Phagocytosis, Degranulation and Extracellular Traps Release by Neutrophils—The Current Knowledge, Pharmacological Modulation and Future Prospects

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Neutrophils are crucial elements of innate immune system, which assure host defense via a range of effector functions, such as phagocytosis, degranulation, and NET formation. The latest literature clearly indicates that modulation of effector functions of neutrophils may affect the treatment efficacy. Pharmacological modulation may affect molecular mechanisms activating or suppressing phagocytosis, degranulation or NET formation. In this review, we describe the role of neutrophils in physiology and in the course of inflammatory reaction, illustrating the versatility and plasticity of those cells. This review also focus on the action of plant extracts, plant-derived compounds and synthetic drugs on effector functions of neutrophils. These recent advances in the knowledge can help to devise novel therapeutic approaches via pharmacological modulation of the described processes.

Keywords: neutrophils, phagocytosis, degranulation, extracellular traps, inflammation

NEUTROPHILS IN HEALTH AND DISEASE

Role in Health

Neutrophils are produced in the venous sinuses of the bone marrow, where they derive from a common myeloid progenitor cells (Figure 1; Nerlov and Graf, 1998). Recently, a novel classification of bone marrow neutrophil-lineage cells emerged. Progenitor stem cells evaluate into a pool of preneutrophils, further differentiating into non-proliferative immature neutrophils and, eventually, mature neutrophils, equipped with numerous machineries to combat pathogens. Maturation of this lineage is orchestrated by transcription factors such as GFH1 and Pu.1 (Evvard et al., 2018). Neutrophils are continually generated in the bone marrow (daily production may reach up to 2 × 10¹¹ cells), where also begin the maturation process (Borregaard, 2010). Granulocyte-colony stimulating factor (G-CSF) is the main agent responsible for their development, production and release (Lieschke et al., 1994). In vivo studies showed that neutrophils may circulate in human blood for ~10 h (Athens et al., 1961a; McMillan and Scott, 1968) to 5 days (Pillay et al., 2010). Nonetheless, in the course of the development of inflammatory reaction, neutrophils become active, and their lifespan is extended. This is a result of interactions with various cytokines, (e.g. IL-1 beta, tumor...
necrosis factor (TNF), IFN-gamma), as well as bacterial products, such as lipopolysaccharide (LPS) (Colotta et al., 1992; Summers et al., 2010; Kim et al., 2011). Mature neutrophils (with segmented nuclei and fully formed granules) may also proliferate outside of the bone marrow—in the spleen after infection (Deniset et al., 2017) or when they come in contact with serum amyloid A (De Santo et al., 2010). Longer lifespan allows them to shape many processes: tissue healing, inflammation resolution and modulation of adaptive immune responses, but also may cause adverse effects, including tissue damage (Liew and Kubes, 2019). Neutrophils’ maturation and aging goes in parallel with natural drift in their phenotype and function in the absence of inflammation. These cells undergo daily modifications (Casanova-Acebes et al., 2013), such as expression of adhesion molecules affecting tissue trafficking, (e.g. CD62L, CD11b, and CD49 days), expression of chemokines receptors, (e.g. CXCR2 and CXCR4), pathogen receptors (Toll-like and NOD-like receptors), activity of pathways related to cell activation (such as NF-kB and MAPK signaling), and cell death (as revised in (Adrover et al., 2016). This process is influenced by the microbiome, as its depletion with broad-spectrum antibiotics reduces neutrophil number and aging, which can be reversed by LPS stimulation. Similar results were observed in germ-free and TLR2, TLR4, and Myd88 knock-out mice. Neutrophils not stimulated by microbiota are less responsive, including bone marrow (where they inhibit hematopoietic niches and trigger the release of hematopoietic progenitor cells into the bloodstream), spleen (mainly in the marginal zone and red pulp; supporting B cell maturation and antibody production), liver, lung (where they prevent formation of melanoma metastases and affect lung transcriptome in diurnal pattern), intestine (where phagocytosis of incoming neutrophils by resident macrophages regulates cytokine production and participates in the remote regulation of bone marrow niches), white adipose tissue, skin, skeletal muscle, peripheral lymph nodes, intestine (as isolated patches in the intestine) kidneys, and heart. Only immune-privileged organs, i.e., ovaries, testes, and brain are free from neutrophils. Apoptotic neutrophils located in skin, muscle, and intestine are phagocyted by dendritic cells and macrophages, which inhibits IL-23 expression and thus controls granulopoiesis by regulating G-CSF production (Casanova-Acebes et al., 2013). How neutrophils persist in these organs is not fully elucidated yet. Intravascular transit time in the liver is around 2 min (Peters et al., 1985), in the spleen and bone marrow—10 min (Peters et al., 1985; Ussov et al., 1995). Recently, two pools of neutrophils were identified in the spleen. Mobile neutrophils were roaming around the red pulp, while the stationary ones remained in the perivascular area, both populations differ in terms of Ly6G expression. Mature, (i.e. Ly6G<sup>high</sup>) splenic neutrophils were necessary to eradicate Staphylococci during systemic infection. This process takes place mainly in the red pulp, whereas immature neutrophils, (i.e. Ly6G<sup>int</sup>), residing perivascular area, undergo emergency proliferation and mobilization (Deniset et al., 2017) Using intravital microscopy large quantities of neutrophils resident in the lung under normal conditions were visualized (Kreisel et al., 2010). Other study showed that neutrophils in this location are not simply “stuck” in the capillary bed, but rather are on the move—tethering, crawling or adhered to cell surfaces (Yipp et al., 2017). Lung is an organ with a high number of neutrophils in margined pool, maintained due to expression of adhesion molecules on neutrophils themselves and on endothelial cells. After stimulation, such as infection, profile of adhesion molecules evolve quickly, allowing transmigration (Sibille and Marchandise, 1993). Moreover, the persistence time in this organ is way longer than the mean neutrophil intravascular transit time estimated for other organs (Peters et al., 1985; Lien et al., 1987; Ussov et al., 1995).

Granulocytes’ clearance from the circulation takes place mainly in the liver, spleen, and bone marrow (Shi et al., 2001;
As they grow older, the expression of CXCR4 increases, which probably leads them back to the bone marrow, where they are eliminated (Casanova-Acebes et al., 2013). The same receptor also negatively regulates release of the newly formed neutrophils from the bone marrow (Eash et al., 2009; Summers et al., 2010). CXCR4 is not the only factor responsible for the destruction of neutrophils as cells that lack this receptor have the same half-life as the wild type neutrophils (Eash et al., 2009). Granulocytes may also die in the blood vessels and are later removed by the Kupffer cells (liver-resident macrophages) (Shi et al., 2001). The neutrophils clearance by Kupffer or dendritic cells is regulated, in part, by the IL-23, IL-17 and granulocyte colony-stimulating factor (G-CSF) cytokine axis. These mediators stimulate granulocyte production in the bone marrow as well (Stark et al., 2005). There are still significant gaps in the knowledge about the neutrophil clearance by other cells. Neutrophils die primarily by an intrinsic mechanism of apoptosis, regulated by Bcl-2 protein family. MCL-1 is the key anti-apoptotic agent, antagonizing the pro-apoptotic effects of remaining Bcl-2 family members. Severe neutropenia without affecting other cells circulating in the blood was observed in MCL-1 knock-out mice (Csepregi et al., 2018). During the resolution of inflammation, macrophages dispose apoptotic neutrophils by phagocytosis, releasing lipid mediators. These molecules play a role in re-establishing homeostasis and their absence likely leads to chronic inflammation (Buckley et al., 2014). Some neutrophils sacrifice themselves, releasing neutrophil extracellular traps (NETs) to fight pathogens (Brinkmann et al., 2004; Stark et al., 2005). Neutrophils are also capable of vital netosis—i.e., releasing NETs by a living cell (Yipp and Kubes, 2013).

