Furthermore, strong anti-inflammatory properties of PBMCsec have been demonstrated in the context of myocarditis [35], as well as inflammatory skin conditions [34]. The observed tissue-regenerative effect of PBMCsec is based on a complex interplay of various biologically active agents produced and released by stressed PBMCs [30,32,35]. The broad action spectrum of PBMCsec has been intensively investigated and revealed promising treatment opportunities, where anti-inflammatory [37,38], anti-microbial [33], tissue-regenerative [32], pro-angiogenic [29,30], and vasodilatory [35] properties are important.

Although PBMCsec possesses compelling immunomodulatory effects [35], potential anti-inflammatory and stabilizing effects on (activated) neutrophils have not been investigated so far. Hence, we sought to investigate the effect of PBMCsec on NET formation.

2. Materials and Methods

2.1. Ethics Statement

This study was conducted in accordance with the Declaration of Helsinki and applicable local regulations. Use of human neutrophils was approved by the institutional ethical review board of the Medical University of Vienna (Vienna, Austria) (protocol code 1539/2017). Written informed consent was obtained from all donors.

2.2. Generation of PBMCsec

PBMCsec was produced in compliance with good manufacturing practice by the Austrian Red Cross, Blood Transfusion Service for Upper Austria (Linz, Austria) as previously described [30,34]. Briefly, the PBMCs were enriched using Ficoll-Paque PLUS (GE Healthcare, Chicago IL, USA) density gradient centrifugation. Cell suspensions were adjusted to 2.5×10^7 cells/mL and exposed to 60 Gy γ -irradiation (IBL 437, Isotopen Diagnostik CIS GmbH, Dreieich, Germany). Subsequently, cells were cultured in phenol red-free CellGenix GMP DC medium (CellGenix, Freiburg, Germany) for 24 h. Cells, as well as cellular debris, were removed by centrifugation. The conditioned supernatants containing the secretome were filtered through 0.22 µm filters followed by viral clearance using Theraflex methylene blue technology (MacoPharma, Mouvaux, France). The secretomes were lyophilized and sterilized by high-dose γ -irradiation (25,000 Gy, Gammatron 1500, Mediscan, Seibersdorf, Austria). CellGenix GMP DC medium without cells was used as vehicle control. The GMP batches A000918399131, A00918399136, and A000918399132 were used in this study. The stock concentration of one vial lyophilized secretome equals to 25 units/mL.

2.3. Fractionating PBMCsec

The lipid fraction was purified according to Folch et al. (PMID: 13428781) with minor modifications. Briefly, one part reconstituted PBMCsec was mixed with 9 parts 2:1 (vol/vol) chloroform-methanol and, subsequently, excessively vortexed. Then, 0.7 M of formic acid was used to acidify the emulsion (a one-fourth volume of the chloroform-methanol mix) and homogenized by thorough shaking. Phase separation was obtained by leaving the samples on ice for 30 min. The lower, organic phase was further applied to rotary vacuum evaporation (475 mbar, 100 rpm, 60 °C water bath temperature) in order to eliminate solvents. The protein fraction was isolated by combining four times the volume of ice-cold acetone (VWR Chemicals, PA, USA) to one volume of reconstituted PBMCsec, followed by thorough vortexing and incubation at -20 °C for 60 min, to obtain a protein precipitate. After centrifuging the sample at $18,000 \times g$ for 10 min, ice-cold acetone was added to the protein pellet, briefly vortexed and centrifuged at $18,000 \times g$ for 10 min. Acetone was discarded and remaining acetone was allowed to evaporate at room temperature. Finally, the protein pellet was resuspended in 0.9% NaCl in the initially used volume. DNA was isolated by adding equal amounts of isopropanol (Merck Millipore, MA, USA) as PBMCsec and 1/10 volume of 7.5 M sodium acetate (Merck Millipore) and incubated at -20 °C for 1 h. After centrifugation for 5 min at $18000 \times g$ the DNA pellet was washed twice with 1 mL 70% Ethanol (Merck Millipore) followed centrifugation at 18,000× g for 5 min. The DNA pellet was allowed to dry for 10 min at room temperature prior to resuspension in double distilled, nuclease-free H2O. Extracellular vesicles were obtained by ultracentrifugation at $110,000 \times g$ for 2 h at 4 °C, as previously described [30]. To ensure comparability, all fractions were used in the same concentrations as are present in PBMCsec. All fractions were tested separately and in a combined form. To reconstitute the fractions, equal volumes of each fraction were combined and further diluted to the equivalent concentration of PBMCsec.

2.4. Neutrophil Isolation

Neutrophils were isolated using the MACSxpress Whole Blood Neutrophil Isolation kit (Miltenyi Biotec, Bergisch-Gladbach, Germany) according to manufacturer's instructions. In brief, magnetic beads were resuspended in 2 mL of buffer A. One-fourth of the total amount of processed blood of magnetic beads and buffer B were added to the blood

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and incubated at room temperature for 5 min under constant gentle rotation. Blood and isolation cocktail mix were placed in the MACSxpress Separator (Miltenyi Biotec) and allowed to separate for 15 min. Clear, neutrophil-containing top phase was transferred into a fresh tube and washed with basal RPMI 1640 without phenol red (Thermo Fisher Scientific, Waltham, USA). If required, a red blood cell lysis was performed using a Red Blood Cell Lysis Buffer (Abcam, Cambridge, UK) for 10 to 15 min at room temperature. Neutrophils were resuspended in basal RPMI 1640 in an assay dependent concentration without phenol red for further use.

2.5. Induction of NET Formation

Either isolated neutrophils or whole blood samples after red blood cell lysis were pre-treated with 2 units/mL PBMCsec or equivalent vehicle medium for 20 min at 37 °C. Cells were then stimulated with 5 μ M ionomycin (Sigma Aldrich, St. Louis, MO, USA) or 100 nm phorbol 12-myristate 13-acetate (PMA) (Sigma Aldrich), or 10 μ M Thapsigargin (Abcam) for 2 h at 37 °C unless indicated otherwise.

2.6. Flow Cytometry

After stimulation with indicated compounds, cells were centrifuged and stained with anti-citrullinated Histone H3 antibody (ab5103, Abcam) to detect NETs, anti-CD66b antibody (pacific-blue conjugated mouse anti-human, clone G10F5, BioLegend, San Diego, CA, USA) and anti-CD15 antibody (4hycoerythrin-cyanine 7 conjugated mouse anti-human, clone W6D3, BioLegend) to identify neutrophils. Flow cytometric analysis was performed using BD FACSCanto II and BD FACSDiva software (version 6.1.3) (BD Pharmingen, San Jose, CA, USA).

2.7. Cell Viability Assay

Incucyte Cytotox Dye for Counting Dead Cells (Sartorius, Goettingen, Germany) was used according to the manufacturer's instructions. In brief, cells were treated as indicated, followed by the addition of 250 nm cytotox green dye for staining 100 μ L cell suspension in a concentration of 4 \times 10⁶ cells/mL condition in a 96-well plate. Cell death was assessed over the indicated time periods in a microplate reader at an excitation wavelength of 485 nm and an emission wavelength of 520 nm. Microplate reader (BMG Labtech, FLUOstar OPTIMA) and the BMG Labtech Optima software (software version 2.20Rs, BMG Labtech, Ortenberg, Germany).

2.8. EZ4U Cell Proliferation and Cytotoxicity Assay

EZ4U (Biomedica, Vienna, Austria) assay was performed according to manufacturer's instructions. Briefly, the substrate was dissolved in 2.5 mL activator solution and prewarmed to 37 °C. Then, a 20 μ L substrate was added to 200 μ L cell suspension at a concentration of 4 \times 10⁶ cells per condition in a 96-well plate and incubated for 2 h. Continuous absorbance measurements at 450 nm were performed using a microplate reader (BMG Labtech, FLUOstar OPTIMA) and the BMG Labtech Optima software (software version 2.20Rs).

2.9. ROS Production Measurement

ROS production was measured using the DCFD/H2DCFDA cellular ROS assay kit (Abcam). Cells were treated as indicated and the assay was performed as recommended by the manufacturer.

