

## REVIEW ARTICLE

# The Contribution of Formyl Peptide Receptor Dysfunction to the Course of Neuroinflammation: A Potential Role in the Brain Pathology

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**Abstract:** Chronic inflammatory processes within the central nervous system (CNS) are in part responsible for the development of neurodegenerative and psychiatric diseases. These processes are associated with, among other things, the increased and disturbed activation of microglia and the elevated production of proinflammatory factors. Recent studies indicated that the disruption of the process of resolution of inflammation (RoI) may be the cause of CNS disorders. It is shown that the RoI is regulated by endogenous molecules called specialized pro-resolving mediators (SPMs), which interact with specific membrane receptors. Some SPMs activate formyl peptide receptors (FPRs), which belong to the family of seven-transmembrane G protein-coupled receptors. These receptors take part not only in the proinflammatory response but also in the resolution of the inflammation process. Therefore, the activation of FPRs might have complex consequences.

This review discusses the potential role of FPRs, and in particular the role of FPR2 subtype, in the brain under physiological and pathological conditions and their involvement in processes underlying neurodegenerative and psychiatric disorders as well as ischemia, the pathogenesis of which involves the dysfunction of inflammatory processes.

**Keywords:** Neuroinflammation, glial cells, resolution of inflammation, formyl peptide receptors, new pro-resolving agonists, Alzheimer's disease, depression, ischemia.

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## ARTICLE HISTORY

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Received: June 26, 2019  
Revised: August 01, 2019  
Accepted: October 15, 2019

DOI:  
10.2174/1570159X17666191019170244

## 1. INTRODUCTION

Among the many hypotheses that attempt to explain the causes of CNS diseases, the hypothesis regarding the dysfunction of the immune system is of significance. It is believed that inflammatory processes in the brain, called neuroinflammation, can lead to the development of neurodegenerative diseases (e.g., Alzheimer's and Parkinson's diseases), mental disorders (e.g., depression and schizophrenia) and, to some extent, ischemic diseases. In general, acute inflammation is considered to be a beneficial process, while the dysfunction of the RoI leads to the development of the chronic inflammatory process. For this reason, in recent years, mechanisms and potential strategies that promote RoI have become a focus of interest. Indeed, based on the available literature, it can be postulated that both inflammatory activation and deficits in the RoI in the brain are significant causes of the development of CNS diseases.

The RoI is mediated by endogenous mediators called SPMs through interactions with specific membrane receptors. In the present review, special attention was paid to the role of G-protein-coupled receptors (GPCRs), which are expressed on immunocompetent cells in the brain, particularly formyl peptide receptors, in the RoI process.

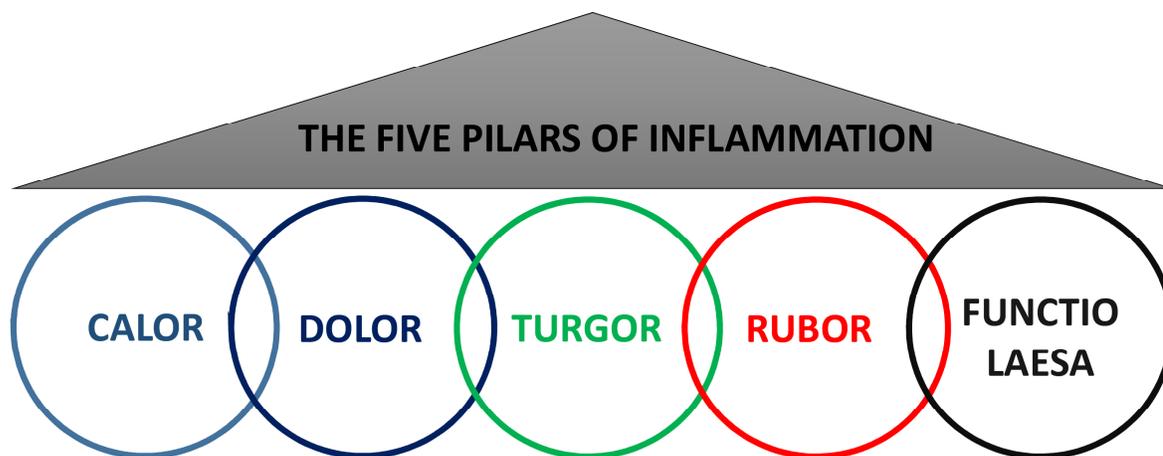
Moreover, new strategies supporting their role in shifting from proinflammatory to anti-inflammatory activation were proposed as a potential tool for the pharmacotherapy of CNS diseases triggered by chronic inflammation.

## 2. OVERVIEW OF THE INFLAMMATORY RESPONSE

Inflammation is a pathophysiological response of the body to tissue dysfunction or homeostatic imbalance triggered by a variety of harmful stimuli, including toxins, infections, trauma, stress or ischemia, that elicit the activation of the immune system. This process is characterized by the classic cardinal signs described by the Roman scholar and encyclopedist Aurelius Celsus (25 – 50 B.C.), namely, pain (*dolor*), increased temperature (*calor*), redness (*rubor*), and swelling (*tumor*) [1]. The fifth symptom of inflammation,

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**Fig. (1).** The classic cardinal signs of inflammation. (A higher resolution / colour version of this figure is available in the electronic copy of the article).

*i.e.*, loss of function and organ damage (*functio laesa*) was described later by Galen (129 - 200 A.D) [2] (Fig. 1).

The molecular and cellular events of the inflammatory response are known and result in increased blood flow, capillary dilatation, leukocyte infiltration, and production of chemical mediators. Generally, several phases of inflammation, including initiation, propagation, and resolution, have been described [3]. Recently, there have been very interesting reports indicating that the phases of inflammation do not develop sequentially but rather overlap; thus, both inflammatory activation and resolution can co-occur. Of course, the intensity of these processes is different, and this determines whether inflammation progresses or is abated. This novel interpretation of the RoI attempts to explain the dual role of some factors in the course of neuroinflammation; for example, it is still an open question whether cyclooxygenase 2 (COX-2) is an inhibitory target or a source of resolution molecules [4].

Acute inflammation is a rapid and self-limiting process. It usually lasts a few days and disappears after the removal of the cause without major damage to the body. When inflammation is properly controlled, it results in a protection against the spread of infection or damage and is followed by a resolution phase in which the affected tissues are restored to their original structural and functional state [5, 6]. Inflammation is mainly characterized by the presence of neutrophils, which quickly migrate to the site of injury or infection, promote the recruitment of inflammatory monocytes, and produce proinflammatory factors, thereby allowing for appropriate neutralization of the harmful factor [7]. The trafficking and homing of neutrophils are mediated mainly by families of GPCRs, one of which recognizes chemokines and the other of which recognizes classic chemoattractants originating from pathogens or damaged host tissues. Although neutrophils are essential for the proper elimination of the harmful factor, an excess influx of leukocytes can be more dangerous than the infection or injury itself; therefore, neutrophils undergo apoptosis after performing their action at the site of inflammation [8] and are subsequently eliminated by macrophages.

Briefly, the major signs of acute inflammation resolution include: the limitation or cessation of blood-borne cell influx, the counter-regulation of chemokines and cytokines, the switching of the signaling pathways associated with leukocyte survival, the induction of leukocyte apoptosis, the subsequent removal of leukocytes through efferocytosis by macrophages, the reprogramming of macrophages from proinflammatory to anti-inflammatory activated cells, the return of nonapoptotic cells to the vasculature or lymphatic system, and finally the initiation of healing processes. Resolution is a multistage and complex process [9, 10], and the failure of one or more steps may be involved in prolonged inflammation and the pathogenesis of chronic inflammatory diseases mainly due to constant stimulation of the immune system, the overproduction of proinflammatory cytokines [*e.g.*, interleukin (IL)-1 $\beta$ , IL-6, and (tumor necrosis factor (TNF)- $\alpha$ ], oxidative stress, the destruction of tissues at the site of the inflammatory process, and impairments in returning to homeostasis [11-14].