Significant heterogeneity and functional versatility are observed among neutrophils (Scapini et al., 2016; Silvestre-Roig et al., 2016; Deniset and Kubes, 2018). Both, in steady state and in the course of inflammation, these cells undertake various roles, not always beneficial. It is still unclear whether this variety results from different programs of neutrophils’ maturation and activation or is somehow influenced by external factors. In patients with autoimmune disorders, low density neutrophils/granulocytes were more frequent, though their functions are not precisely defined (Hackbarth and Kajdacsy-Balla, 1986; Scapini et al., 2016). Polymorphonuclear myeloid-derived suppressor cells (PMN-MDSCs) are responsible for the failure of many cancer therapies and poor clinical outcomes. They derive from pathologically activated neutrophils, however, the exact mechanism is of their action is unclear (Gabrilovich, 2017). Two distinct fractions of tumor-associated neutrophils are identified in cancer patients. N1— a pro-inflammatory and anti-tumor subset, induced by TGF-β blockade and N2—a protumor group, increasing in number, following stimulation by TGF-β (Fridlender et al., 2009). The concept of neutrophils’ heterogeneity cannot be universally summarized. We still see the vast gaps in the knowledge that preclude a succinct explanation for this phenomenon. Nevertheless, this heterogeneity may have a significant impact on choosing potential targets for future therapeutic agents (Nemeth et al., 2020).

To exert their effects, neutrophils must first reach the target tissue. The recruitment of granulocytes into the inflamed site involves tethering, rolling, adhesion, crawling and transmigration (Kolaczkowska and Kubes, 2013). The entire process is initiated via activated endothelium that exposes adhesion molecules, that enable leukocytes recruitment. Activation occurs via stimulation by mediators such as histamine, cysteinyl-leukotrienes and cytokines, usually released by resident leukocytes, when the pathogens are present (Ley et al., 2007). Endothelial cells may also be activated directly. If pattern-recognition receptor (PRR) connects with its ligand, the number of adhesion molecules increases on the endothelium surface. P-selectin and E-selectin are responsible for further neutrophil recruitment steps (Petri et al., 2008). They bind to their ligands, including P-selectin glycoprotein ligand 1 (PSGL1), capturing free-flowing neutrophils to the endothelium surface and promote subsequent granulocytes’ rolling along the vessel. Adhesion, crawling and transmigration depend, to a large extent, on integrin interactions with cell adhesion molecules (CAMs) (Phillipson et al., 2006).

Luminal surface of endothelium exposes chemokines, which activate rolling neutrophils, thus inducing conformational changes and completing the extravasation process. Neutrophils are capable of returning to the bloodstream via a process called reverse transendothelial cell migration (rTEM). Neutrophils that underwent rTEM are characterized by high CD54 and low CXCR1 expression and are identified more frequently in case of systemic inflammation than in healthy donors (Buckley et al., 2006). Other forms of neutrophils’ motility are: 1) reverse luminal crawling—moving along blood vessel against blood flow; 2) reverse abluminal crawling—occurs while seeking essential directional cues to fully breach venular walls and 3) reverse interstitial migration—a movement directed away from inflammation site within interstitium, which may result in remote organ damage, as documented in ischemic injury (Nourshargh et al., 2016).

Role in Disease Neutrophils play key roles in many diseases, protecting against pathogens and regulating innate and adaptive immunity. On the other hand, when hyperactive or abnormally stimulated, they may lead to tissue damage and exacerbate existing pathology.

During sepsis, neutrophils show enhanced responsiveness to chemokines, resulting in their accumulation at the infection site (Angus et al., 2001; Martin et al., 2003; Sônego et al., 2016). Some hypotheses suggest that sepsis is associated with an early overwhelming innate immune response, characterized by dysregulation of the overproduction of cytokines (TNF-α, IL-1β, IL-6, IL-8) (Benjamim et al., 2004; Faik, 2013; Chousterman et al., 2017). Concentrations of circulating pro-inflammatory cytokines are low or undetectable in healthy individuals but their production is stimulated during invasion by pathogenic microorganisms. In human and experimental animal models of sepsis, cytokines are released in a sequential manner resulting in a “cytokine cascade” (Steinhauser et al., 1999). It is initiated when a stimulus, such as Gram-negative bacterial endotoxin, (e.g. lipopolysaccharides released by E. coli), induces production of the “early inflammatory cytokines,” like TNF-α and IL-1β. TNF-α
was shown to be a central factor of immune regulation and mediator of the pathophysiological changes associated with bacteremia and sepsis syndrome (Benjamim et al., 2004). Overproduction of TNF-α correlates with enhanced properties of phagocytes. In contrast, IL-1β serum levels are only slightly increased during sepsis. The release of “early inflammatory cytokines,” intensifies the production of the “late inflammatory cytokines”—IL-6 and IL-8. IL-6 was recognized as a marker of sepsis with high specificity (Kuster et al., 1998). The increased plasma IL-8 concentration in adult sepsis-occurring patients may correlate with mortality (Marty et al., 1994). Although various pro-inflammatory cytokines contribute to the inflammatory cascade, other cytokines also display anti-inflammatory properties, serving to counterbalance a potentially inadequate pro-inflammatory state. In sepsis, interleukin 10 has been shown to act as the primary endogenous modulator of inflammatory response.

The dysregulation of “cytokine balance” influences of the immune cells functions including phagocytosis and NETosis. The question is: how may cytokine profile affect those processes? Where is the balance between “too much” and “not enough”? A similar question was asked by Garner and colleagues (Garner et al., 1996), who showed that the increasing concentration of TNF-α and IL-1β may suddenly upregulate Fc receptor (FcγR)-mediated phagocytosis by human polymorphonuclear neutrophils (PMN). Nevertheless, the mechanisms of this enhanced phagocytosis remain unknown. This same issue was investigated by Erwig and colleagues. They exposed macrophages on single cytokine and cytokine mixture (IFN-γ, TNF-α, TGF-β, IL-4, IL-6, and IL-10). After 48 h their function was evaluated for nitric oxide (NO) generation, uptake of apoptotic neutrophils, and β-glucuronidase expression. The phagocytic properties of leukocytes were augmented by TNF-α (40 vs 29% controls) and decreased by IFN-γ, IL-10, and IL-4 (Erwig et al., 1998).

Similarly to phagocytosis, NET formation (Galluzzi et al., 2018) may be effective anti-microbial mechanism in the course of sepsis, however with time it may contribute to tissue and endothelial damage, finally leading to disseminated intravascular coagulation (DIC), with high mortality rate. Interestingly, both SOFA (Sequential Organ Failure Assessment) score and acute kidney injury (AKI) correlates with cell-free DNA, measured in septic patient serum (Klopf et al., 2021).

Some studies highlight neutrophils contribution to antiviral immunity (Saitoh et al., 2012; Jenne et al., 2013). In the course of COVID-19, neutrophils-to-lymphocytes ratio was established as an independent prognostic factor (Wang et al., 2020). Recent study showed that increased amounts of reactive oxygen species (ROS) and the release of NETs are associated with intensive thrombi formation—one of the most prevalent and serious COVID-19 complications (Arcanojo et al., 2020). In COVID-19 patients, cell-free DNA, MPO-DNA complexes, and citrullinated histone H3 have been assessed and appeared to be elevated. Interestingly, cell-free DNA and MPO-DNA complexes were significantly higher in the blood from mechanically ventilated vs. non-ventilated patients. It was suggested that vascular damage, resulting in acute respiratory distress syndrome (ARDS) and multigorgan dysfunction, may be due to NET formation (Klopf et al., 2021). The abovementioned findings may link cytokine storm (as discussed in Ragab et al., 2020), microangiopathic occlusions and ARDS, and justifies use of steroids, (i.e. dexamethasone, as shown in RECOVERY trial (Horby et al., 2021)) and tocilizumab (Zhang et al., 2020b) in COVID-19 treatment.