2.10. Ca2+ Flux Measurement

Ratiometric calcium flux measurements with Fura Red were performed as described by Wendt et al., with minor modifications [39]. In brief, a cell suspension of 4×10^6 cells per condition, either isolated neutrophils or whole blood, pre-treated with PBMCsec or vehicle for 20 min as indicated, were washed, resuspended in 400 μ L full medium containing 1 μ M

Fura Red (Invitrogen, Thermo Fisher Scientific, Waltham, MA, USA), and incubated for 30 min at 37 °C. Cells were washed once with medium, resuspended in 4 mL medium and incubated for another 30 min at 37 °C. Subsequently, cells were rested on ice for up to 30 min. Data were acquired on a FACSAria III flow cytometer (BD Bioscience, San Jose, CA, USA). Before intracellular calcium flux measurement, 1 mL of Fura Red-loaded cells was transferred to a FACS tube and pre-warmed for 5 min at 37 °C in a water bath. The cells were kept at 37 °C during the whole measurement. The baseline response was recorded for 30 s prior to stimulation with 5 μ M ionomycin. Changes in calcium mobilization were recorded for a total of 120 s. Fura Red was excited using a 405 nm violet laser and a 561 nm green laser and changes in emission were detected with a 635 LP, 660/20 BP, and a 655 LP, 795/40 BP filter set, respectively. The 'Fura Red Ratio' over time was calculated using the Kinetics tool in FlowJo software (version 9.3.3, Tree Star Inc., Ashland, OR, USA) as follows:

 $Fura \ Red \ Ratio = \frac{increase \ of \ 405 \ nm \ induced \ emission}{decrease \ of \ 561 \ nm \ induced \ emission}$

2.11. DNase Activity Measurement

DNase activity was measured by incubating 0.25 μ g/ μ L Lambda DNA with 0.5 M acetate/NaOH at pH 4.8 (Merck Millipore), 50 mM CaCl₂(Merck Millipore), 50 mM MgCl₂ (Sigma-Aldrich, St. Louis, MO, USA), 40 mM 2-mercaptoethanol (Merck Millipore) and either 2 units/mL PBMCsec or equivalent vehicle control or DNase I (Thermo Fisher Scientific) as positive control for 1 h at 37 °C. 10 μ L of each sample were loaded into a 1% agarose gel with gel red (Biotium, Fremont, CA, USA) together with 2 μ L 6x loading dye (Thermo Fisher Scientific). Electrophoresis was performed at 200 V for 35 min. DNase activity was determined by absence of λ -DNA.

2.12. Proteome Profiler

The Human Apoptosis Array kit (R&D Systems, Minneapolis, MN, USA) was used in accordance with the manufacturer's instructions with no modifications. Isolated neutrophils were treated as indicated and cell lysates of 4×10^6 cells per condition of 4 individual donors were pooled.

2.13. PAD4 Inhibitor Assay

The inhibitory capacity of PBMCsec was measured using the PAD4 inhibitor Screening Assay kit (ammonia, Cayman Chemical, Ann Arbor, MI, USA) and performed according to manufacturer's instructions.

2.14. Western Blot Analysis

For Western blot analysis, cells were lysed in 1x Laemmli sample Buffer (Bio-Rad Laboratories, CA, USA) supplemented with Protease Inhibitor Cocktail (Sigma-Aldrich), After sonication SDS-PAGE electrophorese was performed on 4–20% gradient gels (Criterion TGX Precast Gels, Bio-Rad Laboratories). Proteins were electrotransferred onto 0.2 μ M nitrocellulose membranes (Trans-Blot Turbo, Bio-Rad Laboratories) and immunodetected using primary antibodies against pan phospho-PKC (β II Ser660) antibody (#9371, Cell Signaling Technology, Danvers, MA, USA). This antibody detects endogenous PKC α , β I, β II, δ , ε , η and θ isoforms phosphorylated at carboxy-terminal residue homologous to serine 660 of PKC β II. Peroxidase-conjugated secondary antibody were detected with the SuperSignal West Dura Extended Duration Substrate (Thermo Scientific, Rockford, MA, USA), according to manufacturer's instructions. A Ponceau S solution (Sigma-Aldrich) staining served as equal loading control.

2.15. Statistics

Statistical analyses were performed using Prism 8.0.1 (Graph Pad Prism). Data were shown \pm standard deviation (SD). One-way ANOVA and Sidak's multiple comparisons test were performed and * p < 0.0332, ** p < 0.0021, *** p < 0.0002, **** p < 0.0001.

3. Results

3.1. PBMCsec Inhibits NET Formation

Although immunomodulatory properties of PBMCsec have been well described in a variety of different cell types [30,34,40], a potential effect of PBMCsec on neutrophils has not yet been explored. To investigate whether PBMCsec interferes with experimentally induced NET formation, we pre-incubated human whole blood with 2 units/mL PBMCsec or an equivalent dose of vehicle prior to stimulation with 5 µM ionomycin. Unstimulated or PBMCsec-treated samples showed few citH3-positive neutrophils, identified by CD15⁺CD66b⁺citH3⁺ staining (Figure 1A, upper panel and Figure 1B, $2.4 \pm 0.97\%$ and $1.6 \pm 1.22\%$ positive cells, respectively, Figure S1A). The addition of ionomycin strongly induced NET formation, as demonstrated by a significant increase in citH3-positive cells $(54.52 \pm 14.41\%)$ after two hours of incubation (Figure 1A, bottom panel and Figure 1B). This effect was almost completely abolished by pre-incubation with PBMCsec before ionomycin treatment ($3.9 \pm 1.28\%$ citH3-positive neutrophils). A dose titration of PBMCsec revealed that 2 U/mL was the lowest dose with NET inhibiting activity (Figure S1C). By contrast, vehicle treatment showed only a weak reduction in NET-formation (27.94 \pm 17.71% citH3-positive neutrophils) (Figure 1A, bottom panel, and Figure 1B). As DNA is one of the major constituents of NETs [4], we analysed the amount of extracellular DNA in ionomycin-stimulated samples using cytotox green staining (Figure 1C). After two hours, only a weak cytotox green signal was detected in untreated, PBMCsec, or vehicle treated samples. Although stimulation with ionomycin resulted in a drastic increase in extracellular DNA (Figure 1C), the addition of PBMCsec almost completely inhibited the release of DNA after ionomycin stimulation. To exclude the possibility that the observed effect was due to a direct inhibitory effect of PBMCsec on ionomycin, we also used PMA (100 nM), another well-described inducer of NET formation. As shown in Figure 1D, PMA treatment of PBMCsec pre-incubated neutrophils led to a comparable inhibition of DNA release. Additionally, we evaluated the metabolic activity of ionomycin-activated neutrophils [41]. Ionomycinactivation resulted in a prominent decrease in metabolic activity which, in contrast to vehicle treatment, was completely abolished by the addition of PBMCsec (Figure S1B). Immunostaining of neutrophils for citH3 showed classical NET-structures after ionomycin treatment, which were completely absent in the presence of PBMCsec (Figure 1E). Taken together, these findings indicate that treatment of experimentally activated neutrophils with PBMCsec significantly reduces the formation of NETs.

3.2. A Synergistic Effect of Different Substance Classes Inhibits NET Formation

PBMCsec is composed of different substance classes, including free DNA, lipids, proteins, and extracellular vesicles [34,35,38,42] (Figure 2A). Thus, we further aimed to investigate to what extent the individual fractions contribute to the inhibitory effects on NET formation (Figure 2B,C). Therefore, PBMCsec and its fractions were added to whole blood prior to ionomycin stimulation and citH3 levels were analysed (Figures 2B,C and S2). Ionomycin treatment showed a significant increase in citH3⁺ neutrophils, which was almost completely abolished by the addition of PBMCsec. In contrast, purified fractions showed only partial inhibition of ionomycin-induced histone citrullination, indicating that the inhibitory effect of PBMCsec requires the complex interplay of all fractions of the secretome. Stimulation with the reconstituted fractions of PBMCsec fully restores the inhibitory activity of NET formation (Figure 2B,C).

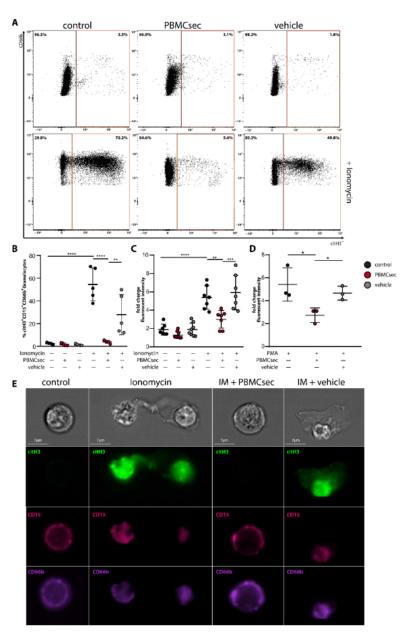


Figure 1. PBMCsec inhibits NET formation. Erythrocyte-lysed blood was treated with PBMCsec or vehicle for 20 min and subsequently stimulated with ionomycin (IM) for 2 h and analysed with flow cytometry, cytotox staining, and image stream analysis. (A) Neutrophils were identified in flow cytometry as CD66b⁺CD15⁺ cells and NET-forming neutrophils were characterized by citH3. Control, PBMCsec, or vehicle treated samples are shown in the top panel and ionomycin-activated neutrophils are shown in the bottom panel. n = 5. One representative sample is shown out of five

replicates summarized in (**B**). (**C**) Extracellular DNA content was measured using cytotox staining of neutrophils after pre-treatment with PBMCsec or vehicle and subsequent activation for 2 h with (**C**) IM or (**D**) PMA. Fold change increase in relative fluorescent intensity is shown after two hours of stimulation relative to time point zero/start of stimulation/minute one after induction of NETs. (**E**) Visualization of IM-activated neutrophils was performed using image stream analyses. Untreated (control) and IM-PBMCsec treated neutrophils did not show citH3⁺ staining. IM and IM-vehicle treated neutrophils showed robust citH3⁺ staining of cells and additional extracellular structures (indicative for NETs). Green, citH3; magenta, CD15; purple, CD66b; *n* = 2. One representative sample is shown. Data are represented as individual values with mean and error bars indicate SD, one-way ANOVA and Sidak's multiple comparisons test. * *p* < 0.0032, ** *p* < 0.0021, *** *p* < 0.0002, **** *p* < 0.0001.