### 3. NEUROINFLAMMATION: THE IMMUNE RESPONSE IN THE BRAIN

Generally, CNS inflammation differs from inflammation that occurs in peripheral tissues because it is the result of the collective effects of various brain cells (microglia, astrocytes, oligodendrocytes, and NG2 glia) and in some cases peripheral immune cells. Moreover, data demonstrated that glial cells play a dual role in this process depending on the course of the disease and inflammatory environment. Interestingly, the double-edged nature of neuroinflammation, especially when the process is acute (transient), suggests that controlling neuroinflammation may have a neuroprotective function in maintaining homeostasis. On the other hand, it is commonly accepted that, while the mechanisms that ultimately lead to brain disturbances are different in various diseases, chronic neuroinflammation is a prominent feature of the progressive nature of neurodegeneration [14,15]. It appears to result from the complexity of neuroinflammation, which, as mentioned above, requires a coordinated response of glial cells and peripheral immune cells through the release of proinflammatory cytokines, including IL-1 and IL-6, as well as caspase activation [16].

### 3.1. The Role of Glial Cells in the Course of Neuroinflammation

In the brain, glial cells play a key role in the inflammatory response. They are the most abundant and widely distributed cells in the CNS, and they interact with neurons and immune cells as well as with blood vessels. They contribute to the monitoring of the physiological milieu and act as the first line of defense when the brain is exposed to different insults [17]. Glial cells participate in the activation, recognition, and modulation of immune reactions as well as the release of different factors (*e.g.*, cytokines and chemokines), which support immune defense and are crucial for the coordination of various immune cells [18].

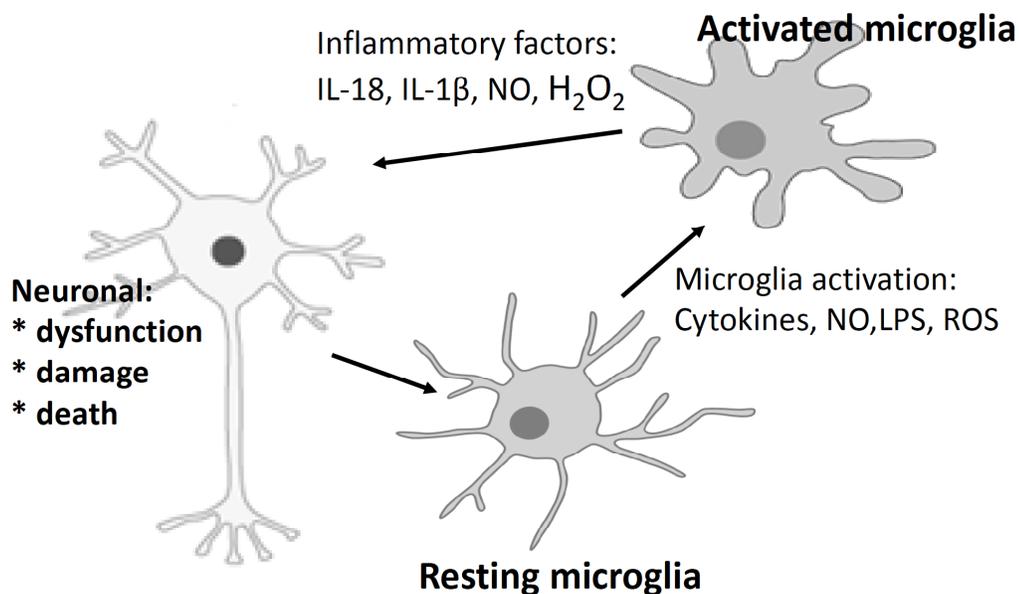
#### 3.1.1. Microglia

Microglia, small phagocytic cells of myeloid origin that comprise 10 – 20% of all cells, are the main immunocompetent cells in the brain. A vast body of data suggested that microglia have a dual character: neuroprotective and neurotoxic. In fact, under normal conditions, the activation of microglia cells plays a protective role by regulating the response to pathogens and promoting tissue repair through the release of anti-inflammatory and neurotrophic factors. Moreover, microglia participate in ontogenetic brain development and in maintaining homeostasis, including being involved in the programmed death of neurons during development, the removal of cellular debris and dying cells, and the regulation of synaptic plasticity [19]. Microglia also play a crucial role in neuron-glia interactions, thus controlling the proper functioning of neurons. In the healthy adult brain, microglial cells are highly dynamic in the resting state. At the same time, they express low levels of markers, including major histocompatibility complex (MHC) class I and II molecules. However, microglial processes are highly active, and once activated, microglia and CNS-infiltrating mono-

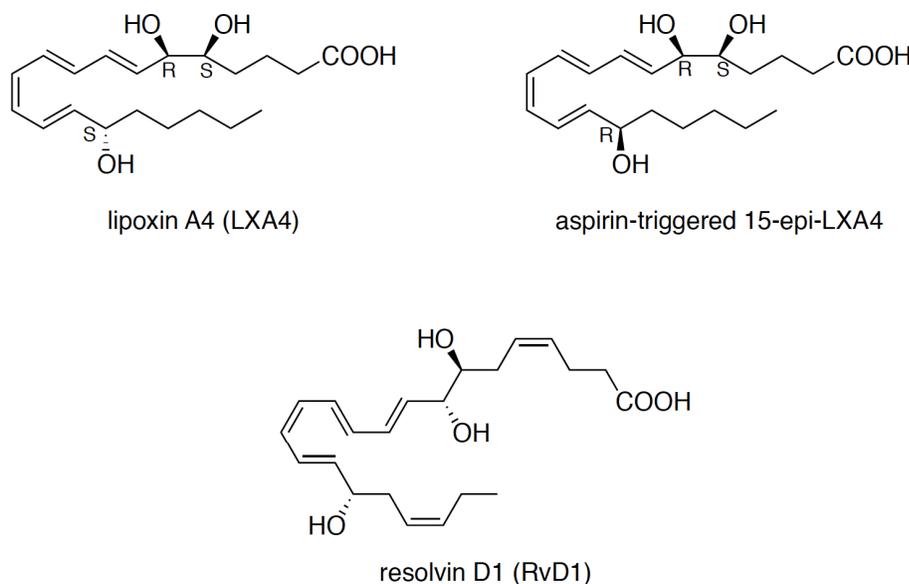
cyte-derived macrophages upregulate many surface molecules, including MHCII proteins and other costimulatory molecules, allowing them to act more efficiently than astrocytes as antigens for T cell presentation. Under pathological conditions, the chronic activation of immune responses can lead to the functional switch of microglia from a regulatory phenotype to a neurotoxic phenotype and the excessive release of proinflammatory cytokines, such as IL-1 $\beta$ , IL-18, TNF- $\alpha$ , and IL-6, chemokines and neurotoxic mediators, such as nitric oxide (NO), prostaglandin E2, superoxide anion, and excitatory amino acids [20]. Interestingly, the process of the acute and prolonged activation of microglia is phenotypically and functionally dynamic and may vary depending on the stage of disease or the status of the brain environment (Fig. 2).

#### 3.1.2. Astrocytes

Astrocytes, the most abundant subtype of glial cells in the CNS, are essential for brain homeostasis, as they provide metabolites and growth factors for neurons and to support synapse formation and plasticity. Furthermore, astrocytes are also able to detect harmful signals, promote cytokine and chemokine secretion, and activate the immune defense. Generally, astrocytes are exposed simultaneously to a plethora of stimuli and activate various intracellular signaling pathways in the course of neuroinflammation. Thus, astrocyte responses are not a reaction to a single event but instead the net result of a complex and diverse network of intracellular activators [21]. Importantly, the activation of astrocytes often follows the primary activation of microglia, of which the IL-1 family of cytokines is a key mediator, in response to pathological conditions such as trauma, stroke, or neurodegenerative disorders [22]. In fact, the release of IL-1 $\beta$ , mainly from microglia, can occur rapidly and may increase the secretion of other cytokines, such as IL-6, from astrocytes to promote inflammation. Moreover, IL-1 $\beta$  might hin-



**Fig. (2).** Microglia activated by stress, infection, and neurotoxins release many proinflammatory factors, such as IL-18, IL-1 $\beta$ , NO, and H<sub>2</sub>O<sub>2</sub>. These mediators, in turn, lead to neuronal dysfunction and, consequently, cell death. (A higher resolution / colour version of this figure is available in the electronic copy of the article).