In the respiratory system, neutrophils act like a double-edged sword. On the one hand, they defend organism against pathogens, but on the other they cause extensive tissue damage in e.g., acute lung injury, chronic obstructive pulmonary disease, or neutrophilic asthma (Grommes and Soehnlein, 2011; Meijer et al., 2013; Krishnamoorthy et al., 2018). Neutrophils also contribute to formation and destabilization of atherosclerotic plaques (Soehnlein, 2012). The accumulation of these cells is associated with a greater risk of plaque rupture (Ionita et al., 2010). Moreover, neutrophils accumulated at the plaque site may release NETs, which leads to smooth muscle cells damage, prompting destabilization (Silvestre-Roig et al., 2019).

In ischemic diseases, such as myocardial infarction or stroke, inflammatory processes cause severe tissue damage, especially after reperfusion (Eltzschig and Eckle, 2011). In myocardial infarction model, Ly6C<sup>high</sup> monocytes and neutrophils contributed comparably to proteolysis, which may result in left ventricle rupture (Anzai et al., 2017). As mentioned above, neutrophils may undergo rTEM, stimulated by LTβ<sub>4</sub> from area of ischemia-reperfusion injury and contribute to distant organ damage (Colom et al., 2015).

The role of NET formation in autoimmune disease was reviewed by Mitsios et al. (Mitsios et al., 2016) and Lee et al. (Lee et al., 2017). Some authors suggest that the anti-DNA immunization in systemic lupus erythematosus may be initiated by NETs. Released nucleic acid activates the dendritic cells, which in turn prompt antibodies production by B cells, causing systemic autoimmunity (García-Romo et al., 2011; Lande et al., 2011). Consistently, citrullinated histones, present in NETs were suggested as the trigger activating anti-citrullinated peptide autoantibodies (ACPs) production, prevalent in rheumatoid arthritis (Khadpur et al., 2013). Release of proteinase 3 (PR3) and myeloperoxidase (MPO) may initiate antineutrophil cytoplasmic antibody-associated vasculitis (Kessenbrock et al., 2009).

Cancer biology is influenced by neutrophils’ behavior. As mentioned before, neutrophils may exert pro-inflammatory or pro-tumoral effects (Fridlender et al., 2009). This phenotypic switch and overall plasticity may explain the vast spectrum of neutrophil activities, both harmful and beneficial to the host (Sagiv et al., 2015). Some of the undesirable effects include suppression of T cell-mediated antitumor immunity (Coffelt et al., 2015; Steele et al., 2016) or elastase-mediated degradation of IRS1 (Houghton et al., 2010), enhancing cancer cell proliferation. On the other hand, some studies reported that neutrophils prevent metastasis formation (Granot et al., 2011) or inhibit early cancer growth (Blausell et al., 2015). In solid tumors, increased neutrophils number in tumor microenvironment is a negative outcome predictor (Templeton et al., 2014; Gentles et al., 2015).
TO PHAGOCYTOSE, TO DEGRANULATE OR TO MAKE EXTRACELLULAR TRAPS?

Neutrophils are crucial elements of innate immune system, which assure host defense via a range of effector functions, such as phagocytosis, degranulation, and NET formation (Figure 2; Kolaczkowska and Kubes, 2013).

Phagocytosis is a fundamental mechanism for the effective elimination of disease-causing agents (Schumann, 2016). It enables clearance of microbes, but also dead cells and tissue debris (Jaumouillé and Waterman, 2020). Thus, phagocytosis is a fundamental component responsible for tissue homeostasis and the innate immune response (Jaumouillé and Waterman, 2020). This process is initialized by internalization of targeted organism/particle. Two different mechanisms have been identified for the internalization of diverse particulate material: 1) the trigger mechanism where discreet signaling initiates formation of plasma membrane protrusions, shaped by actin, that surround nearby material, and 2) the zipper mechanism where cell surface receptors sequentially bind to ligands on the target particle, which leads to a complete wrapping of the particle by the plasma membrane (Nordenfelt and Tapper, 2011). Only a small number of pathogens, (e.g., Salmonella or Shigella) is able to initiate the process, whereas the zipper mechanism, involving a wide spectrum of phagocytic receptors, allows to successfully bind many species of pathogen/particles (Jaumouillé and Waterman, 2020). The essential phagocytic receptors are Fcγ class receptors, which recognize IgG, the complement receptor 3 (αMβ2 integrin). While TLRs and NOD receptors are not classified as phagocytic receptors, their activation may increase phagocytosis (Nordenfelt and Tapper, 2011). Receptor activation initiates signaling cascades that re-model lipids in the cell membrane and leads to rearrangement of actin cytoskeleton in order to extend the cell membrane around the particle. After reconstructing the plasma membrane of the phagocyte, bacteria are ultimately internalized in phagosome (Fairn and Grinstein, 2012). The dynamics of phagosome maturation is highly dependent on the several regulators like Rab proteins synthesis. Among known Rab proteins, Rab5 and Rab7 are directly involved in regulation of phagosome functions (Mottola, 2014).

Unlike macrophages, maturation of neutrophilic phagosomes is dependent on fusion with preformed granules (Nordenfelt and Tapper, 2011; Gierlikowska et al., 2020b). Membrane proteins and granule contents are directed to distinct locations by precise trafficking and fusion/fission processes (Nordenfelt and Tapper, 2011). Activation of granules is assisted by recruitment of NADPH oxidase (Johansson et al., 1995). It is suggested that an early alkalization of neutrophil phagosomes leads to the oxidative burst in the phagosomes (Segal et al., 1981).
Besides ROS production, neutrophils eliminate pathogens via production of microbicidal molecules following degranulation (Mortaz et al., 2018).

Degranulation is important for pathogen killing, but also modulates the immune response during infectious or non-infectious diseases (Mortaz et al., 2018). Different granule types are released sequentially (Kolaczkowska and Kubes, 2013). At the beginning, tertiary granules are released, (e.g. cathepsin, gelatinase, known as metalloprotease 9), then secondary granules, (e.g. lactoferrin, alkaline phosphatase, lysozyme, cathelicidin, NADPH oxidase and collagenase). At the end, primary granules are secreted containing the most pro-inflammatory and antimicrobial proteins, such as elastase, MPO, cathepsin G, defensins (HNP-1, HNP-2 and HNP-3 peptides) and bacterial permeability increasing protein (BPI) (Faurschou and Borregaard, 2003). Normal neutrophil degranulation involves the release of primary and secondary granules mainly into the phagosome, which prevents host tissue damage (Sengeløv et al., 1995). Bacteria are able to disrupt, dysregulate, or induce excessive neutrophil degranulation, in order to diminish the protective effects of neutrophil degranulation in a way that ultimately benefits the pathogen and extend the disease course, thus pharmacological modulation may potentially bring promising results for recovering (Eichelberger and Goldman, 2020).

Neutrophils use a few strategies to counteract infections. Two of those strategies have been thoroughly studied and are described above: phagocytosis and degranulation. They can also utilize NADPH oxidase to form ROS which have antimicrobial potential (Mantovani et al., 2011). A new mechanism that combats infections—NET formation—has been described in 2004 (Brinkmann et al., 2004). Once the process is initiated, the chromatin in neutrophils’ DNA loosens up and it forms complexes with numerous granular and cytoplasmic proteins, which are subsequently released into the extracellular space. Depending on the origin of proteins, they could by classified as histones (H1, H2A, H2B, H3, and H4), azurophilic granules (MPO, elastase, defensins, cathepsin G), specific granules (lactoferrin, alkaline phosphatase, NADPH oxidase, lysozyme, collagenase), tertiary granules (gelatinase, cathepsin) and cytosol proteins (LL-37, proteinase 3, neutrophil serine protease 4 and tryptase) (Brinkmann et al., 2004). The biological function of extracellular traps is to sequester a specific space, enabling accumulation of high concentration of antimicrobial agents, preferably at the site of infection. Thus extracellular traps immobilize and kill microorganisms, which prevent them from dissemination (Manda et al., 2014).