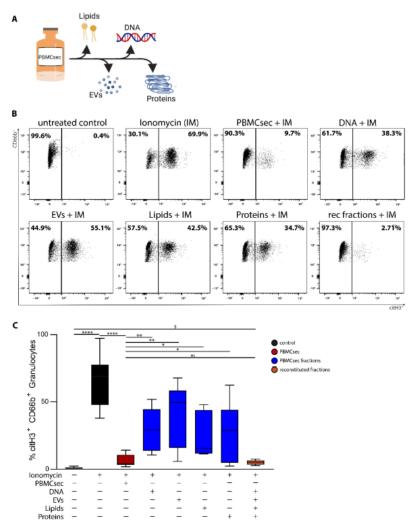


Figure 2. Isolated substance classes of PBMCsec show a synergistic effect on NET-inhibition. (A) Schematic depiction of the isolated and tested substance classes of PBMCsec. This scheme was

created with BioRender.com, accessed on 2 June 2022 (**B**) Neutrophils were identified in flow cytometry as CD66b⁺CD15⁺ cells and NET formation was characterized by citH3+ signal. rec fractions + IM = reconstituted fractions + ionomycin. n = 6. One representative sample out of six is shown, summarized in (**C**) one-way ANOVA and Sidak's multiple comparisons test. Data are represented as mean and error bars indicate SD. § = ANOVA without multiple comparison tests, p = < 0.0001; * p < 0.032, ** p < 0.0021, **** p < 0.0001.

3.3. PBMCsec Does Not Show DNase-Activity

As digestion of NETs by DNAses is the main NETs-clearing mechanism [43], we next investigated whether PBMCsec displays DNAse activity. Therefore, we incubated λ DNA with PBMCsec and analysed DNA degradation (Figure 3A). Compared to recombinant DNAse I, which completely digested λ DNA, PBMCsec showed no DNA degrading activity (Figure 3A). Since this in vitro assay was optimized for DNAse I only, we further tested potential NETs-degrading properties of PBMCsec in whole blood ex vivo. For this purpose, we stimulated whole blood with ionomycin and applied PBMCsec either prior to or two hours after ionomycin treatment (Figure 3B). In contrast to neutrophils treated with PBMCsec prior to their activation, treatment two hours after induction of NET formation did not reduce the amount of citH3 positive neutrophils (Figure 3C,D and Figure S3). These data demonstrate that PBMCsec does not degrade pre-formed NETs by DNases, suggesting an active intervention in the NET-forming signalling cascade.

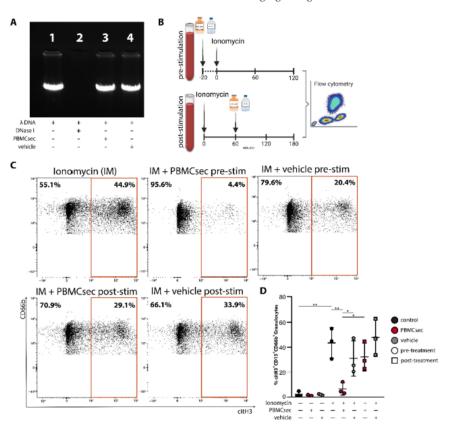


Figure 3. PBMCsec inhibits NET formation by a DNase-independent mode of action. (A) DNase

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activity was measured in a cell-free assay by co-incubation of PBMCsec or vehicle with λ -DNA. DNase I was used as positive control. n = 3, one representative sample is shown. (**B**) Schematic depiction of the adapted neutrophil stimulation protocol for the measurement of potential DNase activity in a cell-based assay. This scheme was created with BioRender.com, accessed on 2 June 2022 (**C**) Neutrophils were identified as CD66b⁺CD15⁺ cells and citH3+ signal was used to characterize NET formation. n = 3, one representative experiment is shown. (**D**) Statistical summary of all biological donors is shown. Data are represented as individual values with mean and error bars indicate SD. One-way ANOVA and Sidal's multiple comparisons test. * p < 0.0332, ** p < 0.0021.

3.4. PBMCsec Inhibits NET Formation by Preventing ROS Production and PAD4 Activity

Induction of NETs requires an increase in intracellular calcium levels [8,9,11]. We, therefore, first investigated whether PBMCsec interferes with ionomycin-induced calcium influx. Analysis of intracellular calcium signalling, using a Fura Red based flow cytometry approach, revealed that pre-treatment of whole blood with PBMCsec only marginally reduced calcium influx after addition of ionomycin (Figure 4A,B). The decline in calcium flux was only transient and returned to control values rapidly. No significant difference was observed between PBMCsec and vehicle treatment, suggesting that the observed decrease in the calcium influx is not sufficient to affect NET formation. In addition, we also tested Thapsigargin, an irreversible inhibitor of the sarco-endoplasmic reticulum Ca²⁺-ATPase (SERCA) pumps, which prevents the storage of excess intracellular calcium in the endoplasmic reticulum [44]. The addition of 10 µM Thapsigargin induced NET formation comparably to ionomycin (Figure S4). Furthermore, PBMCsec was able to counteract Thapsigargin-induced NET formation (Figure S4) very efficiently. Together, these findings indicate that PBMCsec does not interfere with calcium flux in neutrophils and suggest that its NET-inhibiting action is mediated downstream of calcium flux. As increased intracellular calcium concentrations promote the activity of PKC [9], we next investigated PKC phosphorylation by Western blot analysis, using a pan phosphor-PKC antibody. As this antibody detects several phosphorylated isoforms of PKC, an assignment to a specific isoform was not possible. However, we were not able to detect differences in the amount of phosphorylated PKC after pre-treating neutrophils with PBMCsec or vehicle (data not shown), suggesting that the inhibitory action of PBMCsec is also downstream of PKC.

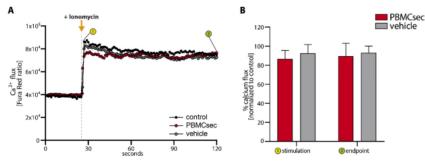


Figure 4. PBMCsec inhibits NET formation without interfering with calcium flux. (A) Ratiometric calcium flux was measured with Fura Red in neutrophils. Neutrophils were observed for approximately 30 s to record a baseline Fura Red ratio indicating homeostatic calcium flux prior to the addition of IM and subsequent analysis for a total of 120 s. (B) Statistical analysis of the percent reduction in calcium flux in PBMCsec or vehicle treated neutrophils compared to control samples is shown for the time points of stimulation (①, addition of IM) and the endpoint (②,120 s) n = 4. Error bars indicate SD. No statistically significant reduction was observed for PBMCsec or vehicle treatment.

Since activation of the NET signalling pathway down-stream of NADPH leads to the production of ROS and activation of PAD4 [8,10,11], we next investigated whether PBMCsec exerts its inhibitory activity by modulating these processes. We, therefore, investigated ionomycin-induced production of ROS using the cell permeant reagent 2',7'-dichlorofluorescin diacetate (DCFDA). Ionomycin treatment of PBMCsec-stimulated neutrophils resulted in a significant decrease in ROS production, as compared to ionomycin treatment alone (Figure 5A). By contrast, pre-treatment with vehicle showed no inhibitory effect (Figure 5A). Analysis of protein expression revealed that PBMCsec inhibited ionomycin-induced down-regulation of known anti-oxidative factors, including hemoxygenase-1 (HO-1) [45] and hypoxia inducible factor 1 alpha (HIF-1 α) [46] in purified human neutrophils (Figures 5B and S5B), which was not observed in vehicle-treated cells.

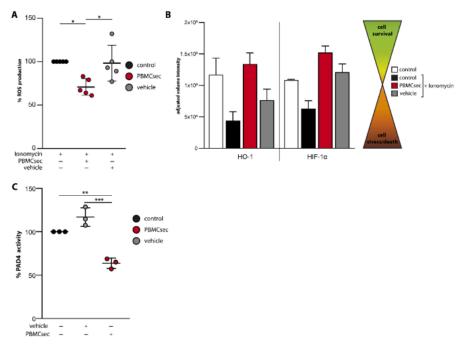


Figure 5. PBMCsec prevents ROS production and PAD4 activity. (**A**) ROS production in IM stimulated and PBMCsec or vehicle treated neutrophils normalized to IM-induced ROS production. n = 5(**B**) Analysis of protein levels of HO-1 and HIF-1 α of isolated neutrophils stimulated with ionomycin and treated with PBMCsec or vehicle, using a proteome profiler is shown. Cell lysates of four individual donors and experiments were pooled. Error bars indicate SD of two technical replicates. (**C**) Enzymatic activity of PAD4 was measured in a cell-free assay upon co-incubation of PBMCsec or vehicle with the PAD4-substrate and normalized to untreated control. n = 3, data are represented as individual values with mean and error bars indicate SD, unless indicated otherwise. One-way ANOVA and Sidak's multiple comparisons test was performed where significances are indicated. * p < 0.0332, ** p < 0.0021, *** p < 0.0002.