**Fig. (3).** Structures of SPMs that bind at FPR2.

der the ability of astrocytes to reabsorb glutamic acid and promote the release of free radicals [23, 24]. Notably, IL-1Ra prevents astrocytes from causing pathological damage [25], showing that microglia might indirectly affect the function of astrocytes. In addition, microglial activation stimulates astrocytes to secrete IL-10 and TGF- $\beta$ 1 (transforming growth factor) [26], of which high levels initiate a feedback loop to reduce IL-1 $\beta$  release, thus inhibiting microglial activation and resulting in the RoI. Moreover, astrocytes may secrete growth substances, such as neural growth factor (NGF), cerebral growth factor (BGF), and basic fibroblast growth factor (bFGF), which have a significant role in the repair and growth of neurons as well as in brain development.

It should be mentioned that neuroinflammation is contingent upon the influx of immune cells from the peripheral blood to the brain. Experimental data have shown that a disturbance in the interaction between astrocytes and microglia plays a crucial role in blood-brain barrier (BBB) dysfunction [27]. The overexpression of inflammatory stimuli in the neurovascular unit may initiate a response that leads to the destruction of the BBB and to neuronal damage. In addition, reactive forms of microglia and astrocytes contribute to damage within the BBB, which becomes more permeable to peripheral leucocytes and activates glial cells to produce proinflammatory cytokines, chemokines, and reactive oxygen species (ROS) to become an additional source of inflammation-supporting molecules.

#### 4. THE ROLE OF PRO-RESOLVING MOLECULES IN THE RESOLUTION OF INFLAMMATION

Accumulating evidence has suggested that chronic neuroinflammation plays a relevant role in the onset and progression of neurodegenerative diseases as well as psychiatric disorders. Furthermore, the combined dysregulation of glial activation and proinflammatory cytokine production may also be an urgent driver in the pathogenesis of ischemia and traumatic brain disorders, as they are associated with neuro-

immunological abnormalities that contribute to neurodegeneration. To date, many potential neuroprotective agents, such as plant extracts, and strategies (*e.g.*, autophagy activation), have been established [16, 28]. Generally, the prevention of neuropathologies crucially depends on the termination of inflammation, which is traditionally considered a passive process through which inflammation spontaneously subsides. However, as mentioned before, there is a growing appreciation that, like the initiation of inflammation, the RoI is a complex, active process aimed at restoring tissue integrity and function [5, 29]. The progression of this process requires proper endogenous activation that induces a switch from the release of proinflammatory molecules to the secretion of pro-resolving mediators, which comprise a wide variety of compounds, including gases, proteins, and lipids [30]. The latter group is most often studied and is termed SPMs (Fig. 3).

SPMs are produced through the oxidation of arachidonic acid (AA), 3-eicosapentaenoic acid (EPA), and docosahexaenoic acid (DHA). The first class of SPMs to be described was termed lipoxins and originates from AA. E-series resolvins are derived from EPA, while D-series resolvins comprise resolvins, maresins, and protectins generated from DHA. In humans, the production of lipid SPMs is catalyzed by epithelial cell-, eosinophil-, and monocyte-derived 15-lipoxygenase (15-LOX), leukocyte-derived 5-LOX, platelet-derived 12-LOX, and COX-2. These enzymes have been proven to be present in various cell types and to generate precursors of aspirin-triggered lipoxins, resolvins, and protectins in the acetylated form. SPMs exert their biological effects *via* G-protein-coupled receptors, including the following:

- Formyl peptide receptor 2 (also called lipoxin A4 receptor, FPR2/ALX) for lipoxin A4 (LXA4), annexin A1 (AnxA1), and resolvin D1 (RvD1).
- G-protein-coupled receptor 32 (GPR32) for lipoxin A4 and RvD1.

In addition, RvD1s act through chemokine-like receptor 1 (ChemR23), and RvE1s act through the leukotriene B4 receptor (LTB4R or BLT1) [29, 31, 32].

The main advantage of the biological activity of SPMs is that they do not block signal transduction within the inflammatory cascade; instead, they trigger processes that reduce the expression of proinflammatory molecules. Moreover, SPMs activate cascades that induce remodeling within sites damaged by inflammatory processes. Importantly, anti-inflammatory effects rely mostly on an inhibitory/blocking action, while pro-resolving effects are mediated by the activation and stimulation of specific inherent processes (e.g., apoptosis and efferocytosis); however, the RoI is the final result of both [33-35]. In addition, Peretti *et al.* recently noted that SPMs elicit “mild to moderate effects”, which, by establishing an equilibrium between proinflammatory and anti-inflammatory reactions, help to strike a balance [35].

According to the literature, ideal SPM's, which combine both anti-inflammatory and pro-resolving activities, should fulfill the following fundamental criteria [3, 31, 36]: the cessation of inflammatory cell recruitment (e.g., tissue infiltration by neutrophils), the regulation of the secretion of cytokines and chemokines, the switching of macrophages from the classically activated phenotype to the alternatively activated phenotype, the induction of efferocytosis of apoptotic neutrophils (by macrophages), the removal of nonapoptotic cells, dendritic cells and macrophages from the site of inflammation, the modulation of the immune response, the instruction of suppressive immune cells and the adaptive immune response, the induction of tissue repair (the final step of resolution), and a return to homeostasis.

In the context of our article, lipoxins (LXs), which are metabolites of AA, are crucial. Briefly, LXA4 and lipoxin B4 (LXB4) are produced by the oxygenation of AA by 15-LOX and 5-LOX and subsequent enzymatic hydrolysis. On the other hand, new lipoxin analogs termed aspirin-triggered lipoxins (AT-LXA4) are derived from the aspirin-triggered pathway, in which aspirin alters COX-2 activity by increasing the acetylation of COX-2 [37]. The inactivation of lipoxin catalyzed by the enzyme 15-prostaglandin dehydrogenase can result in the synthesis of a series of other stable lipoxin analogs [38] that retain all of the biological functions of lipoxins. It is now widely recognized that the synthesis of lipoxins is enhanced during the inflammatory response, and their activation leads to the RoI through the promotion of phagocytosis of apoptotic neutrophils by macrophages, which potentiates the abatement of the inflammatory activation [39].

The concept that the biological action of lipoxins is mediated by GPCRs was developed in 1990 based on studies with pertussis toxin [40]. The first studies indicating a specific recognition site for LXA4 on polymorphonuclear leukocytes (PMNs) were conducted through the use of tritium-radiolabeled LXA4 in 1992 [41]. A research group led by Serhan identified cDNA for the human 7-transmembrane receptor (pINF114), which codes for functional high-affinity receptors for lipoxin A4 [42]. These receptors are termed FPR2/ALXR [43-48].

## 5. FORMYL PEPTIDE RECEPTOR FAMILY

Pattern recognition receptors (PRRs) located on immune cells are a type of receptor that plays a significant role in the mechanisms of innate immunity. These receptors recognize pathogen-associated molecular patterns (PAMPs) as important mechanisms for the elimination of various pathogens and their components by innate immunity [49]. *N*-formyl-methionyl-leucylphenylalanine (*f*MLP) is one of them, and it induces the chemotaxis of neutrophil granulocytes and monocytes [50, 51]; these effects are associated with interactions with formyl peptide receptors.

In humans, FPRs have been detected in the spinal cord, brain, anterior horn cells, hypoglossal nucleus neurons, choroid plexus, and epithelium [52]. FPRs have also been found in the cerebellar system, in the sensory system, on reticular activating system neurons, and in the ependyma. Hippocampal neurons, pyramidal cell neurons, end-plate pyramidal cells, astrocytes, Schwann cells of the peripheral nervous system, and cells in the parasympathetic system show a high level of FPR expression [53]. On the other hand, microglial cells isolated from normal adult humans express the gene for FPRs, but the level of the receptor protein is low, and no functional observation has been reported [54, 55]. In rodents, microglial cells lack a chemotactic response to *f*MLP [56], suggesting that these cells express a low level or no *f*MLP receptors on their surface. Cui and collaborators found that both murine primary microglial cells and the N9 cells constitutively express low levels of both the *Fpr2* and *Fpr1* genes in the resting state, and this expression is enhanced by treatment with TNF- $\alpha$ . However, resting murine microglial cells do not respond to *f*MLP or to other agonists of either FPR2 or FPR1. TNF- $\alpha$  treatment enables both primary and N9 cells to develop a potent chemotaxis response to agonists that act only on FPR2 in association with the surface expression of low-affinity *f*MLP binding sites [57].