Modulations of killing mechanisms (like phagocytosis, degranulation, or NET formation) could be of interest, as it may potentially boost pathogen killing or protect host from own tissue damage.

**PHARMACOLOGICAL MODULATION OF PHAGOCYTOSIS VIA MEDICAL PLANTS**

Therapies targeted at innate immune response modulation via medicinal plants and metabolites isolated from them experience a renaissance in recent years (Ríos, 2010; Licciardi and Underwood, 2011; Grigore et al., 2020; Behl et al., 2021).

After performing a literature search using relevant MeSH terms and keywords, we identified 53 plant extracts, which were tested on neutrophil model and their influence on phagocytic properties were evaluated (Table 1). Selected plants belonged to diverse families: Anacardiaceae, Araceae, Araliaceae, Arecaceae, Apocynaceae, Asphodelaceae, Asteraceae, Boraginaceae, Capparaceae, Celastraceae, Convolvulaceae, Cucurbitaceae. Elaeocarpaceae, Euphorbiaceae, Fabaceae, Geraniaceae, Hypericaceae, Lamiaceae, Lythraceae, Malvaceae, Melastomataceae, Meliaceae, Menispermaceae, Moraceae, Moringaceae, Phyllanthaceae, Pooaceae, Ranunculaceae, Rosaceae, Rubiaceae, Santalaceae, Sapindaceae, Saxifragaceae, Theaceae, Zingiberaceae, and Zosteraceae. Plant materials were collected in Europe, Asia, Southern and Northern America and Africa. The geographic location directly determines the phytochemical composition of plants and their pharmacological activity (Czerwińska and Melzig, 2018; Kiss et al., 2020). Thus, evaluation of the medical plants collected from different locations, may highlight future directions for searching other plants (with similar chemical composition), and then select them for compounds isolation (Gierlikowska et al., 2020a). We noticed that 33 of the selected plant extracts stimulated phagocytic activity of neutrophils, 18 decreased phagocytic activity and 2 extracts did not affect the process.

Pharmacological modulation of phagocytosis may affect the uptake of pathogens, biochemical changes inside of phagos- and lysosomes, phagolysosomes formation and modulation of intracellular killing via oxygen-dependent and oxygen-independent degradation (Table 1).

Among selected plant extracts, two of them, obtained from *Tinospora crispa* (L.) and *Heliotropium sarmentosum* (Lam.), modulated expression of surface receptors. The neutrophils treated with *T. crispa* as well as *H. sarmentosum* extracts overexpressed MAC-1 cell surface receptor leading to increased phagocytic activity (Chen et al., 2014; Ahmad et al., 2015). MAC-1 is a complement receptor (“CR3”) consisting of CD11b (integrin αM) and CD18 (integrin β2) (Todd, 1996). The integrin CD11b is responsible for direct binding to intercellular adhesion molecule-1 (ICAM-1) leading to firm adhesion to endothelium and transmigration of phagocytes to infected tissues (Ross and Větvička, 1993). Considering that *T. crispa* has been used traditionally in the treatment of rheumatic arthritis, urinary tract infections, fever, inflammation, fracture, and hypertension (Ahmad et al., 2015), the modulation of phagocytic properties of neutrophils creates new research perspectives for further evaluation. Based on the use in traditional medicine, *H. sarmentosum* has demonstrated anti-inflammatory, antinociceptive and antipyretic activities (Chen et al., 2014), thus similarly to *T. crispa* modulation of phagocytosis may point to unknown immunomodulatory properties of this extract.

Pharmacological stimulation of C3b and Fcγ expression on neutrophil surface was proved for *Echinacea purpurea* (L.) *Moench* (Melchart et al., 1995; Isaykina et al., 2008).
TABLE 1 | List of selected plant material with documented impact on phagocytosis.

| Plant family    | Species                        | Plant material | Cellular target and mechanism of action                                                                 | Effect on phagocytosis                                      | References                                      |
|-----------------|--------------------------------|----------------|----------------------------------------------------------------------------------------------------------|-----------------------------------------------------------|------------------------------------------------|
| Anacardiaceae   | Semecarpus anacardium L.f      | Plant extract  | Increase of reactive oxygen species production and lysosomal enzymes activity (acid phosphatase and cathepsin D) | Increased phagocytic activity                              | Ramprasath et al. (2006)                        |
| Araceae         | Lemna minor L                  | Plant extract  | n.d                                                                                                     | Increased phagocytic activity                              | Hu et al. (1995)                                |
| Araliaceae      | Panax ginseng C.A.Mey          | Plant extract  | n.d                                                                                                     | Increased phagocytic activity                              | Hu et al. (1995)                                |
| Areceae         | Areca catechu L                | Plant extract  | Suppression of CR1, CR3, CR4 and fcy-receptors expression, decrease of F-actin stability                  | Inhibition of phagocytic activity                           | Hung et al. (2006); Lee et al. (2014)            |
| Apocynaceae     | Leptadenia pyrotechnica        | Plant extract  | n.d                                                                                                     | Increased phagocytic activity                              | Rasheed et al. (2016)                           |
| Asphodelaceae   | Aloe vera (L.) Burm.f          | Plant extract  | Reduction the availability of intracellular free Ca^{2+} and inhibition of reactive oxygen species production | Increased phagocytic activity                              | Ou, (1991)                                      |
| Asteraceae      | Echinacea purpurea (L.) moench | Plant extract  | Stimulation of C3b and fcy receptors                                                                       | Increased phagocytic activity and digestive capacity       | Meichart et al. (1995); Yang et al. (2002); Isaykina et al. (2008) |
| Boraginaceae    | Echium amoenum fisch. and C.A.Mey | Plant extract  | Increase of reactive oxygen species production                                                           | Increased phagocytic activity                              | Figueiredo-Rinzel et al. (2017)                 |
| Boraginaceae    | Heliotropium seratensum (lam.) | Plant extract  | Increase of MAC-1 cell surface expression and activation of AKT signaling pathway                         | Increased phagocytic activity                              | Chen et al. (2014)                              |
| Elaeocarpaceae  | Capparis zeylanica L           | Plant extract  | n.d                                                                                                     | Increased phagocytic activity                              | Ghule et al. (2006)                             |
| Elaeocarpaceae  | Elaeocarpus angustifolius      | Plant extract  | n.d                                                                                                     | Increased phagocytic activity                              | Chang et al. (1997)                             |
| Fabaceae        | Guilandina bonduc L            | Plant extract  | Stimulation of phagosome-lysosome fusion                                                                | Increased phagocytic activity                              | Miyazaki et al. (2005)                          |
| Fabaceae        | Vigna mungo (L.) hepper        | Plant extract  | Stimulation of syk phosphorylation                                                                     | Increased phagocytic activity                              | Sudam et al. (2017)                             |
| Geraniaceae     | Geranium sanguineum L          | Plant extract  | Stimulation of syk phosphorylation                                                                     | Increased phagocytic activity                              | Sudam et al. (2017)                             |
| Hypericaceae    | Hypericum perforatum L         | Plant extract  | Stimulation of lysosomal enzymes activity                                                                | Increased phagocytic activity                              | Fidan et al. (2008)                             |
| Lamiaceae       | Ocimum tenuiflorum L           | Plant extract  | Stimulation of lysosomal enzymes activity                                                                | Increased phagocytic activity                              | Mukherjee et al. (2005)                         |
| Lamiaceae       | Ajuga reptans L                | Plant extract  | Inhibition of reactive oxygen species production                                                         | Increased phagocytic activity                              | Toiu et al. (2018)                              |