The activation of PAD4 with subsequent histone-citrullination represents one of the final steps in the NETs signalling pathway [11]. Thus, we next investigated PAD4 activity in a cell-free assay. Although vehicle treatment did not show PAD4 inhibiting properties, PBMCsec inhibited PAD4 activity by approximately 40% (Figure 5C). Together, these data demonstrate that PBMCsec exerts its inhibitory activity on NET formation by reducing intracellular ROS production and preventing PAD4 activation.

4. Discussion

The formation of NETs is a highly effective first line defence mechanism against invading pathogens [17]. However, there is growing evidence that excessive NETs formation contributes to tissue damage and the induction of auto-immune diseases [16,47–49]. Although several substances, including acetylsalicylic acid, cyclosporine A [50], metformin [51] and chloroquine [52], have been shown to influence NET formation, therapeutic drugs targeting NET formation are so far not available. In the current study, we provide evidence that PBM-Csec effectively inhibited NET formation by reducing ROS production and PAD4 activation, thereby providing a novel potential cell-derived but cell-free therapeutic intervention for NET-associated diseases.

The tissue regenerative and anti-inflammatory action spectrum of PBMCsec is multifaceted [29,32,34,37,38,53,54], and most of its beneficial effects have been shown to require the interplay of several components of the secretome [30,55]. Indeed, we also found that NET formation was only fully inhibited when neutrophils were treated with the whole secretome or reconstituted purified fractions. Since we observed NETs inhibition at different steps of the NETs signalling pathway, we hypothesize that individual secretome fractions act on different signalling molecules. Recently, Laggner et al. showed that lipids present in PBMCsec attenuate skin inflammation and allergic reactions by targeting dendritic cell function [34], as well as mast cell and basophil activation, respectively [56], suggesting that lipids are mainly responsible for the anti-inflammatory activities of PBM-Csec. Several lipid species have been detected in PBMCsec, including phosphatidylserines, lysophosphatidylcholines, lysophosphatidylethanolamines, phosphatidylcholines, phosphatdiylethanolamines, and resolvins [34]. Interestingly, several studies described a NET-inhibitory or NET-resolving action of resolvins [57-59]. Spinosa et al. demonstrated a decreased NET burden accompanied by reduced abdominal aortic aneurysm in reslovintreated mice [57]. In addition, neutrophils derived from resolvin-treated mice showed less susceptibility to ionomycin-induced NET formation [58]. Although both of these studies identified less NET formation in the presence of resolvins, Chiang et al. showed that NETs formed after Staphylococcus aureus infection were more efficiently cleared by macrophages after treatment with reslovins [59]. These data indicate an important function of resolvins in the prevention of NET formation and/or resolution of NETs. Therefore, it is conceivable that the partial inhibition of NET formation observed by PBMCsec-lipids may be explained by the variety of resolvins found in PBMCsec [34]. However, whether PBMCsec-derived resolvins or other lipid classes are indeed involved in PBMCsec-induced NETs inhibition needs to be determined in future studies.

In addition to the lipid fraction, PBMCsec-derived proteins also showed a strong inhibitory action on NET formation. Previous studies demonstrated that addition of either bovine or human serum albumin to ionomycin-treated cells almost completely blocked NET formation by chelating calcium [60]. However, as we only detected a slight decrease in calcium influx after treatment with PBMCsec and vehicle, a sole albumin-dependent effect is unlikely. Additionally, PBMCsec treatment also abrogated Thapsigargin-induced NETosis, which indicates that PBMCsec-mediated NETosis inhibition occurs without interfering with calcium flux or the cells' capability to store excess calcium into intracellular storage units, such as the endoplasmic reticulum. Furthermore, we also showed comparable effects when NET formation was induced with PMA, which induces NETs in a calcium-independent manner. Together, our data suggest a calcium- and albumin-independent mode of action of the protein fraction of PBMCsec. Further in-depth proteomics analyses of PBMCsec-derived proteins are required to elucidate whether a single protein or a combination of proteins is responsible for the inhibition of NET formation.

Our data suggest the inhibition of ROS production and PAD4 activation as the two major modes of action for the reduction in NET formation by PBMCsec. Oxidative stress, especially the generation of ROS, is a hallmark of NET formation [10,61] and HSPs are known to effectively block excessive ROS production [62]. Our study revealed that PBM-Csec inhibited hemeoxygenase 1 (HO-1 or HSP32) and HIF-1 α downregulation during

ionomycin-induced NET formation. Both HO-1 and HIF-1 α have been shown to promote neutrophil survival by reducing ROS levels [63] and via Akt and NF κ B signalling under stress, respectively [46,64]. These data suggest that PBMCsec contributes to the stabilization of the delicate balance of pro- and anti-oxidative processes by regulating the expression of HSPs, thereby preventing neutrophil-induced tissue damage. Since HO-1 is also known to downregulate adhesion molecules and chemokines required for neutrophil infiltration [45], PBMCsec may alleviate inflammatory responses by reducing neutrophil infiltration in damaged and inflamed tissue. However, further studies are required to unravel the exact mechanism by which PBMCsec counteracts ROS production as it is not yet clear whether it functions as ROS-scavenger, inhibits the liberation of ROS from mitochondria or if it interferes with the functional assembly of NADPH oxidase subunits.

PAD4 is one of the most prominently investigated factors critical for NET formation, and PAD inhibitors have been extensively studied in the context of a broad variety of diseases, including multiple sclerosis [65], myocardial infarction [66], and rheumatoid arthritis [67]. However, the exact mechanism of PAD4 inhibition is not yet fully understood [67]. Our data indicate that stressed PBMCs secrete factors that serve as PAD4 inhibitors. Interestingly, Yost et al. identified a group of peptides in umbilical cord blood with strong PAD4-inhibiting effects, leading to inhibition of NETs [68]. Sequence analyses identified α 1-antitrypsin, a serine protease inhibitor, known to possess immunomodulatory and anti-inflammatory properties [69], as the main PAD4-inhibiting factor. Interestingly, α 1-antitrypsin is synthesized by circulating monocytes and, therefore, a component of PBMCsec (Figure S6) [70]. According to our quantification analysis, 25–30 ng/mL SER-PINA1 are present in two units of PBMCsec. This enzyme inhibitor has been considered as an acute phase protein, which contributes to the inhibition of NET formation by targeting a vast array of factors contributing to NET formation [71]. Further studies are needed to identify the PAD4-inhibiting factor(s) in PBMCsec.

5. Conclusions

In summary, we have demonstrated a strong NETs-inhibitory activity of PBMCsec via a dual mechanism. Specifically the identification of a PAD4 inhibitor, produced naturally in the human body, as well as the prevention of ROS production might strongly improve the treatment of diseases associated with excessive NET formation, such as rheumatoid arthritis [67], multiple sclerosis [65], sepsis [17], heart failure, and myocardial infarction [66]. Pre-clinical toxicological evaluation of PBMCsec has already been performed without the occurrence of major adverse events after topical and intravenous application (LPT, study number 35015). Therefore, our study has paved the way for a clinical study in humans, assessing the potency of PBMCsec in NETs-associated diseases in vivo.

Supplementary Materials: The following supporting information can be downloaded at: https:// www.mdpi.com/article/10.3390/antiox11081559/s1, Graphical abstract: PBMCsec inhibits NET formation; Figure S1: Gating strategy, PBMCsec improves neutrophil metabolic activity; Figure S2: Flow cytometric analysis of spontaneous NET formation; Figure S3: Flow cytometric analysis of unstimulated neutrophils; Figure S4: PBMCsec inhibits Thapsigargin-induced NETosis; Figure S5: Gating strategy and purity of isolated neutrophils; Figure S6: SERPINA1 abundance in PBMCsec.

Author Contributions: Conceptualization, K.K., M.M., and H.J.A.; methodology, K.K., M.M., A.S.O., and T.M.H.; validation K.K., M.M., A.S.O., T.M.H.; formal analysis, K.K.; investigation, K.K., A.S.O., T.M.H., and K.P.; resources, H.J.A.; data curation, K.K., D.C., M.D., and D.B.; writing—original draft preparation, K.K. and M.M.; writing—review and editing, K.K., M.M., M.L., and A.M.; visualization, K.K.; supervision, M.M.; project administration, K.K. and M.M.; funding acquisition, H.J.A. All authors have read and agreed to the published version of the manuscript.

Funding: This research project was financed in part by the Austrian Research Promotion Agency grant "APOSEC" (852748 and 862068, 2015–2019), by the Vienna Business Agency "APOSEC to clinic," (2343727, 2018–2020), and by the Aposcience AG under group leader H.J.A., M.M. was funded by Aposcience AG.

Institutional Review Board Statement: The study was conducted in accordance with the Declaration of Helsinki, and approved by the Institutional Ethical Review Board of the Medical University of Vienna (protocol code 1539/2017).

Informed Consent Statement: Informed consent was obtained from all subjects involved in the study.

Data Availability Statement: Data is contained within the article or Supplementary Materials.