### 5.1. Structural Characteristics of FPRs

FPRs belong to the largest and functionally diverse family of G-protein-coupled receptors. These receptors, irrespective of their function, have the same structural design. They are long single-chain proteins that span the plasma membrane seven times and are therefore often referred to as heptahelical (7-transmembrane) receptors (7-TMs) [58-60]. Their polypeptide chain is composed of the extracellular N-terminal domain, seven transmembrane helices (TM1-7) linked by intracellular and extracellular loops (IL1-3, EL1-3), and an intracellular C-terminus. In some receptors, an additional helix that is parallel to the inner surface of the plasma membrane has been detected [61, 62]. According to the GRAFS (glutamate, rhodopsin, adhesion, frizzled/taste2, and secretin) system recommended by the International Union of Basic and Clinical Pharmacology (IUPHAR), in which GPCRs are classified based on phylogenetic criteria, FPRs belong to the rhodopsin-like receptor family (class A) [63, 64]. Receptors of this family are characterized by relatively short N-terminal parts and transmembrane regions that participate in ligand recognition. To date, two very conserved and key motifs have been discovered in FPRs: the NPXXY motif in TM7, which is a part of a molecular switch

that induces a conformational change that activates the receptor, and the E/DRY motif, which is located at the border between TM3 and TM2. The E/DRY motif acts as an ionic blockade and links TM3 and TM6 [65].

## 5.2. Human Formyl Peptide Receptors

To date, three members belonging to the *FPR* family have been identified, namely, *FPR1*, *FPR2*, and *FPR3* [66-70]. *FPR1* and *FPR2* show high homology and common functions.

The human formyl peptide receptor 1 (*hFPR1*), which is an *fMLP* target, was isolated from HL-60 cells, which are highly responsive to treatment with N-formylated peptides. *hFPR2* is the result of gene amplification. However, evolutionarily, *hFPR3* is the youngest member of this gene family and was created by gene duplication in the lineage leading to primates [71].

All members of the human *FPR* gene family are located on chromosome 19. Pairs of paralogous receptors exhibit a basic structure highly similar to that of human *FPR*. Human *FPR2* has a 69% sequence homology with *hFPR1*. *FPR3* sequence analysis has demonstrated that the receptor is most strongly related to *FPR2*, and this discovery suggests that *FPR3* was created by *FPR2* gene duplication [71, 72]. A comparison of human *FPRs* with their counterparts in primates has revealed a high level of homology that reaches 95-99% [73]. This finding highlights the functional significance of this receptor family. Interestingly, the highest variability levels of the *FPR* sequence are detectable in the extracellular loops of the *FPR* sequence, which suggests different ligand binding affinities/preferences of particular receptors and species [73,74]. *FPR1* and *FPR2* are expressed both on monocytes and neutrophils, whereas *FPR3* expression is found only on monocytes. *FPR1* is also expressed on astrocytes, microglia, and immature dendritic cells. *FPR2* exhibits even greater expression both on cells in the CNS and in the periphery, including tumor cells [53, 52,76].

## 5.3. Mouse Formyl Peptide Receptors

Since the *FPR1* gene was cloned in humans, this gene has also been identified in other mammals including rabbits, guinea pigs, horses, rats, and mice [44]. In spite of the general sequence homology, the genes coding for the *FPR* family in humans and other species significantly differ in terms of the number and sequence [71]. For instance, the mouse *Fpr* gene family comprises 8 known members (*mFpr1*, *mFpr2*, *mFpr-rs1*, *mFpr-rs3*, *mFpr-rs4*, *mFpr-rs6*, *mFpr-rs7*, and *mFpr-rs8*) located on mouse chromosome 17A3.2. Studies on *mFprs* have mostly focused on *mFpr1* and *mFpr2* gene products, which are expressed on mouse phagocytic leukocytes and are highly similar to their human counterparts. Targeted *mFpr1* or *mFpr2* deletion makes mice more susceptible to bacterial infections without affected viability or fertility [77, 78]. Under unstimulated conditions, mice with *mFpr1* or *mFpr2* gene knockout do not show behavioral disturbances. However, an array of models of diseases has shown that the development of disturbances in *mFpr1*<sup>-/-</sup> and *mFpr2*<sup>-/-</sup> mice is induced, suggesting a regulatory role for these receptors in inflammation. Similar to human *FPR1*, *mFpr1* has been identified as a receptor with low affinity for

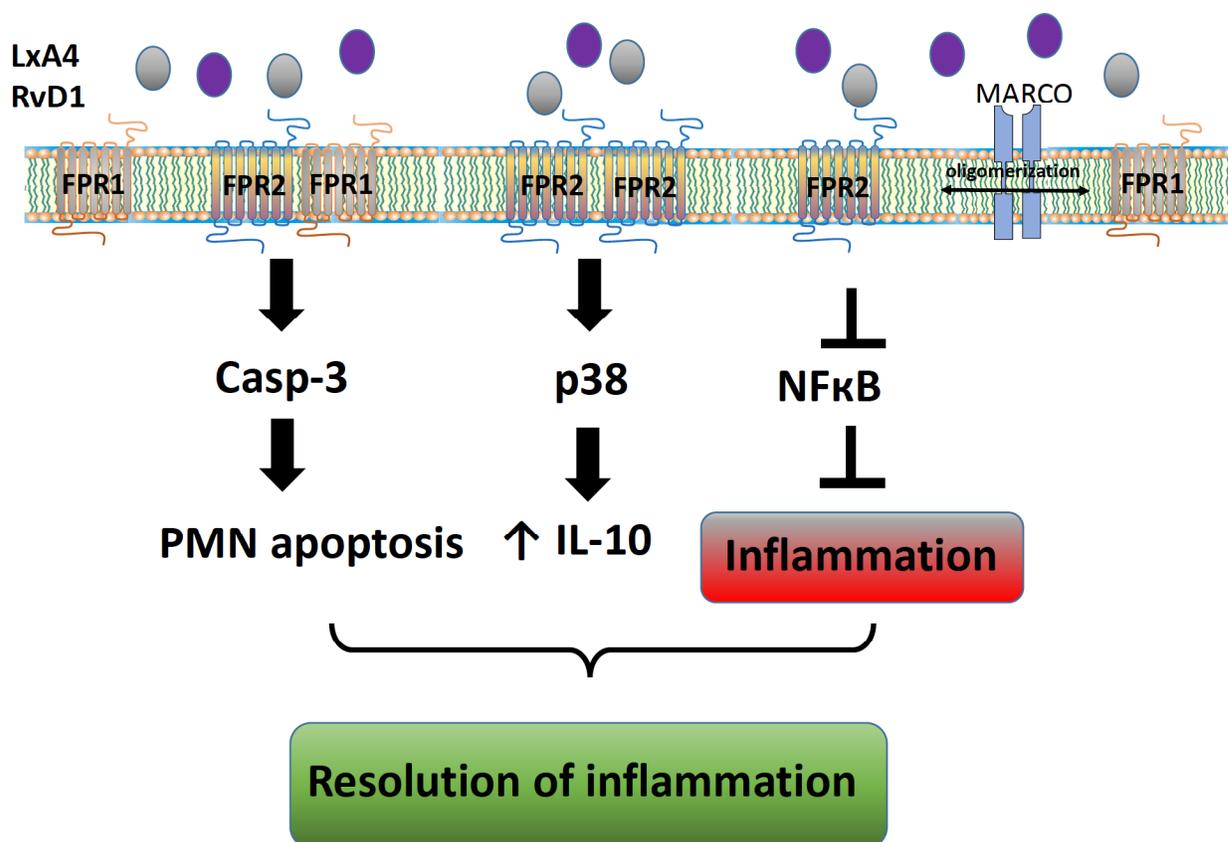
classical *fMLP* [79]. In general, *mFpr2* is not a high-affinity receptor for *fMLP* or other formyl peptides tested thus far [80]. However, it interacts with endogenous *FPR2* agonists, including the amyloidogenic proteins SAA (serum amyloid A) [81] and  $\beta$ -amyloid plaques ( $A\beta$ ) 42 [82]. In addition, *mFpr2* has also been described as a receptor for F2L [83], which is a strong agonist of human *FPR3* [84]. The gene products of five other *mFprs*, including *mFpr-rs1*, *mFpr-rs3*, *mFpr-rs4*, *mFpr-rs6*, and *mFpr-rs7*, have been described as mouse vomeronasal sensory receptors [79, 85]. The functions of *mFpr-rs1* are still unclear. Until recently, it was believed that *mFpr-rs1* has similar functions as those of human *FPR2* because it was reported that an *mFpr-rs1* variant encodes the mouse lipoxin A4 receptor [86]. However, functional and pharmacological tests on stably transfected cell lines showed that *mFpr-rs1* barely responds to most of the agonists of human and mouse *FPRs* [87]. These observations indicate that mouse *Fprs* (especially *mFpr1* and *mFpr2*) present many structural and pharmacological similarities to human *FPRs*.