(Continued on following page)
TABLE 1 | (Continued) List of selected plant material with documented impact on phagocytosis.

| Plant family | Species | Plant material | Cellular target and mechanism of action | Effect on phagocytosis | References |
|--------------|---------|----------------|----------------------------------------|-------------------------|------------|
| Lamiaceae    | Ajuga genevensis L | Plant extract | Inhibition of reactive oxygen species production | Inhibited phagocytic activity | Tolu et al. (2019) |
| Lythraceae   | Punica granatum L | Plant extract | n.d | No influence on phagocytosis | Oliveira et al. (2010) |
| Malvaceae    | Grewia asiatica L | Plant extract | Stimulation of reactive oxygen species production | Increased phagocytic activity | Mesak et al. (2013) |
| Malvaceae    | Osbeckia octandra DC. | Plant extract | Inhibition of reactive oxygen species production | Inhibited phagocytic activity | t Hart et al. (1990) |
| Meliaceae    | Azadirachta indica A.Juss | Plant extract | n.d | Inhibited phagocytic activity | van der Nat et al. (1988) |
| Menispermaeae | Tinospora cordifolia (wild.) Hook.f. and thomson | Plant extract | Stimulation of lysosomal enzymes activity | Increased phagocytic activity | Rege et al. (1993); Mukherjee et al. (2010); Sharma et al. (2012) |
| Menispermaeae | Tinospora crispa (L.) Hook.f. and thomson | Plant extract | Increase of MAC-1 cell surface expression | Increased phagocytic activity | Ahmad et al. (2015) |
| Moraceae     | Ficus racemosa L | Plant extract | n.d | Increased phagocytic activity | Heroor et al. (2013) |
| Moraceae     | Moringa oleifera lam | Plant extract | n.d | Increased phagocytic activity | Gupta et al. (2010); Sudha et al. (2010) |
| Phyllanthaceae | Phyllanthus debils J.G.Klein ex wild | Plant extract | Inhibition of reactive oxygen species production | Inhibited phagocytic activity | t Hart et al. (1990) |
| Phyllanthaceae | Phyllanthus amarus schumach. and thonn. | Plant extract | Inhibition of MAC-1 cell surface expression | Increased phagocytic activity | Ilranguan et al. (2015) |
| Poaceae      | Saccharum officinarum L | Plant extract | Stimulation of reactive oxygen species production | Increased phagocytic activity | El-Abasy et al. (2002); Lo et al. (2009); Chen et al. (2012) |
| Ranunculaceae | Nigella sativa L | Plant extract | Inhibition of superoxide anion generation | Increased phagocytic activity | Haq et al. (1995) |
| Rosaceae     | Crataegus laevigata (poir.) DC. | Plant extract | Inhibition of reactive oxygen species production | Inhibited phagocytic activity | Dalli et al. (2008) |
| Rubiaceae    | bixa cocinea L | Plant extract | n.d | Increased phagocytic activity | Wickramaisinghe et al. (2014) |
| Santalaceae  | Viscum album L | Plant extract | n.d | Increased phagocytic activity | Hajito et al. (1989); Fidan et al. (2008) |
| Sapindaceae  | Acer pictum subsp. Monica (maxim.) H.Ohashi | Plant extract | Stimulation of reactive oxygen species production | Improved neutrophil uptake capacity | An et al. (2013) |
| Saxifragaceae | Bergenia crassifolia (L.) fritsch | Plant extract | Improved neutrophil uptake capacity | Increased phagocytic activity | Popov et al. (2005) |
| Theaceae     | Camellia sinensis (L.) kuntze | Plant extract | Reduction of TLR-4, iK and CD11 b expression | Inhibited phagocytic activity | Albuquerque et al. (2016) |
| Zingiberaceae | Zingiber zerumbet (L.) roscoe ex sm | Plant extract | Decreased oxidative burst and ROS production | Inhibited phagocytic activity | Akhtar et al. (2019) |
| Zosteraceae  | Zostera marina L | Plant extract | n.d | Increased phagocytic activity | Zaporozhets et al. (1991) |

n.d - not defined

E. purpurea has a well-documented biological effect in a variety of diseases, particularly bacterial and viral infections. Recently, many studies are focused on immunomodulatory effects of echinacea. The phytochemical analysis confirmed that alkamides, flavonoids, derivatives of quercetin, kaempferol and caffeic acid, are considered important for biological activity (Barnes et al., 2005).

Popov and colleagues showed that Bergenia crassifolia (L.) treatment resulted in stimulation of neutrophilic uptake capacity (Popov et al., 2005). This observation justifies the traditional use of B. crassifolia as a treatment of bacterial infections (Shikov et al., 2014). The plant presents a multitude of bioactive agents, including tannins (pedunculagin, tellimagrandin I), polysaccharides, flavonoids (bergenin, kaempferol, and...
quercetin derivatives) aldehydes (2,4-heptadienal, benzaldehyde), terpenes (thymol, geraniol), phenolics (arbutin, ellagic acid, hydroquinone), phenolic acids (caffeoyl quinic acid, gallic acid, quinic acid) and other compounds (bergenan) (Shikov et al., 2014). The selected compounds are typical for Bergenia species and exerts anti-inflammatory, antimicrobial, antioxidant, and immunomodulating properties (Shikov et al., 2014). A well-studied compound is bergenin, which stimulates the uptake of apoptotic human neutrophils and then intensifies the ROS production by mouse macrophages (Shikov et al., 2014).

**Tinospora cordifolia** (Willd.), *Ocimum tenuiflorum* L. and *Semecarpus anacardium* L. stimulated lysosomal enzymes activity increasing phagocytic properties of neutrophils (Mukherjee et al., 2005; Ramprasad et al., 2006; Sharma et al., 2012). *T. cordifolia* extract was tested on *Escherichia coli*-induced peritonitis in a mice model—both phagocytic and intracellular bactericidal capacities of neutrophils were enhanced by extract treatment (Thatte et al., 1992). The reduction of bacterial colonization in mice model was also observed after *O. tenuiflorum* treatment (Saími et al., 2009).

*Ipomoea batatas* (L.) was identified as an extract capable of stimulating phagolysosome formation inside of neutrophils (Miyazaki et al., 2005).

Inside of newly formed phagolysosomes, ROS are produced more intensively (Nathan, 2006; Urban et al., 2006). One of the major functions of phagocytes is to produce ROS during the oxidative burst (Brungs et al., 2015). Thus, stimulating this process may have beneficial effects during bacterial infections. *Echium amoenum* Fisch (Asadollahi et al., 2015), *Grewia asiatica* L. (Mesaik et al., 2013), *Saccharum officinarum* L. (El-Abasy et al., 2002) and *Acer pictum* subsp. mono (Maxim.) stimulated ROS production and enhanced phagocytosis efficiency. These plants have documented antioxidant and immunomodulatory properties (Abed et al., 2012; Akram et al., 2014; Mehmoood et al., 2020).

The signaling pathway leading to ROS release is highly dependent on spleen tyrosine kinase (Syk) activation (Brungs et al., 2015). Three plant extracts (*Cuscuta epithymum* (L.), *Ipomoea batatas* (L.) and *Euphorbia hirta* L.) stimulated Syk phosphorylation resulting in a more pronounced ROS production and an increased phagocytic activity (Sudam et al., 2017).