Acknowledgments: We thank H.P. Haselsteiner and the CRISCAR Familienstiftung for their belief in this private-public partnership to augment basic and translational clinical research.

Conflicts of Interest: The Medical University of Vienna has claimed financial interest. H.J.A. holds patents related to this work (WO 2010/079086 A1; WO 2010/070105 A1).

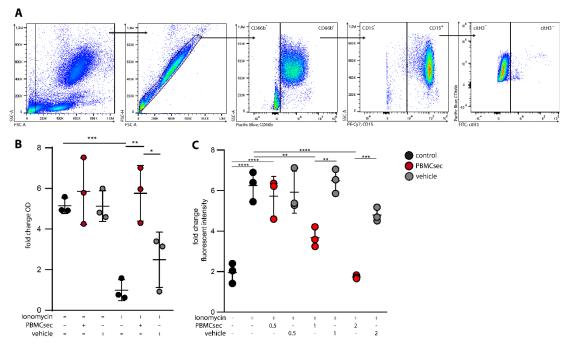
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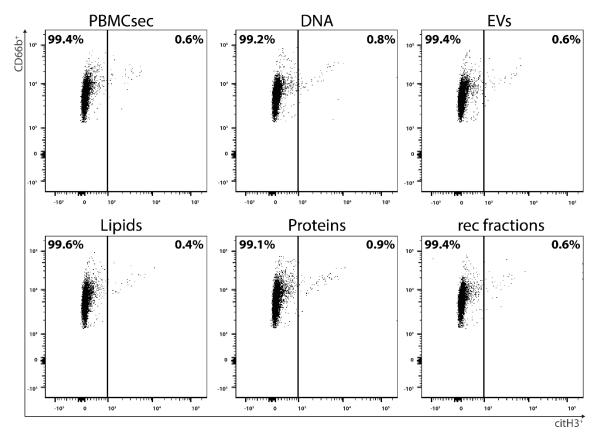
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1.2 Supplementary Figures

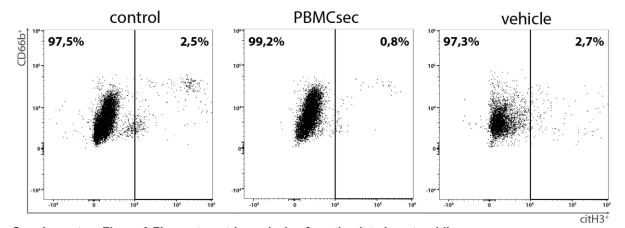
Supplementary Figure 1 PBMCsec improves neutrophil metabolic activity and inhibits NETosis in a dosedependent manner

A) Flow cytometry gating strategy for erythrocyte lysed blood samples is shown. (B) Metabolic activity of neutrophils was measured using an absorbance based assay (EZ4U). Vehicle treated neutrophils did not show altered metabolic activity compared to untreated control samples. PBMCsec treatment appeared to partially promote metabolic activity of non-activated neutrophils. IM treatment resulted in a significant reduction of metabolic activity of neutrophils which was abolished by PBMCsec treatment. Vehicle treatment could not restore homeostatic metabolic activity in IM-activated neutrophils. C) Extracellular DNA content was measured using cytotox staining of neutrophils after pre-treatment with PBMCsec or vehicle in a dose dependent manner. Data are represented as mean ± SD, one-way ANOVA and Sidak's multiple comparisons test. *p<0.0322, **p<0.0021, ***p<0.0002

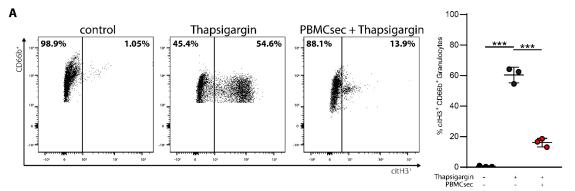


Supplementary Figure 2 Flow cytometric analysis of spontaneous NET formation

Flow cytometric analysis of untreated control neutrophils and neutrophils treated with PBMCsec derived substance classes in absence of an activating stimulus is shown. Neutrophils were identified as CD66b+CD15+ cells and NET formation was characterized by additional citH3+ signal. n = 3, one representative experiment is shown.

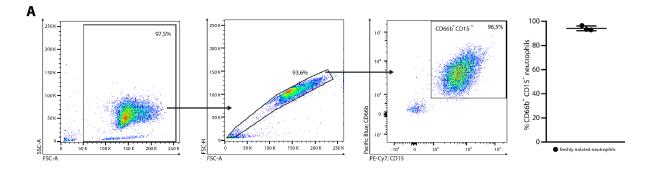


Supplementary Figure 3 Flow cytometric analysis of unstimulated neutrophils Flow cytometric analysis of untreated control neutrophils and neutrophils treated with PBMCsec or vehicle in absence of an activating stimulus after two hours is shown. Neutrophils were identified as CD66b+CD15+ cells and NET formation was characterized by additional citH3+ signal. n = 3, one representative experiment is shown.



Supplementary Figure 4 PBMCsec inhibits Thapsigargin-induced NETosis

Neutrophils were analysed using flow cytometry and identified by CD66b⁺CD15⁺ signal. NETosis was identified by citH3⁺ positive signal. Untreated control samples were compared to Thapsigargin-activated cells as well as Thapsigargin-activated and PBMCsec-treated cells. Data are represented as mean ± SD, one-way ANOVA and Sidak's multiple comparisons test. ***p<0.0002



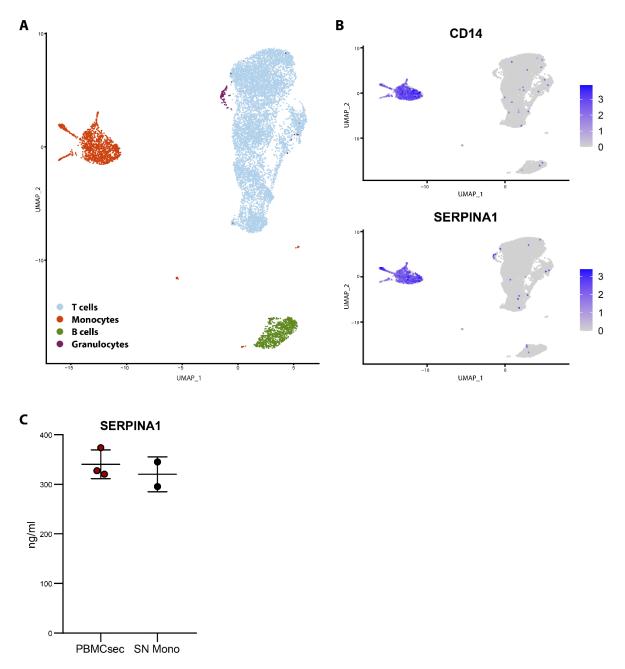
В

control	lonomycin	PBMCsec + IM	vehicle + IM
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ΗΙΕ-1α ΗΟ-1	ΗΙΕ-1α ΗΟ-1	ΗΙΕ-1α ΗΟ-1	ΗΙΕ-1α ΗΟ-1

Label	Туре	Adjusted volume	# of analysed
Laber		intensity	pixels
control	HIF-1α	109 825	13
control	HIF-1α	108 123	13
lonomycin	HIF-1α	72 267	13
lonomycin	HIF-1α	54 072	13
PBMCsec + IM	HIF-1α	160 245	13
PBMCsec + IM	HIF-1α	145 165	13
vehicle + IM	HIF-1α	112 189	13
vehicle + IM	HIF-1α	130 932	13
control	HO-1	136 363	13
control	HO-1	98 343	13
lonomycin	HO-1	34 494	13
lonomycin	HO-1	54 476	13
PBMCsec + IM	HO-1	121 447	13
PBMCsec + IM	HO-1	147 245	13
vehicle + IM	HO-1	64 068	13
vehicle + IM	HO-1	89 974	13

Supplementary Figure 5 Gating stragety and purity of isolated neutrophils and protein analysis of HIF-1α and HO-1

A) Flow cytometry gating strategy for the analysis of neutrophil purity after magnetic bead isolation is shown. Neutrophil purity was assessed by the percentage of CD66b⁺CD15⁺ cells and ranged from 92.9% to 96.5%. B) Analysed images of the proteome profiler of the protein levels of HO-1 and HIF-1 α . Isolated neutrophils of four individual donors were stimulated with ionomycin and treated with PBMCsec or vehicle. Cell lysates of four individual donors and experiments were pooled. Raw analysis values are shown in the table below.



Supplementary Figure 6 SERPINA1 abundance in PBMCsec

Single cell RNA sequencing analysis of erythrocyte lysed whole blood shows (A) UMAP cluster depiction of captured cell populations. (B) Monocyte cluster were identified by the expression of CD14. SERPINA1 expression was almost exclusively found in the CD14⁺ monocyte cluster. (C) SERPINA1 concentration in PBMCsec was analysed using ELISA. Supernatants of monocytes (SN Mono), cultured for 24 hours, were used as positive controls.