## 5.4. Nomenclature of FPR Receptors

In the past, members of the *FPR* family were designated by different names (e.g., *FPR1*, *FPR2*, *FPRL1*, *FPRH1*, *FPR*-related receptor, *HM63*, *FMLP*-related receptor II for *FPR2*, *FPRL2*, *FPRH2*, and *RMLP*-related receptor I for *FPR3*). To standardize the terminology, the IUPHAR introduced a new nomenclature based on agonist-receptor interactions. Based on this system, members of the human *FPR* family are designated *FPR1*, *FPR2*, and *FPR3* [44]. However, in the literature, human *FPR2* is often termed *LXA4R*, *ALX*, or a combination of both of these terms (*FPR2/ALX*) due to its interaction with lipoxins A4 [42, 45, 47].

## 5.5. Conformational Changes Determine the Pro-resolving Activity of FPRs

A growing body of evidence has indicated that GPCRs are allosteric proteins that can adopt many conformations, some of which may promote continuous endocytosis in the absence of stimulation. More attention has been dedicated to processes that appear to be a crucial aspect of receptor regulation [88]. In many cases, *FPR2* is constitutively internalized via the  $\beta$ -arrestin-dependent pathway and undergoes clathrin-dependent constitutive internalization [89]. Phosphorylation and internalization seem to be independent events, suggesting that constitutive endocytosis may not be a consequence of the basal activity of the receptor. Moreover, *FPR2* is phosphorylated in an agonist-dependent manner, but the phosphorylation sites have not yet been identified [90]. Emerging literature has indicated that the formation of higher-order structures fulfills a vital role in the regulation of *FPR* family receptors. For instance, *FPR1* forms homodimers and interacts with both *FPR2* and *FPR3* [88]. It is worth mentioning that the pro-resolving ligand *AnxA1* and its N-terminal peptide *Ac2-26* potentiate the formation of *FPR1/FPR2* heterodimers and/or *FPR2* homodimers, while proinflammatory ligands and antagonists do not exert such action. It is thought that the stabilization of the *FPR2* oligomer may be significant for p38/MAPK/Hsp27 pathway activation [88, 91]. A conformational change in the *ALX/FPR2* receptor (ligand-dependent) determines its pro-



**Fig. (4).** Ligand-biased signaling *via* formyl peptide receptors. A variety of ligands (*e.g.*, lipoxin A4 (LXA4) and resolvin D1 (RvD1)) induce FPR2/ALX homodimerization and FPR2/FPR1 dimerization and lead to the activation of different downstream signaling cascades. Oligomerization is not limited to FPR family members and includes even non-GPCR surface receptors, such as the scavenger receptor MARCO. PMN: polymorphonuclear leukocyte. (*A higher resolution / colour version of this figure is available in the electronic copy of the article.*)

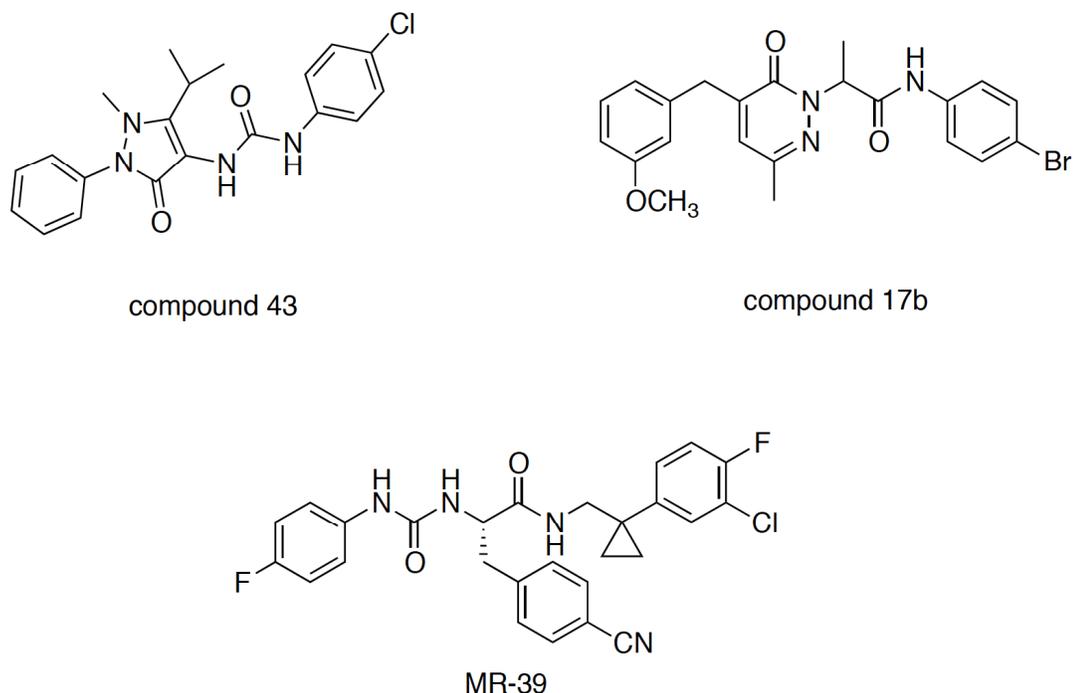
resolving action [88]. Moreover, the oligomerization of these receptors is not limited only to FPR family members but also involves surface receptors, such as the scavenger receptor MARCO (macrophage receptor with collagenous structure). Interactions between FPR1, FPR2, and MARCO have been demonstrated by bioluminescence and coimmunoprecipitation studies and are associated with agonist-induced changes in cyclic adenosine monophosphate (cAMP) levels and extracellular signal-regulated kinases (ERK1/2) phosphorylation. They can also play an important role in A $\beta$ 1-42-induced signal transduction in glial cells [92] (Fig. 4). Recently, in an excellent review, Raabe *et al.* discussed biased perspectives on FPRs. In the authors' opinion, the unique character of FPR2 relies on the ability of its ligands to selectively activate subsets of downstream signaling pathways coupled to the receptor and inhibit others. This interpretation provides an elegant explanation as to why different FPR2 agonists do not cause the same effects; this receptor is exceptional due to its ability to shift from a proinflammatory to anti-inflammatory response while maintaining the former at a low but possibly life-saving level [93].

#### 5.6. New FPR Agonists as a Promising Tool for the Pharmacotherapy of CNS Diseases

The vital and fascinating characteristics of FPRs result from their ability to interact with structurally diverse ligands

(lipids, proteins, and peptides), which can stimulate various cell type- and ligand-specific cellular responses downstream of receptor activation. This ability is caused by the biased agonism of FPRs rather than the existence of a variety of FPR subtypes. This feature reflects the ability of GPCRs to assume different conformational states, each linked to discrete cell effects. Therefore, the biased agonism provides an opportunity to potentially promote expected on-target signal transduction without on-target adverse outcomes. This paradigm has an impact on the current search for new drugs. In addition, the search of FPRs, finding new FPR ligands that are potentially more resistant to degradation than endogenous peptide ligands appears to be of special interest.