Besides the plant extracts that stimulated phagocytosis, we also presented plant materials that inhibited this process or had no effect on it (Table 1). As mentioned above, 18 plant extracts decreased phagocytic activity of neutrophils and 2 extracts did not affect this process. Only few molecular mechanisms responsible for suppression of phagocytosis were identified. They were dependent on the inhibition of MAC-1 cell surface expression (*Phyllanthus amarus*, *Camellia sinensis* (L.)) (Ilangkovan et al., 2015; Albuquerque et al., 2016), suppression of CR1, CR3, CR4, and Fcy receptors expression (*Areca catechu* L.) (Hung et al., 2006) and inhibition of ROS production (*Cucumis maderaspatanus* L, *Ajuga reptans* L, *Ajuga genevensis* L. Osbeckia octandra DC, *Cedrela fissilis* Vell, *Trichilia glabra* L, *Phyllanthus debilis*, *Crataegus laevigata* (Poir.), *Ixoera coccinea* L, *Zingiber zerumbet* (L.)) (t Hart et al., 1990; Benencia et al., 1995; Benencia et al., 2000; Dalli et al., 2008; Wickramasinghe et al., 2014; Akhtar et al., 2019; Toiú et al., 2019).

Two plant extracts: *Punica granatum* L. (Oliveira et al., 2010) and *Melia azedarach* L. (Benencia et al., 1994) did not affect phagocytosis. *P. granatum*, is known as a medical plant used for hypertension, diabetes, as well as a few types of cancer, but not for bacteria-induced diseases (Ge et al., 2021). *M. azedarach* was also used for the cancer-related disease treatment (Ervina et al., 2020; Shrestha et al., 2021).

We identified a few 9) plant-derived compounds regulating the neutrophils’ phagocytosis (Table 2). Seven compounds stimulated this process (gingerol, aracoline, 11-hydroxymustakone, N-methyl-2-pyridolidone, N-formyllonannain, magnoflorine, and tinocordiside) (Hung et al., 2006; Sharma et al., 2012; Jin et al., 2016) and two did not affect phagocytosis (cordifolioside A, syringin) (Sharma et al., 2012). Gingerol was isolated from *Pinellia pedatisecta* Schott, aracoline from *Areca catechu* L. and the rest of the selected compounds were isolated from *Tinospora cordifolia* (Willd.). *P. pedatisecta* as well as *A. catechu* are well-known in Chinese medicine as anti-inflammatory agents (Li et al., 2017; Wang et al., 2019). *T. cordifolia* is also known in Asian traditional medicine as an anti-inflammatory agent (Yates et al., 2021).

**PHARMACOLOGICAL MODULATION OF PHAGOCYTOSIS BY SELECTED DRUGS**

Some papers reported the effects of synthetic drugs on phagocytic function of neutrophils, but the data are scarce (Table 3). Numerous macrolide antibiotics stimulate phagocytosis performed by macrophages (Hodge et al., 2006; Xu et al., 2019), at least for some of the compounds using phosphatidylserine receptor-dependent pathway (Yamaryo et al., 2003). These results encouraged researchers to test influence of erythromycin (Cuffini et al., 2009) and azithromycin (Pohl et al., 2020) on neutrophil’s phagocytosis, which revealed that the former indeed stimulates phagocytosis, whereas the latter does not affect it. Fosfomycin was shown to enhance intracellular killing of *S. aureus* in both macrophages and neutrophils (Shen et al., 2016). Among antibiotics, chloramphenicol was also tested showing no significant results (Bystrzycka et al., 2017). Clinical significance of those findings remains uncertain, nevertheless testing the influence of other antibiotics, especially macrolides, on phagocytic function should be encouraged.

The interaction between melatonin and immune system is being widely explored, as discussed previously (Carrillo-Vico et al., 2013). Among others, its influence on neutrophil-mediated phagocytosis was tested, showing its inhibitory role in this phenomenon (Xu et al., 2019). The data on beneficial role of melatonin against polymicrobial sepsis also exist (Xu et al., 2019), thus further studies on the exact role of melatonin are needed.

**PHARMACOLOGICAL MODULATION OF DEGRANULATION VIA MEDICAL PLANTS**

We have identified 14 plant extracts, whose impact on degranulation was tested (Table 4). All the extracts inhibited
the degranulation, but only in 5 cases the mechanism of action was explained. Decreased degranulation resulted from inhibiting the elastase and MPO synthesis and release. An elastase-inhibitory effect was observed after Typhonium roxburghii treatment (Korinek et al., 2016), as well as Panax notoginseng (Jin et al., 2007), Lobelia chinensis Lour (Kuo et al., 2011) and Tamarindus indica (Paula et al., 2009). Selected plants were widely used to treat cancer and inflammatory-related diseases (Jin et al., 2007; Kuo et al., 2011; Korinek et al., 2016).

Anti-MPO activity was observed for Anvillea garcinii subsp. radiate (Boukemara et al., 2016). This plant was used for symptomatic treatment of cold (Perveen et al., 2020), but the successful treatment was related to the presence of flavonoids and sesquiterpene lactones, which show significant antibacterial and antifungal properties (Perveen et al., 2020). Thus, based on traditional use of A. garcinii, further evaluation of phagocytosis seems to be worth exploring.

For the rest of tested extracts (Cnidium monnieri (L.), Angelica gigas Nakai, Carthamus tinctorius L. Cinnamomum cassia (L.), Guarea kunthiana A. Juss. Prunus persica (L.), Rehmannia glutinosa (Gaertn.), Eugenia aurata, and Eugenia paniculata) authors documented only final effect on degranulation without identifying the molecular basis of the observed effect (Park et al., 2004; Costa et al., 2016; Jerjomiceva et al., 2016).

Chung and colleagues identified Hypericum Hemsli as a source of compounds modulating degranulation (Table 5). Geumchalcone A and gemichalcone B. isolated from H. germinal and Hypericum Hemsli decreased the degranulation of neutrophils (Chung et al., 2002). Plants belonging to the Hypericum species were used as diuretics, analgetics, antiphlogistics and antidepressant agents (Zhang et al., 2020a). Immunomodulatory, anti-inflammatory or antimicrobial properties of Hypericum species are unknown. Another example of a plant-derived compound decreasing degranulation of neutrophils is quercetin (Kanashiro et al., 2007; Pečivová et al., 2012). Although quercetin is widespread in many plant species, in both mentioned articles quercetin was used as a synthesized compound. Quercetin is a plant molecule that has shown many pharmacological activities, such as antimicrobial, antiviral, anticancer, having potential for treating metabolic, allergic, and inflammatory disorders, cardiovascular diseases, and arthritis (Batiha et al., 2020). The documented pharmacological effects encouraged to introduction of quercetin to the pharmaceutical market, however, low solubility in water, which is a key factor in drug absorption and its bioavailability, limits its use.

### Table 2 | List of selected compounds isolated from plant material with documented impact on phagocytosis.

| Plant family | Species | Compound | Cellular target and mechanism of action | Effect on phagocytosis | References |
|--------------|---------|----------|----------------------------------------|------------------------|------------|
| Araceae      | Pinellia pedatisecta schott | Gingerol | n.d | Increased phagocytic activity | Jin et al. (2016) |
| Arecales     | Areca catechu L | Arecolone | Increased uptake of bacteria | Increased phagocytic activity | Hung et al. (2006) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | 11-hydroxymustakone | Increased oxidative burst and ROS production | Increased phagocytic activity | Sharma et al. (2012) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | N-methyl-2-pyridolone | Increased oxidative burst and ROS production | Increased phagocytic activity | Sharma et al. (2012) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | N-formylmannosan | Increased oxidative burst and ROS production | Increased phagocytic activity | Sharma et al. (2012) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | Cordifoloside A | n.d | No influence on phagocytosis | Sharma et al. (2012) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | Magnoflorine | Increased oxidative burst and ROS production | Increased phagocytic activity | Sharma et al. (2012) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | Tinocordiside | Increased oxidative burst and ROS production | Increased phagocytic activity | Sharma et al. (2012) |
| Menispermaceae | Tinospora cordifolia (wild.) Hook.f. and thomson | Syringin | n.d | No influence on phagocytosis | Sharma et al. (2012) |

n.d. - not defined.