DISCUSSION

1 General discussion

Neutrophils belong to the most powerful immune cells mainly due to their capacity to form NETs as line defence mechanism of the innate immune system.¹⁶ Despite their important function in host protection and pathogen elimination, accumulating evidence proves a more complex involvement of neutrophil effector functions in health and disease.^{15,157} Particularly NET formation has been associated to detrimental host tissue damage and was shown to contribute to the progression of several diseases and pathologic conditions.^{15,157} A broad range of studies was performed aiming to pharmacologically target NET formation or NET products, however, therapeutic drugs designed to specifically target NET formation are not yet available for in-patient use.^{158,377,425} In this thesis, the effect of PBMCsec on NET formation in primary human neutrophils was investigated in an *ex vivo* setting. Our results provide evidence that PBMCsec effectively inhibits NET formation.

1.1 NET-derived extracellular DNA and histones

The overall reported inhibition of NET formation due to the application of PBMCsec treatment of activated neutrophils is indispensably associated with reduced extrusion of extracellular DNA and (modified) histones. Approximately 70% of the NETs components account for core histones and DNA.¹⁵⁶ Both, exhibit a dual role in host defence and possess potent antimicrobial as well as pro-inflammatory properties.⁴²⁶ However, extracellular DNA serves as sign for tissue damage or programmed cell death and histones promote pro-inflammatory responses as they serve as danger associated molecular patterns if present in the extracellular space.^{319,427} Modified histones, specifically citH3, were shown to majorly contribute to the pathogenesis of several pathologic conditions including acute lung injury, the disruption of the microvascular endothelial barrier and trigger positive feedback mechanisms which further potentiates NET formation.⁴²⁸ Considering these detrimental effects of NET-derived DNA and histones, our findings suggest PBMCsec as potent treatment option for pathologic conditions strikingly influenced by increased availability of citH3 and extracellular DNA within tissues and circulation.

1.2 Synergy of different PBMCsec-substance classes

Previous studies already reported that several beneficial effects of PBMCsec treatment depend on the interplay of multiple subfractions of the secretome.^{418,421} In concordance with these observations, we also observed that full inhibition of NET formation was only achieved in activated neutrophils upon treatment with the whole PBMCsec. As several PBMCsec derived fractions exerted partial inhibitory capacity we suggest that either individual factors within the protein and lipid fractions or a combination of factors found in these fractions act on different signalling molecules at different steps of the NETs signalling pathway.

It was previously shown that particularly the PBMCsec-derived lipid fraction possesses highly immunomodulatory functions as treatment of skin inflammation and allergic reactions with PBMCsec-lipids dampened dendritic cell function and reduced mast cell and basophil activation.^{422,423} Furthermore, more in depth analysis of PBMCsec components revealed that it contains several lipid species such as phosphatidylserines, phosphatidylcholines, lysophosphatidylcholines, lyso-phosphatidylethanolamines, phosphatidylethanolamines and resolvins.⁴²² Analysis of the resolvins in particular revealed that several resolvins including resolvin D1 (RvD1), RvD2, RvD3, RvD4 and RvE1, were present at increased levels in PBMCsec compared to the secretome of non-irradiated PBMCs (unpublished data).

Resolvins have been previously associated with decreasing NETosis and alleviating abdominal aortic aneurism (AAA) disease burden.⁴²⁹ Particularly RvD1 treatment was identified as potent NETosis inhibitor and was accompanied by decreased levels of the proinflammatory cytokines IL-1beta and IL-6. In parallel, the anti-inflammatory cytokine IL-10 was found to be increased upon RvD1 treatment. The specific NETs marker, citH3, was further found to be drastically decreased in RvD1 treated mice.⁴²⁹ RvD4 was therapeutically administered in an experimental animal model of deep vein thrombosis and resulted in significantly enhanced resolution of thrombosis.⁴³⁰ It was shown that RvD4 treatment reduces the number of neutrophils within the thrombus while promoting the recruitment of monocytes. RvD4 was attributed to exert anti-inflammatory effects by various mechanisms including the downregulation of adhesion molecules on immune cells, reducing the production of pro-inflammatory cytokines and diminishing leukocyte-endothelial cell interactions.⁴³⁰ Several resolvins of the resolving T series (RvTs) were shown to act on neutrophils in a dual mechanism. On the on hand, they actively target yet unidentified mediators of the NET-triggering stimuli. On the other hand, RvTs enhance NET-clearance by macrophages *in vitro* and *in vivo*.⁴³¹

Taken together, accumulating evidence indicates that certain resolvins specifically target NETformation, while others target neutrophil function and infiltration and some resolvins promote NET clearance by inducing increased phagocytosis in macrophages.⁴²⁹⁻⁴³² These studies led us to hypothesize that the NETosis inhibiting effect is in part due to the several resolinvs present in PBMCsec, and they might favour a shift in neutrophil effector functions as resolvins were observed to promote phagocytosis in macrophages.⁴³²

Furthermore, besides the lipid fraction, we also observed an inhibitory effect on NET formation by the proteins derived from PBMCsec. Previous studies revealed that calcium chelation by bovine or human serum albumin was sufficient to block NET formation in ionomycin-treated cells.⁴³³ However, other stimuli, such as PMA, were shown to robustly induce NETosis despite the addition of bovine or human serum albumin.⁴³³ Nevertheless, PMA-induced NET formation was similarly inhibited by PBMCsec treatment as calcium ionophore-induced NET formation. To further unravel which PBMCsec-derived protein or group of proteins is responsible for the observed partial inhibition of NET formation, proteomics analyses are required.

Since we observed only a partial inhibition of NET formation with the individual fractions but a strong inhibitory effect with reconstituted fractions, our data clearly indicate that this effect of PBMCsec is mediated by an interplay of the subfractions yielding a synergistic effect.

1.3 Inhibition of NETosis by a DNase-independent mechanism

Attempting to unravel the mode of action by which PBMCsec inhibits NET formation, we first had to investigate whether the observed NETs reduction was due to an active mechanism or a bystander result of DNA-degradation by DNases present in PBMCsec. Endogenous DNases function as potent host defence mechanism to maintain tissue integrity and degrade the DNA backbone of NETs.³⁷⁸ Therefore, we tested whether PBMCsec possesses active DNases capable of degrading DNA and NETs. We neither observed DNA degradation in a cell-free assay nor could detect a reduction of citH3, indicating that PBMCsec does not degrade preformed NETs. It was previously shown that the variability of endogenous DNase activity critically affects for example vascular occlusion and autoimmune diseases as reduced DNase activity is associated with severely worsened patient prognosis.^{378,434} Presence of DNase

inhibitors, genetic mutations, or the loss of endogenous DNases was observed in SLE resulting in decreased NETs clearance and aggravated disease progression.⁴³⁴ Pharmacological targeting of DNase activity has been previously suggested as potential means to alleviate NETs-induced disease burden in SLE patients. However, this approach is at a preliminary state and specific circulating DNase inhibitors in SLE patients are not yet identified.⁴³⁴ In the context of other diseases such as respiratory diseases, cancer, sepsis, and neurological disorders, DNase treatment resulted in a thoroughly positive effect on disease prognosis.⁴³⁵ However, several studies fail to prove, that the observed beneficial effects of DNase treatment are due to degradation of NETs or the prevention of coagulation. Furthermore, neither timing, route of administration or dose are coherent in the different studies.⁴³⁵ While beneficial effects of therapeutically administered DNases were partially reported, PBMCsec offers therapeutic means to inhibit NET formation rather than degrading NETs, thereby preventing host damage by other factors released during NET formation.

1.4 Calcium flux in neutrophils

Calcium represents a second messenger which is of tremendous importance for a great assortment of intracellular signalling cascades.⁴³⁶ Previous studies have highlighted its importance in immune cells and particularly in neutrophils, as it was shown that calcium is critically involved in oxidative stress, neutrophil activation, inflammatory processes and cell death.⁴³⁷⁻⁴³⁹ Independent of the mode of induction, NETosis relies on calcium as either initial inducer or downstream second messenger.^{158,166} Considering the important role of this molecule we investigated the calcium flux in ionomycin-activated and PBMCsec-treated neutrophils. Our data indicate that the observed inhibitory effect of PBMCsec is not due to calcium scavenging as there was only a marginal decrease in calcium flux observed, which was quickly restored to similar levels as detected in control samples. While store operated calcium entry (SOCE) has been comprehensively investigated, another yet poorly understood receptor-dependent mechanism has been suggested.436 SOCE, a two-step process, is accompanied by the depletion of intracellular calcium stores from the endoplasmic reticulum via sarco-endoplasmic reticulum calcium ATPase (SERCA) pumps. Subsequently, stromal interacting molecules, together with Orai proteins and transient receptor potential channels, forward this information to plasma membrane channels, which results in opening of calcium channels in order to replenish the depleted stores. ⁴³⁶ Thapsigargin, an irreversible inhibitor of SERCA pumps, has previously been used to force intracellular calcium store depletion.^{436,440} We used Thapsigargin as stimulus to exclude that PBMCsec-treated neutrophils circumvent

NETosis by sequestration of excess calcium into intracellular calcium stores. Our data revealed that Thapsigargin not only serves as potent NETosis inducer itself, but also that the addition of PBMCsec prevented NET formation. Together, these data indicate that the mode of action of PBMCsec is mediated by influencing other factors than calcium that are involved further downstream in the NETosis signalling cascade.