Several studies attempted to elucidate the molecular basis of this feature of FPRs. By using chimeric receptors, it has been demonstrated that distinct receptor domains might interact with distinct ligands, thus eliciting different downstream responses. In particular, lipoxin A4 has been shown to interact with the third extracellular loop of FPR2 to induce pro-resolving responses [94]. The interaction of AnxA1 with the N-terminal domain and the second extracellular loop is required for inducing Ca<sup>2+</sup> mobilization and the modulation of gene expression. On the other hand, the proinflammatory ligand SAA might activate Ca<sup>2+</sup> flux and ERK phosphorylation through its interaction with the first and second extracellular loop. Small-molecule FPR ligands, such as compound



**Fig. (5).** Structures of some “small-molecule” nonpeptidic FPR ligands.

43 (Fig. 5), which can penetrate into the transmembrane domains, have been proposed to interact with the first extracellular loop, the third helix, and the second intracellular loop [95]. In addition to naturally occurring peptides/proteins and endogenous lipids, various synthetic peptides, as well as nonpeptidic ligands belonging to different chemical classes, have been reported to date. The reader can refer to excellent reviews of FPR ligands reported in the literature [30, 96]. Here, we briefly discuss small-molecule nonpeptidic ligands that may be of interest for the development of future innovative therapies for chronic inflammatory conditions.

Qin *et al.* [97] studied, for the first time, the effect of the biased agonism of two small-molecule FPR agonists, namely, compound 43 and compound 17b, on the cardioprotective profile in myocardial infarction. It was demonstrated that compound 17b, unlike compound 43, induces a lower  $\text{Ca}^{2+}$  flux relative to that induced by ERK1/2-Akt signal transduction. Since increased intracellular  $\text{Ca}^{2+}$  contributes to cardiomyocyte damage and is a key contributor to the influx of inflammatory neutrophils and macrophages, the biased agonism of compound 17b provides superior outcomes in *in vivo* models of myocardial infarction [97]. This study clearly suggests that exploiting biased signaling can offer new possibilities in balancing proinflammatory and anti-inflammatory pathways to retain the homeostatic environment in complex inflammatory diseases, and this approach can also be applied for drug development for CNS diseases.

We contributed to the field by identifying a series of ureidopropanamide derivatives as FPR2 agonists. The compounds formally originate from the gastrin-releasing peptide receptor antagonist PD-175266, which has been found to be a potent FPR1/FPR2 agonist [48, 98, 99]. Through appropriate structural modifications, we ultimately identified MR39 as

an FPR2 agonist (Fig. 5). The *in vitro* pharmacokinetic properties of MR39 are favorable, as the compound is stable to oxidative metabolism in rat liver microsomes ( $t_{1/2} = 48$  min) and displays good passive permeability through a monolayer of hCMEC/D3 cells, immortalized human brain microvascular endothelial cells that are considered an *in vitro* model of the BBB. In addition, MR39 reduced IL-1 $\beta$  and TNF- $\alpha$  levels in LPS-stimulated rat primary microglial cell cultures and thus showed protective and anti-inflammatory properties [100]. Considering that LXA4 is rapidly inactivated *in vivo* [101] and that there is no direct evidence that LXA4 can pass the BBB, MR39 represents a prospective tool for studying the therapeutic potential of FPR2 agonists for the pharmacotherapy of CNS diseases (Fig. 5).

## 6. FORMYL PEPTIDE RECEPTORS AS A NEW TARGET FOR THERAPY FOR SOME BRAIN PATHOLOGIES

Despite many years of multicenter studies, the efficacy of therapeutic interventions for some CNS diseases remains unsatisfactory. This seems to be due to the complex and multifactorial nature of their pathological basis. In this review, we focused our attention on Alzheimer’s disease, depression, and ischemic disease because they show different symptomatology and different therapeutic requirements, although prolonged inflammation undoubtedly plays a key role in their etiopathogenesis. Moreover, the lack of appropriate pharmacological therapy for neuroinflammation is believed to hamper the efficacy of pharmacotherapy when it is available. Based on the available data, the multidimensional analysis was carried out to answer the question of whether supporting the endogenous RoI process through the regulation/modulation of FPR activity, including the use of pro-

resolving and anti-inflammatory ligands, may be a promising therapeutic option for treating these diseases.

### 6.1. RoI Deficits in Alzheimer's Disease

Dementia is a clinical syndrome that develops as a result of neurodegenerative processes in the brain. Alzheimer's disease (AD), the most common form of dementia worldwide, has been reported for more than 100 years thanks to the studies of the German neuropathologist and psychiatrist Alois Alzheimer [102]. According to the World Health Organization (WHO) estimates, at present, 50 million people suffer from AD worldwide, and 10 million new cases are recorded every year. It has been projected that the number of AD patients will reach 82 million in 2030 and 152 million in 2050 [103].

The major histopathological hallmarks of AD include  $\beta$ -amyloid ( $A\beta$ ) plaque formation, neurofibrillary tangles (NFTs), and progressive neuronal loss. However, a third core feature of AD was recently identified; *postmortem* studies of AD patient brains demonstrated an exaggerated inflammatory response [104-114]. It should be mentioned that the first studies in the 1980s revealed the presence of immune cells and their mediators in the vicinity of  $\beta$ -amyloid plaques [107] and demonstrated that medications used for diseases such as rheumatoid arthritis exhibit protective potential against AD. Moreover, the risk of developing AD is reduced to 50% in patients treated with nonsteroidal anti-inflammatory drugs (NSAIDs) in the long term [115-117]. Therefore, the results of these investigations support the hypothesis that chronic immune activation plays a key role, although indirectly, in the development of AD.