### Table 3 | List of selected synthetic drugs with documented impact on phagocytosis.

| Synthetic drug/s | Cellular target and mechanism of action | Effect on phagocytosis | References |
|------------------|----------------------------------------|------------------------|------------|
| Azithromycin     | n.d | No influence on phagocytosis | Pohl et al. (2020) |
| Chloramphenicol  | n.d | No influence on phagocytosis | Bystrzycka et al. (2017) |
| Erythromycin     | n.d | Increased phagocytic activity | Cuffini et al. (2009) |
| Fosfomycin       | Increased ROS production and strengthened oxidative burst process | Increased phagocytic activity | Shin et al. (2016) |
| Melatonin        | n.d | Decreased phagocytic activity | Xu et al. (2019) |

n.d. - not defined.
Further attempts to increase the quercetin solubility are urgently needed. Interestingly, quercetin, as well as described above gemichalcone A and gemichalcone B belong to flavonoids (Panche et al., 2016), which may indicate a specific chemical group of plant-derived compounds that modulate neutrophil degranulation.

Another source of compounds modulating degranulation is Salvia officinalis – it contains salvianolic acid capable to stimulate degranulation (Wang et al., 2010). Although mentioned data refer to mast cells the observed pharmacological effect is noteworthy, and in our opinion further studies on neutrophil model may also bring promising results. S. officinalis was used for the treatment of inflammation, gout, paralysis, ulcers, rheumatoid arthritis, dizziness, tremor, diarrhea, and hyperglycemia (Ghorbani and Esmaeilizadeh, 2017). Salvianolic acid is considered a chemo-preventive agent which suppresses oxidative stress and apoptosis (Wang et al., 2021). It was also documented that salvianolic acid may also be an antioxidant and anti-inflammatory agent which modulates PI3K/Akt/mTOR signaling pathway (Jin et al., 2021).

### PHARMACOLOGICAL MODULATION OF DEGRANULATION VIA DRUGS

Number of synthetic drugs known to regulate neutrophils degranulation is even lower than that affecting phagocytosis.

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**TABLE 4 | List of selected plant material with documented impact on degranulation of neutrophils.**

| Plant family | Species | Plant material | Cellular target and mechanism of action | Effect on degranulation | References |
|--------------|---------|----------------|----------------------------------------|-------------------------|------------|
| 1            | Apiaceae| Cnidium monnieri (L.) cusson | Plant extract | n.d | Decreased degranulation | Park et al. (2004) |
| 2            | Apiaceae| Angelica gigas nakai | Plant extract | n.d | Decreased degranulation | Park et al. (2004) |
| 3            | Araceae | Typhonium roxburghii schott | Plant extract | Inhibition of elastase release | Decreased degranulation | Korinek et al. (2016) |
| 4            | Aralilaceae | Panax notoginseng (burkil) F.H.Chen | Plant extract | Inhibition of elastase release | Decreased degranulation | Jin et al. (2007) |
| 5            | Asteraceae | Carthamus tinctorius L | Plant extract | n.d | Decreased degranulation | Park et al. (2004) |
| 6            | Asteraceae | Anvillea gacnica subsp. Radiata (coss. and durieui) | Plant extract | Inhibition of myeloperoxidase (MPO) release | Decreased degranulation | Boukemara et al. (2016) |
| 7            | Campanulaceae | Lobelia chinensis lour | Plant extract | Inhibition of elastase release | Decreased degranulation | Kuo et al. (2011) |
| 8            | Fabaceae | Tamarindus indica L | Plant extract | Inhibition of elastase release | Decreased degranulation | Paula et al. (2009) |
| 9            | Lauraceae | Cinnamomum cassia (L.) J.Presl | Plant extract | n.d | Decreased degranulation | Park et al. (2004) |
| 10           | Meliaceae | Guarea kunthiana A.Juss | Plant extract | n.d | Decreased degranulation | Jerjomiceva et al. (2016) |
| 11           | Myrtaceae | Eugenia aurata | Plant extract | n.d | Decreased degranulation | Costa et al. (2016) |
| 12           | Myrtaceae | Eugenia punicifolia | Plant extract | n.d | Decreased degranulation | Costa et al. (2016) |
| 13           | Orobanchaceae | Rehmannia glutinosa (gaertn.) DC. | Plant extract | n.d | Decreased degranulation | Park et al. (2004) |
| 14           | Rosaceae | Prunus persica (L.) batsch | Plant extract | n.d | Decreased degranulation | Park et al. (2004) |

n.d. - not defined.

**TABLE 5 | List of selected compounds isolated from plant material with documented impact on degranulation of neutrophils.**

| Plant family | Species | Compound | Cellular target and mechanism of action | Effect on degranulation | References |
|--------------|---------|----------|----------------------------------------|-------------------------|------------|
| 1            | Hypericaceae | Hypericum guminforum hemsl | Gemichalcone A | n.d | Decreased degranulation | Chung et al. (2002) |
| 2            | Hypericaceae | Hypericum guminforum hemsl | Gemichalcone B | n.d | Decreased degranulation | Chung et al. (2002) |
| 3            | Lamiaceae | Salvia officinalis L | Salvianolic acid | n.d | Increased degranulation | Wang et al. (2010) |

n.d. - not defined.
Similarly, two macrolide antibiotics were tested, and it was shown that erythromycin stimulates degranulation (Abdelghaffar et al., 1997), whereas azithromycin does not affect this process (Pohl et al., 2020). Chloramphenicol was also investigated, but exerted no effect on this function (Bystrzycka et al., 2017). Again, whether these effects are of clinical significance, remains unclear.

**PHARMACOLOGICAL MODULATION OF NETOSIS VIA MEDICAL PLANTS**

A few plant extracts were tested for their influence on NETosis (Table 7), showing that extracts obtained from *Anacardium occidentale* L. (Hollands et al., 2016), *Verbesina erstediana* Benth (Yaseen et al., 2017), and *Guarea kunthiana* A. Juss (Jerjomiceva et al., 2016) increased it, whereas *Salvia*
miltiorrhiza Bunge (Tao et al., 2018), Eugenia aurata (Costa et al., 2016), and Eugenia punicifolia (Costa et al., 2016) decreased it. A. occidentale has shown significant antibacterial activity and might have potential applications as new antibacterial drug (Shabeeba and Rathinasamy, 2018). A. occidentale extract has been tested in mice and rat models as a therapy of gastric-related conditions (Konan and Bacchi, 2007; Morais et al., 2010; Goulart da Silva et al., 2021). Thus, further evaluation of its immunomodulatory properties may open novel prospects.

V.verstediana and G. kunthiana were used in the folk medicine in treatment of bacterial infections, especially S. aureus (Jerjomiceva et al., 2016; Yaseen et al., 2017). Thus, documented antimicrobial properties justify the traditional use.

It is unknown why E. aurata and E. punicifolia extracts decreased NET release. For S. miltiorrhiza it has been speculated that decreased NET formation may results from the inhibition of MPO and NADPH oxidase (NOX) activity (Tao et al., 2018).

Six plant-derived compounds were investigated (Table 8). Anacardic acid, as the only substance, was able to increase NET production (Hollands et al., 2016), whereas catechin hydrate, epicatechin, rutin trihydrate (Kirchner et al., 2013), celastrol (Yu et al., 2015), and polydatin (Liao et al., 2018) decreased it. Anacardic acid belongs to phenolic lipids (Hollands et al., 2016), a diversified group of compounds derived from mono and dihydroxyphenols (Stasiuk and Kozubek, 2010). Anacardic acid shows antioxidant capacity, what may be related with suppression of superoxide generation (Trevisan et al., 2006). According to Hollands et al. inhibition of ROS production (inter alia superoxide generation) directly affects NET formation (Hollands et al., 2016).