1.5 Akt and NFkB signalling

Both Akt and NFkB signalling have been associated with NETosis. It was shown that activation of Akt is critical for both, PMA and calcium ionophore-induced NETosis.⁴⁴¹⁻⁴⁴⁴ Interestingly, our group has previously demonstrated that PBMCsec itself functions as potent inducer of Akt activation in a diverse set of cell types, such as primary human keratinocytes, fibroblasts, endothelial cells, Schwann cells as well as astrocytes.^{412,414} Akt is known to suppress caspase signalling thereby promoting cell survival by blocking apoptosis.⁴⁴³ The previously reported cytoprotective effects mediated by PBMCsec have partially been attributed to Akt activation. However, in terms of neutrophils and particularly NETosis, our data indicate that, despite mediating Akt activation in a celltype independent manner, the NETosis inhibiting activity by PBMCsec is most likely due to the modulation of other factors further downstream of the Akt signalling cascade or generally independent of this pathway.

However, it was previously demonstrated that activation of the Akt signalling pathway inhibits Raf-MEK-ERK signalling.^{445,446} Interestingly, it was shown that pharmacological inhibition of Raf, as well as other members of the Raf-MEK-ERK signalling pathway, prevented NETosis.¹⁸¹ Furthermore, Hakkim et al. demonstrated that Raf-MEK-ERK inhibitors also block the production of ROS due to the inability of ERK to phosphorylate p47^{phox}, thus impairing functional NADPH oxidase assembly.¹⁸¹

Furthermore, immunofluorescence staining of NF κ B in ionomycin-activated neutrophils was marked by strong activation of NF κ B. Interestingly, treatment of activated neutrophils with PBMCsec did not prevent or interfere with NF κ B activation, however, also the immunofluorescence images clearly demonstrated that PBMCsec-treatment results in the inhibition of NET formation (data not shown). Together, these findings suggest that PBMCsec modulates other players involved in NETosis, that either overrule, function independently, or further downstream of NF κ B.

1.6 Prevention of ROS production

One hallmark feature of NETosis is oxidative stress, and particularly the production of ROS.¹¹⁷ Our data indicate, that one of the critical modes of action of PBMCsec in the inhibition of NET formation is the prevention of ROS production. Furthermore, our study revealed that HSPs were strongly down-regulated upon ionomycin-induced activation of NETosis. However, PBMCsec treatment of activated neutrophils counteracted the downregulation of heme oxygenase 1 (HO-1 or HSP32) as well as hypoxia inducible factor 1 alpha (HIF-1α).

1.6.1 Heme oxygenase 1

Previous studies have revealed that HO-1 upregulation occurs upon pathophysiological stimuli including endotoxemia, oxidative stress, ischemia, trauma-haemorrhage and inflammation.447-⁴⁵⁰ Furthermore, it has been reported, that this enzyme critically contributes to the protection against oxidative tissue injury.⁴⁵¹ HO-1 is known to reduce factors such as p47^{phox} and p67^{phox}, which are two important cytosolic proteins required for the assembly of functional NADPH oxidase.^{451,452} Particularly the release of superoxide anion from neutrophils requires activation of functional NADPH oxidase.⁴⁵¹ During oxidative burst, activated neutrophils are capable of producing approximately 10nmol per minute of superoxide anion per one million neutrophils.⁴⁵³ Neutrophil-derived ROS are known to directly act on endothelial cells reducing cell integrity and barrier function. During the development of lesions in atherosclerosis, ROS were reported to contribute to endothelial cell apoptosis. Additionally, ROS produced by neutrophils, particularly if excessively produced, have been linked to directly damage tissue in inflammatory bowel disease and potentially inducing gastrointestinal cancer.⁹¹ Furthermore, HO-1 knockout in a murine model of renal ischemia-reperfusion injury was reported to exacerbate disease burden by inducing an upregulation of vascular cell adhesion molecule-1, thereby favouring increased neutrophil adhesion and the activation of inflammatory responses.⁴⁵⁴

Thus, our data suggest that PBMCsec mediated maintenance of high levels of HO-1, similar to those observed in neutrophils in a non-activated state, indirectly contributes to impaired ROS production by preventing the translocation of the proteins p47^{phox} and p67^{phox} required for the assembly of the NADPH oxidase complex. Moreover, PBMCsec treatment may further mitigate inflammatory responses by impeding neutrophil adhesion.

1.6.2 Hypoxia inducible factor

The transcriptional regulator HIF-1 α is known to induce metabolic switches required for cell survival in response to hypoxic, stressed, infected or inflamed tissues. HIF-1α stabilization is required for the regulation of cellular metabolism and the mediation of gene expression.^{455,456} We observed a robust decrease of HIF-1a protein content upon neutrophil activation with ionomycin, which was counteracted by PBMCsec treatment. Interestingly, several studies have reported contradictory results on the influence of ionomycin and other calcium ionophores in regard to the increased or decreased protein levels of HIF-1a.457-460 It was previously suggested, that HIF-1a protein increase is a result of decreased intracellular calcium concentrations and attenuated proline hydroxylase activity.^{459,460} Conversely, other studies reported that HIF-1α protein abundance was not affected by ionomycin. However, the same group reported an increase in transcriptional activity of HIF-1a.458 Furthermore, it was reported that ionomycin-induced elevation of calcium concentration lead to the activation of a degradation pathway of HIF-1a.⁴⁵⁷ Nevertheless, it is noteworthy, that these inconsistent and in part contradictory findings have to be interpreted carefully as these results are based on different cell types and calcium inducing agents and are not consistent with respect to normoxia or hypoxia.

It was previously shown that the loss of HIF-1 α lead to tremendously impaired glycolysis and energy generation in macrophages and neutrophils, thereby resulting in heavily impaired effector functions including phagocytosis, intracellular killing of phagocytosed pathogens and migration.⁴⁶¹ Furthermore, evidence was provided, that HIF-1a critically contributes to neutrophil survival by suppressing apoptosis via NFB signalling.⁴⁶² Together, these observations lead us to hypothesize that the observed PBMCsec-induced upregulation of HIF- 1α in activated neutrophils is a NETosis independent mode of action. We suggest, that while PBMCsec treatment impairs NET formation, it contributes to the maintenance of a functional immune response in neutrophils via HIF-1a promoted phagocytosis and intracellular killing pathogens, accompanied by reduced neutrophil apoptosis. Together, despite not sufficiently proven yet, this mode of action would provide a potent anti-NETs specific therapeutic approach with beneficial effects in terms of wound healing where proper pathogen clearance is essential while NET formation may impair wound healing.^{305,314} Furthermore, inhibited NET formation in combination with other effector functions remaining functional would be beneficial particularly in pathologies marked by the presence of autoantibodies targeting neutrophils or NETsspecific components.⁴⁶³

1.7 PAD4 inhibition

PAD4 mediated histone citrullination represents one of the main features during NETosis, distinguishing it precisely from other forms of cell death.¹⁹⁵ PAD4 has been of high interest in research as a potential therapeutic target to inhibit NETosis due to its critical involvement in various pathologies including RA, multiple sclerosis (MS), cancer, sepsis, ischemiareperfusion injury, heart failure and myocardial infarction.⁴⁶⁴⁻⁴⁶⁶ Nevertheless, despite intense research, the particular mechanism how PAD4 inhibition is achieved, is not yet fully understood.⁴⁶⁴ Besides the prevention of ROS production, our data indicate another mode of action by acting on PAD4 activity. We observed a PAD4 inhibition by PBMCsec in a cell-free assay. This finding suggests that as a response of stress, PBMCs secrete paracrine factors that function as PAD4 inhibitors. Interestingly, umbilical cord blood derived neutrophils from preterm and term infants fail to undergo NETosis upon stimulation.⁴⁶⁷ This lacking immunecompetency is quickly gained at day three post-delivery. Sequence analysis of proteins present in day 0 umbilical cord blood revealed several protein and peptide clusters to be different from venous blood plasma at day 28 post-delivery.⁴⁶⁷ The sequence of one particular peptide was identified to be identical to the carboxy terminus of α -1-antitrypsin. This peptide showed strong NETosis inhibitory capacity even if applied to LPS-stimulated adult neutrophils.⁴⁶⁷ The mode of action of α -1-antitrypsin was reported to be via PAD4 inhibition.⁴⁶⁷ Furthermore, α -1antitrypsin, also referred as SERPINA1, is known to inhibit neutrophil elastase and proteinase 3 as well as other intracellular and cell surface proteases.⁴⁶⁸ It was previously reported, that circulating monocytes synthesize α -1-antitrypsin.⁴⁶⁹ In line with these findings, using single cell RNA sequencing analysis, we found significantly high SERPINA1 expression levels in monocytes from healthy donors. Furthermore, we observed similar amounts of SERPINA1 in PBMCsec as in the supernatant of cultured monocytes. Together, these findings indicate, that PBMCsec derived SERPINA1 contributes to the inhibition of PAD4 and ultimately the prevention of NET formation.