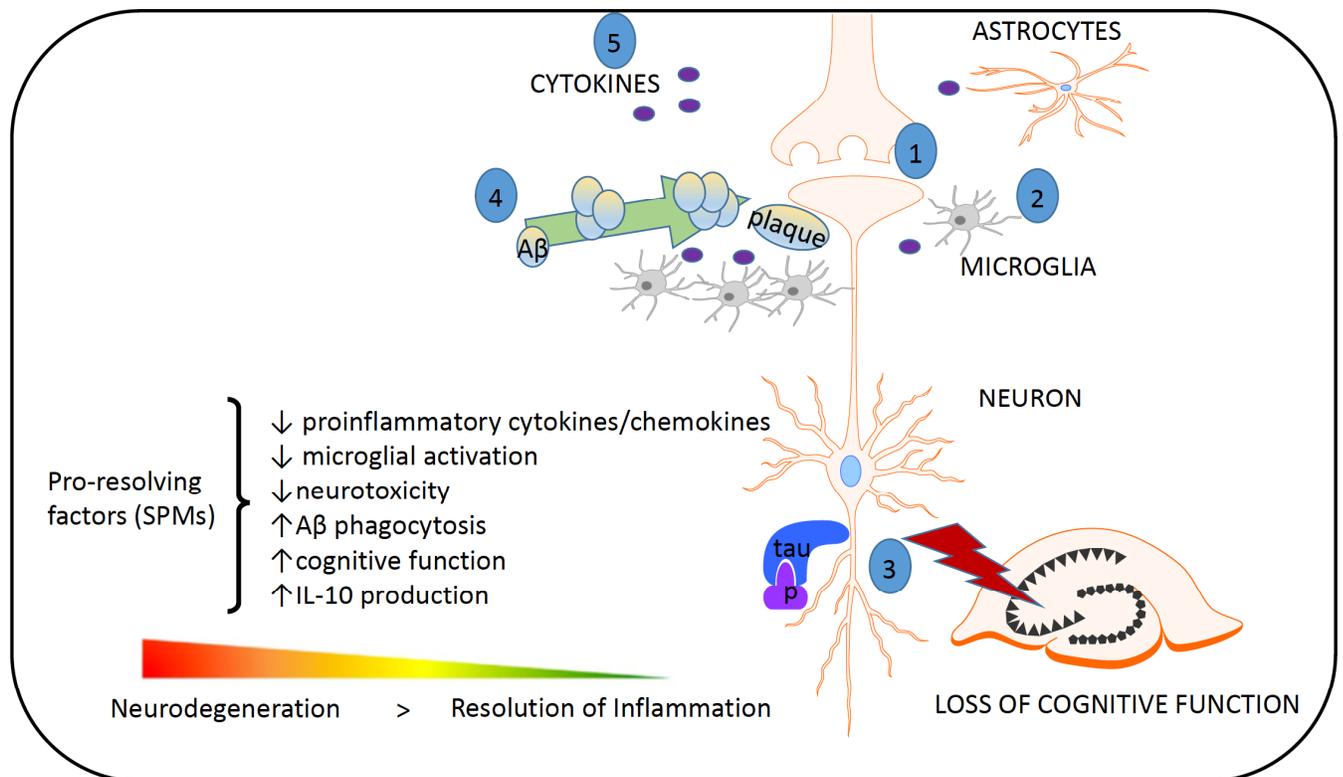
In fact, undisputed evidence of immune system activation in the brains of AD patients comes from *postmortem* studies of patients that suffered recent head trauma; the  $A\beta$  level and the IL-1 $\beta$  level are increased 1-3 weeks postinjury. These changes lead to the further elevation of the production of  $\beta$ -amyloid precursor [118, 119]. Moreover, it has been shown that increased IL-1 $\beta$  levels induce the synthesis of other cytokines, including IL-6, which stimulates the activation of cyclin-dependent kinase 5 (CDK5), a kinase associated with tau phosphorylation [120]. Since CDK5 is an autophagy-regulating kinase and its dysfunction is crucial in the development of neurodegenerative disorders, autophagy-targeted therapeutic approaches can be considered novel therapeutic strategies for AD treatment [28]. Advanced studies have indicated that neuroinflammation in AD patients is mostly related to microglial activation [121-123] and elevated levels of proinflammatory cytokines not only in the brain but also in the serum [124-126]. At the same time, *postmortem* studies documented a downregulation of anti-inflammatory factor expression in the brain tissue [127]. Interestingly, some data indicated that  $\beta$ -amyloid-induced neuronal death and synaptic impairment are in part mediated by astrocyte activation induced by numerous factors, including free saturated fatty acids, pathogens, and oxidative stress. As in the case of microglial activation, the activation of astrocytes leads to the release of proinflammatory cytokines as well as cyclooxygenase-2 and the receptor for advanced glycation end products/nuclear factor- $\kappa$ B (NF- $\kappa$ B) axis activation. These observations suggest a role for advanced glycation end products in

age-related cognitive changes as well as a potential beneficial role of nutraceuticals in the prevention of chronic neuroinflammation and AD-related pathology [128].

Notable hallmarks of AD indicate that prolonged neuroinflammation is a consequence of mitochondrial dysfunction. In fact, data demonstrated that, in AD, the activity of various mitochondria-localized enzymes is decreased and that glucose utilization in the brain is reduced, which may indirectly reflect a consequence of mitochondrial impairment and bioenergetic failure [129]. Furthermore, in many ways, mitochondria represent the remnants of proteobacteria, and they contain immunogenic molecules, including mitochondrial DNA (mtDNA), adenosine triphosphate, cardiolipin, cytochrome c, and formyl peptides; these molecules act mainly through FPR1, which displays a high affinity for bacterial- and mitochondrial-derived peptides and may potentiate the immunological response [130]. Moreover, mitochondrial lysates increase the mRNA and protein expression of  $\beta$ -amyloid precursor protein (APP). Considering these observations and the fact that data concerning the impact of brain mitochondria dysfunction on the FPR-mediated pathways are scarce, such studies should provide a new understanding of this field and are strongly recommended [131].

Recently, an increasing body of data seemed to indicate that excess inflammatory processes in the brains of AD patients are linked to disturbances in the RoI and that these deficits are intensified during the aging processes. This hypothesis appears to be supported by preclinical data obtained in Balb mice, in which aging was shown to be associated with dysfunctional RoI and a greater increase in and slower clearance of recruited neutrophils following acute inflammatory challenge, resulting in higher levels of proinflammatory cytokines and deficits in pro-resolving factor production [132]. Wang *et al.* demonstrated in senescence-accelerated mice prone 8 (SAMP8), which is a murine model of accelerated aging that spontaneously exhibits  $\beta$ -amyloid overproduction, tau hyperphosphorylation, oxidative stress damage, and cognitive decline [133], that aging is indeed associated with a proinflammatory state [134]. Furthermore, experimental data indicate that changes in FPR2 activation play a role in these mechanisms. For instance, SAMP8 mice show increases in receptor levels and inflammatory process aggravation. Deficits in the levels of SPM's for this receptor have also been observed. Moreover, in 9-month-old SAMP8 mice, there is a reduced level of leukocyte-type 12-lipoxygenase, which is positively correlated with the elevation of tau phosphorylation at Ser202/Thr205 (AT8), in the hippocampus. This finding suggests that the disturbance of pro-resolving processes and changes in FPR2 activation may influence the formation of  $A\beta$  and tau pathology.

The most recent data obtained in transgenic mice demonstrate that the application of FPR2 agonists may be a new approach for AD therapy through modulating the RoI process and FPR2 activation. In this study, lipoxin and its analogue aspirin-triggered lipoxin A4 were used. In Tg2576 mice bearing the Swedish double mutation in the human amyloid precursor protein, which shows the accelerated development of  $A\beta$ -related pathologies, ATL treatment reduces NF- $\kappa$ B activation and proinflammatory cytokine levels and increases anti-inflammatory cytokine IL-10 levels, thus en-



**Fig (6).** The most important targets of pro-resolution factors in Alzheimer's disease. (1) Neurotoxicity, *e.g.*, dysregulated neurotransmission, glutamate and calcium signaling; (2) microglia and astrocyte activation, which leads to irreversible inflammation and neurodegeneration; (3) the phosphorylation of tau and the formation of neurofibrillary tangles; (4) the formation of Aβ plaques. Proresolving molecules increase the resolution of spontaneous reactions. (5) Aβ plaques can lead to the accumulation and activation of microglia, resulting in an increase in the synthesis of proinflammatory cytokines such as IL-1β and TNF-α. These cytokines can lead to the hyperphosphorylation of tau and a pathological cycle; increased Aβ formation and the persistent activation of microglia ultimately lead to chronic neuroinflammation and neurodegeneration. This schematic was adapted from Filep *et al.*, 2018 [183]. (A higher resolution / colour version of this figure is available in the electronic copy of the article).

hancing the activation of the alternative microglia phenotype [135]. This change in the microglial phenotype is correlated with decreased synaptotoxicity and improvement of cognitive functions in Tg2576 mice.

Dunn *et al.* [136] reported that, in triple transgenic AD (3xTg-AD) mice, a transgenic mouse strain that expresses the Aβ-processing related mutations APP<sup>swe</sup> and PS1M146V as well as mutant aggregation tau (tauP301L), the formation of senile plaques and neurofibrillary tangles is hastened, while the LXA4 level in the brain is declined. Interestingly, 8 weeks of treatment with ATL leads to the restoration of cognitive functions and a reduction in Aβ levels and tau phosphorylation [136].

It has also been recently observed that human CHME3 microglia incubated with Aβ<sub>42</sub> [(compared with those stimulated with lipopolysaccharide (LPS)] exhibit reduced phosphorylation of 5-lipoxygenase [137], a key enzyme involved in the regulation of leukotriene and lipoxin production, at Ser523. This diminished phosphorylation at Ser523 induces higher production of leukotrienes and reduces the formation of lipoxin, and these results are correlated with disturbances in the resolution of inflammation. Interestingly, neither Aβ<sub>42</sub> nor LPS alter LXA4 or RvD1 levels in the cell culture medium. Moreover, unlike LPS, which induces an increase in

FPR2, Aβ<sub>42</sub> does not influence the FPR2 level. Thus, it appears that, compared with the effects of a proinflammatory factor (*e.g.*, LPS), Aβ<sub>42</sub> formation is associated with changes that lead to the distortion of the pro-resolving processes and to the development of chronic inflammation (Fig. 6).