Other compounds (catechin hydrate, epicatechin, rutin trihydrate, celastrol, and polydatin) classified as NET formation inhibitors belong to different chemical groups, thus an unambiguous assignment of a chemical structure to a biological function is not possible. Interestingly, excluding celastrol, compounds decreased NET formation via modulation of ROS production. Although

### Table 9: List of selected synthetic drugs with documented impact on NET formation.

| Synthetic drug/s | Cellular target and mechanism of action | Effect on NET formation | References |
|------------------|----------------------------------------|-------------------------|------------|
| 1 Amoxicillin    | n.d                                    | Increased NET formation | Bystrzycka et al. (2016) |
| 2 Amphotericin B | n.d                                    | Decreased NET formation | Decker et al. (2018) |
| 3 Azithromycin   | n.d                                    | Decreased NET formation | Bystrzycka et al. (2017) |
| 4 Cefotaxime     | n.d                                    | No effect on NET formation | Manda-Handzik et al. (2017) |
| 5 Clarithromycin | n.d                                    | Increased NET formation | Konstantinidis et al. (2016); Arampatziglou et al. (2018) |
| 6 Chloramphenicol| n.d                                    | Decreased NET formation | Bystrzycka et al. (2017) |
| 7 Clozapine      | n.d                                    | No effect on NET formation | Irizarry-Caro et al. (2018) |
| 8 Enrofloxacin   | Increased expression of PAD (peptidylarginine deminase) 4 protein, increased presence of citrullinated H3 | Increased NET formation | Jerjomiceva et al. (2014) |
| 9 Erythromycin   | n.d                                    | Decreased NET formation | Zhang et al. (2019) |
| 10 17-β-estradiol| n.d                                    | Increased NET formation | Yasuda et al. (2019) |
| 11 Gentamicin     | n.d                                    | Decreased NET formation | Manda-Handzik et al. (2017) |
| 12 Hydralazine   | n.d                                    | Increased NET formation | Irizarry-Caro et al. (2018) |
| 13 Mannitol hypertonic saline | Suppresses NOX2-dependent NETosis is via neutrophil dehydration | Decreased NET formation | Nadesalingam et al. (2018) |
| 14 Melatonin     | n.d                                    | Increased NET formation | Xu et al. (2019) |
| 15 Memantine     | Stimulates the production of MPO       | Increased NET formation | Peng et al. (2020) |
| 16 Metformin     | Reduction of elastase, proteinase-3 and histones concentration | Decreased NET formation | Menegazzo et al. (2018) |
| 17 Minocycline   | n.d                                    | No effect on NET formation | Irizarry-Caro et al. (2018) |
| 18 Procainamide  | n.d                                    | Increased NET formation | Irizarry-Caro et al. (2018) |
| 19 Voriconazole  | n.d                                    | Decreased NET formation | Decker et al. (2018) |

n.d.- not de

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anacardic acid also decreased ROS production, the biological effect was opposite to the one observed for other compounds. Because ROS production is crucial for efficient pathogen killing and NET formation, the dependency between chemical structure and pharmacological effect, as well as potential cytotoxicity need to be evaluated in the future.

PHARMACOLOGICAL MODULATION OF NET FORMATION VIA DRUGS

We identified 19 drugs, whose impact on NET formation was tested, mostly antibiotics and antimycotics (Table 9). β-lactams belong to the most widely used class of antibiotics (Bush and Bradford, 2016), but only two drugs from this group were tested. Amoxicillin increased NET formation (Bystrzycka et al., 2016), whereas no such effect was observed for cefotaxime (Manda-Handzlik et al., 2017). Three representatives of macrolides were investigated with different results. Clarithromycin increased NETosis (Konstantinidis et al., 2016; Arampatzigouli et al., 2018), erythromycin decreased it (Zhang et al., 2019), whereas azithromycin exerted no effect (Manda-Handzlik et al., 2017). Among other groups single particles were tested. Enrofloxacin (fluroquinolone used in veterinary medicine (López-Cadenas et al., 2013)) increased NET formation (Jerjomiceva et al., 2014), chloramphenicol (Bystrzycka et al., 2017) and gentamicin (Manda-Handzlik et al., 2017) decreased it, whereas minocycline (tetracycline used in acne vulgaris, but with some safety issues, as revised in (Garner et al., 2012)) did not affect this effect (Irizarry-Caro et al., 2018).

Neutrophils play an essential role in counteracting mycoses, including NET release, as recently revised (Urban and Nett, 2019). Both amphotericin B, usually used in severe cases (Hamill, 2013), and voriconazole, a drug of choice in invasive aspergillosis (Malani et al., 2015), decreased NET formation (Decker et al., 2018).

Above-mentioned data clearly reveal the complexity of the issue. It obvious that stimulation of NET production during an infection is beneficial, as it increases pathogen clearance and thus infection control, on the other hand it can participate in host tissue’s damage, as it is observed in acute respiratory distress syndrome (Grégoire et al., 2018), or even promote bacteria proliferation (Czaikoski et al., 2016). The effects of antimicrobial drugs are not easy to interpret due to difficulties in separating direct antimicrobial and immune-modulating properties. To further investigate this matter, in vivo studies on combinations of antibiotics or antimycotics with NET-modulating particles are needed. The substances discussed below can be used as such particles.

It was demonstrated that acidification inhibits ROS-dependent NET release (Behnen et al., 2017), whereas more alkaline pH stimulates NET formation mediated by both NOX2-independent (Naffah de Souza et al., 2017) and NOX2-dependent (Khan et al., 2018) pathways. These results are of interest since hypoperfusion-induced lactic acidemia is common in sepsis and septic shock and should be managed accordingly (Rhodes et al., 2017). It is worth verifying whether increasing pH affects NET release during the treatment.

Mannitol and hypertonic saline, agents administered to patients with intracranial edema (Burgess et al., 2016) may suppress of NOX2-dependent NET production (Nadesalingam et al., 2018). Again, the clinical significance of this observation is not clear.

Memantine is a drug approved for Alzheimer’s disease and tested for other applications (Lu and Nasrallah, 2018). It was shown that through stimulation of MPO production, it increases NET formation (Peng et al., 2020). Clozapine, the first atypical antipsychotic drug, whose side effects involve agranulocytosis (Khokhar et al., 2018) was also tested and was shown to exert no influence on NET formation (Irizarry-Caro et al., 2018).

Drug-induced lupus is a condition similar to systemic lupus erythematosus and triggered by initiation of a certain drug, usually resolving after the medication’s withdrawal (Borchers et al., 2007). Carmona-Rivera et al. demonstrated a role of NETosis in this pathology (Carmona-Rivera et al., 2017). In this context, an interesting approach was proposed by (Irizarry-Caro et al., 2018), who focused on drugs related to drug-induced lupus. They investigated hydralazin, a direct-acting vasodilator (Kandler et al., 2011) and procainamide, and showed that both drugs indeed increase NET production, which may contribute to autoimmunity.

CONCLUSION AND PROSPECTS

The Chinese and European Pharmacopeia convincingly report possible modulations of the immune response via medical plants. So far, natural products classified as immunomodulators were cursorially explored, without comprehensive identification of molecular mechanisms responsible for the observed effects. Firstly, there is a need to perform detailed phytochemical analyses of plant extracts, then select pure compounds for further biological evaluation. A specific assessment of purified compounds could help to identify cellular targets, determine bioactive dosage, biodistribution and kinetics. The knowledge of molecular mechanisms regulated by plant-derived compounds may potentially help to identify therapeutic targets, as well as potentially limit the spread of an infection. Moreover, the chemical structures of selected secondary metabolites can
serve as lead structures for synthesis of new substances. Similar conclusions could be drawn for synthetic drugs. Only few synthetic drugs were tested for their influence on immunomodulatory neutrophil functions and most of the data brought interesting results, thus further studies are urgently needed. For synthetic drugs, two areas seem to be the most important: identifying new substances, which affect effector functions of neutrophils, and testing approved drugs, especially antibiotics, antimycotics and those responsible for autoimmune reactions, for their impact on immunomodulatory functions of neutrophils.

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