2 Conclusion

In summary, the observations of this study attribute a strong NETs-inhibitory activity to PBMCsec. The identification of a dual mechanism by impairing two of the major key events during NETosis, ROS production and PAD4 activity (Figure 5), provides future therapeutic opportunities for a diverse set of diseases associated with NETosis, including for example myocardial infarction, heart failure, sepsis and rheumatoid arthritis.¹⁶ PBMCsec has already been subjected to a pre-clinical toxicological assessment including intravenous and topical application, without any reported major adverse events (LPT, study number 35015). Hence, this study has paved the way for the therapeutic administration of PBMCsec in the context of NETs-associated diseases.

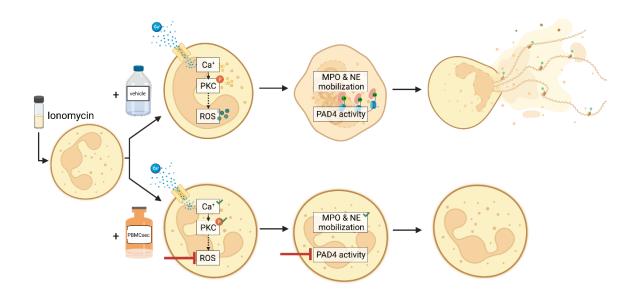


Figure 5 Graphical overview on the suggested inhibition of NETosis by PBMCsec.

Treatment of ionomycin activated neutrophils results in increased calcium influx, thereby initiating the NETosis cascade. While PBMCsec does not influence the activation and phosphorylation of Protein Kinase C (PKC), it prevents reactive oxygen species (ROS) production by a yet undefined mechanism. Myeloperoxidase (MPO) as well as neutrophil elastase (NE) activity remain unaltered by PBMCsec treatment. One of the hallmarks of NETosis, histone hypercitrullination by protein arginine deiminase 4 (PAD4), is most likely impaired by SERPINA1, present in PBMCsec.

3 Future prospective

This study has demonstrated the potent therapeutic potential of PBMCsec in terms of NETosis. Nevertheless, future studies are required to fully delineate the precise mode of action of PBMCsec. These studies should involve a comprehensive investigation of the potential cross-regulation of the Akt and Raf-MEK-ERK pathway. This will help to understand whether the observed prevention of ROS production may result from impaired ERK signalling. Furthermore, analysis of the translocation and activation of NADPH oxidase subunits such as p67^{phox} and p47^{phox}, the latter being a target of ERK, would offer further valuable insight into the modulated NETosis signalling pathway. Lastly, as the lipid and protein fraction were observed to exert the most promising effects as single substance classes, meticulous identification of individual factors present in the secretome should be performed. Lipidomics and proteomics analyses of the secretome will most likely contribute to the understanding how and why PBMCsec exerts its cytoprotective and anti-inflammatory actions in such a broad spectrum of different cell populations.

MATERIALS & METHODS

The materials and methods used for this study are described in the aforementioned manuscript.

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APPENDIX

Curriculum Vitae

Katharina KLAS, BSc, MSc

18.03.1996, Darmstadt Germany Adress: Agnesstraße 27, 3400 Klosterneuburg, Austria phone: +43 680 322 99 36 e-mail: klaskatharina@gmx.at

PROFESSIONAL EXPERIENCE

09.2022 – to date	Sanofi Aventis GmbH Medical Scientific Liaison, Immunology (Dermatology)
11.2019 - 08.2022	
11.2019 - 08.2022	Aposcience AG, Vienna Conducting PhD project with company's product (Aposec)
09.2018 – 09.2019	Internship (Master Thesis Project: Distinct distribution of RTN1A in Mouse Skin and lymphoid organs) Laboratory of Prof. Dr. Erwin Tschachler Department of Dermatology, Medical University of Vienna
07.2018 – 08.2018	Internship at Christian Doppler Laboratory for Molecular Stress Research in Peritoneal Dialysis (CDL-MSRPD), Ap.Prof. Priv.Doz. DI Dr. Klaus Kratochwill, Medical University of Vienna
06.2016 – 02.2017	Internship (Bachelor Thesis Project: Involvement of basement membrane laminins in proliferation and migration of mammary epithelial cancer cells) Laboratory of Assoc. Prof. Pekka Katajisto Institute of Biotechnology, University of Helsinki
2015	PR activities for The Asia Pacific Early Mobilization Network 1 st Annual Conference on Early Mobilization & Rehabilitation in the ICU 2015, Tokyo Japan
2013	Organizational collaboration at the conference 1st European Conference on Waning & Rehabilitation in Critically ill Patients, International Early Mobilization Network, Vienna
EDUCATION	
11.2019 – to date	Medical University of Vienna, PhD Candidate PhD Programme Vascular Biology; Pharmacological inhibition of NETosis by Aposec (in cooperation with Aposcience AG, Vienna)
2017-2019	University of Veterinary Medicine, Vienna Master's Programme Comparative Biomedicine Focus: Infection Biomedicine and Tumour Signalling Pathways

2014-2017	IMC FH Krems, University of Applied Science
	Bachelor's Programme Medical and Pharmaceutical Biotechnology
2009-2014	Hertha Firnberg Schulen for Economics and Tourism, Vienna

ADDITIONAL QUALIFICATION

2021	Certificate of the Federation of European Laboratory Science (FELASA) for
	experimental biomedical studies in animals

Educational focus: Buisness Responsibility Management

CONGRESSES AND MEETINGS

02.2022	48th ADF Annual Meeting, Arbeitsgemeinschaft Dermatologische Forschung
	Poster presentation
06.2021	ÖGDV Science Days
	Poster presentation

LANGUAGE SKILLS

German	native language (Level C2)
English	fluent in writing and speaking (Level C1-C2)
French	capable of regular conversations and writing (Level B2)
Spanish	basic language skills (A1)

IT SKILLS

MS Office, Adobe Illustrator, GIMP, RStudio/Programming language "R"

SOCIAL COMMITMENT & HOBBIES

Make-A-Wish Foundation Austria Cooking for people in need at "Kochen für die Gruft", Caritas *Riding, Biking, Photography*

PUBLICATIONS

The Effect of Paracrine Factors Released by Irradiated Peripheral Blood Mononuclear Cells on Neutrophil Extracellular Trap Formation

Klas K, Ondracek AS, Hofbauer TM, Mangold A, Pfisterer K, Laggner M, Copic C, Direder M, Bormann D, Ankersmit HJ, Mildner M. Antioxidants (Basel) 2022 Aug 11;11(8):1559; doi: 10.3390/antiox11081559

Paracrine Factors of Stressed Peripheral Blood Mononuclear Cells Activate Proangiogenic and Anti-Proteolytic Processes in Whole Blood Cells and Protect the Endothelial Barrier Copic C, Direder M, Schossleitner K, Laggner M, **Klas K**, Bormann D, Ankersmit HJ, Mildner M. Pharmaceutics 2022, 14(8), 1600; https://doi.org/10.3390/pharmaceutics14081600 - 30 Jul 2022

The secretome of irradiated peripheral blood mononuclear cells attenuates activation of mast cells and basophils.

Laggner M, Acosta GS, Kitzmüller C, Copic D, Gruber F, Altenburger LM, Vorstandlechner V, Gugerell A, Direder M, **Klas K**, Bormann D, Peterbauer A, Shibuya A, Bohle B, Ankersmit HJ, Mildner M. EBioMedicine. 2022 Jul;81:104093. doi: 10.1016/j.ebiom.2022.104093. Epub 2022 Jun 4. PMID: 35671621

Schwann cells contribute to keloid formation.

Direder M, Weiss T, Copic D, Vorstandlechner V, Laggner M, Pfisterer K, Mildner CS, **Klas K**, Bormann D, Haslik W, Radtke C, Farlik M, Shaw L, Golabi B, Tschachler E, Hoetzenecker K, Ankersmit HJ, Mildner M. Matrix Biol. 2022 Apr;108:55-76. doi: 10.1016/j.matbio.2022.03.001. Epub 2022 Mar 10. PMID: 35278628

Severity of thermal burn injury is associated with systemic neutrophil activation. Laggner M, Lingitz MT, Copic D, Direder M, **Klas K**, Bormann D, Gugerell A, Moser B, Radtke C, Hacker S, Mildner M, Ankersmit HJ, Haider T. Sci Rep. 2022 Jan 31;12(1):1654. doi: 10.1038/s41598-022-05768-w. PMID: 35102298

Secretome of Stressed Peripheral Blood Mononuclear Cells Alters Transcriptome Signature in Heart, Liver, and Spleen after an Experimental Acute Myocardial Infarction: An In Silico Analysis. Mildner CS, Copic D, Zimmermann M, Lichtenauer M, Direder M, **Klas K**, Bormann D, Gugerell A, Moser B, Hoetzenecker K, Beer L, Gyöngyösi M, Ankersmit HJ, Laggner M. Biology (Basel). 2022 Jan 13;11(1):116. doi: 10.3390/biology11010116. PMID: 35053121

Transcriptional Differences in Lipid-Metabolizing Enzymes in Murine Sebocytes Derived from Sebaceous Glands of the Skin and Preputial Glands.

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