Although age is one of the most relevant risk factors of late-onset AD, researchers have only recently begun to estimate the impact of aging on the resolution of inflammation in humans. Gangemi *et al.* measured the level of LXA4 in the urine and the levels of proinflammatory leukotrienes in 30 healthy volunteers divided into three age groups [138]. Compared with the youngest group, both older age groups showed much lower levels of LXA4 in the urine. Furthermore, the LXA4-to-leukotriene ratio was considerably decreased in the older groups. These studies suggest that the ability of our innate systems to shift proinflammatory leukotriene production towards the formation of pro-resolving lipoxins from arachidonic acid declines with age in humans.

In 2014, Wang and coworkers carried out one of the first trials in patients. They evaluated SPM levels and receptor expression in human CSF and brain tissue and found that neurodegenerative disturbances are related to dysfunctional RoI [139]. LXA4 and RvD1 levels were measured in CSF from patients with AD, mild cognitive impairment (MCI),

and subjective cognitive impairment (SCI). Interestingly, LXA4 levels in the CSF and hippocampal tissue were significantly lower in the AD group than in the MCI and SCI groups. The researchers noted no changes in RvD1 levels in both the hippocampus and CSF. It is worth noting that they observed a positive correlation between LXA4 and RvD1 levels in the CSF and cognitive efficiency, which was measured with the mini-mental state examination [139]. In the same study, immunohistochemical staining of hippocampal tissue from AD patients (compared with that from control subjects) revealed higher levels of SPM receptors, FPR2, and ChemR23 (a RvE1 receptor). The levels of 15-LOX-2, a key enzyme engaged in LXA4 production, were elevated, while those of IL-10 were reduced [140, 141]. The most recent studies by Zhu and coworkers provided additional evidence of disturbed RoI in AD patients [142]. SPM levels in the entorhinal cortex of AD patients and control subjects were assessed 18-21 hours *postmortem*. Compared with those of age-matched controls, the levels of SPM maresin-1, protectin-1, and resolvin D5 were reduced, and these changes were correlated with the dysfunction of the RoI and chronic inflammation.

Recently, it has been postulated that chronic inflammation and RoI dysfunction in elderly AD patients may be potentiated by infectious (bacterial, viral, or fungal) agents, leading to the increased production and deposition of  $\beta$ -amyloid plaques as well as neurofibrillary tangles [143]. In fact, persistent and chronic infections play a role in inducing and amplifying chronic inflammation in AD. Annexin A1, one of the multiple ligands of FPR2, is incorporated into the budding virus membrane of the influenza A virus (IAV). Therefore, once the IAV infects a host cell, FPR2 signaling is activated, leading to an increase in viral replication and the dysregulation of the host immune response and inflammatory response. Interestingly, preclinical studies have proven that FPR2 modulators efficiently protect mice against influenza infections by inhibiting viral replication and deleterious inflammation, which is responsible for tissue injury at later stages of infection. In combination with antiviral drugs (*e.g.*, oseltamivir), FPR2 antagonists might also have a much stronger effect in blocking IAV replication [144]. Therefore, antiviral (*e.g.*, oseltamivir) monotherapy, as well as combined treatment with FPR2 modulators, may provide a new approach for the prevention of AD and/or AD-like pathologies as well as RoI normalization [134, 145].

Unfortunately, the crucial limitation of these promising studies is the low stability and bioavailability of SPMs; therefore, we recently focused on identifying new ureidopropanamide compounds with better pharmacokinetic profiles and evaluating their efficacy for the modulation of FPR2 in transgenic *Fpr2*<sup>-/-</sup> knock-out (KO) mice. Therefore, we decided to compare the potential of new ureidopropanamide-based FPR2 agonists for modulating neurodegenerative and inflammatory changes in an *ex vivo* model of inflammation. We tested their ability to modulate lactate dehydrogenase (LDH) release and metabolic activity in hippocampal organotypic cultures (OHCs). In fact, we evaluated the impact of MR39 (Fig. 5) on LDH release stimulated by LPS and/or oligomeric A $\beta$ 1-42 in both C57BL/6J (wild-type mice - WT) and *Fpr2*<sup>-/-</sup> OHCs (unpublished data). In these conditions,

we found that LPS stimulated LDH release in WT and *Fpr2*<sup>-/-</sup> cultures, while oligomeric A $\beta$ 1-42 potentiated LDH release only in the OHCs obtained from WT animals. Furthermore, co-stimulation (LPS + oligomeric A $\beta$ 1-42) did not potentiate the impact of LPS on the LDH level. Importantly, we observed the pro-resolving properties of MR39 (1  $\mu$ M dose) in WT OHCs only.

These studies are very promising and will be continued because finding new alternative modulators that reduce the levels of endogenous SPMs and an approach for promoting RoI mechanisms seem to be fully justified and are anticipated.

## 6.2. Does Depression Result from RoI Deficits?

In contrast to the wealth of studies supporting the presence of RoI dysfunction in AD, data on abnormalities in SPMs function and/or metabolism in other psychiatric disorders, such as depression and schizophrenia, which are also characterized by neurodegenerative processes, are quite scarce. However, for many years, it has been postulated that the pathogenesis of these diseases are complex, and that, in addition to changes in certain neurotransmitters (*e.g.*, serotonin), the dysfunction of the endocrine system and inflammatory responses, including neutrophil infiltration into the brain and the increased expression of proinflammatory mediators, play a key role [146, 147].

In this context, much attention has been focused on depression. In fact, the first reports that revealed the role of inflammatory processes in the pathogenesis of depression were published by Maes *et al.* (1995, 2009), who demonstrated increased levels of inflammatory biomarkers, such as proinflammatory cytokines, IL-1 $\beta$ , IL-6, TNF- $\alpha$ , and IFN- $\gamma$  in depressed patients [148, 149]. Following these initial reports, other articles confirmed the involvement of inflammation in the pathogenesis of depression. Among them, a study by Zorilla and collaborators [150], as well as one by Dowlati and collaborators [151], showed increased levels of IL-6, TNF- $\alpha$ , and C-reactive protein (CRP) in blood samples from depressed patients. Clinical studies confirmed that, in patients suffering from depression, plasma and cerebrospinal fluid (CSF) levels of IL-1 $\beta$  are elevated and that there is a positive correlation between the serum concentration of this cytokine and depression severity [152, 153]. On the other hand, decreases in the levels of IL-4, TGF- $\beta$  [154], and IL-10 were observed [155], which suggests the dysfunction of the anti-inflammatory response.

Moreover, studies in animal models of depression revealed changes in immune system function. For instance, in rats exposed to repeated intermittent LPS injections (a neurobehavioral model of chronic depression), thymus weight and the proliferative activity of lymphocytes are significantly reduced, and these changes are accompanied by changes in interferon- $\gamma$  (IFN- $\gamma$ ) and IL-10 synthesis in the periphery [156]. Experiments in a mouse model of restraint stress have revealed a higher expression of IL-1 $\beta$  and TNF- $\alpha$  in the hippocampus [157]. Moreover, in an animal model of depression based on chronic mild stress (CMS), in which adult animals are exposed to stress, the levels of IL-1 $\beta$  and IL-6 in the brain and IL-6 and TNF- $\alpha$  levels in the serum are increased [158]. In line with these observations, using a prena-

tal stress model of depression, we demonstrated enhanced microglial activity, increased expression of neurotoxic factors, including proinflammatory cytokines and chemokines (IL-1 $\beta$ , TNF- $\alpha$ , IFN- $\gamma$ , and CCL2) and deficits in the neuronal-glial interaction (e.g., CX3CL1-CX3CR1 and CXCL12-CCR4) in adult rats [159]. It should be noted that changes in immune activity indicative of neuroinflammation are prolonged and present both in young and adult animals and precede the manifestation of depressive behavioral deficits.

Importantly, the intracerebral administration of proteins that possess beneficial homeostatic properties (including IGF-1 and CX3CL1) normalize not only neuroinflammation but also behavioral changes [160]. Therefore, it can be suggested that the immune response of the brain preserves its ability to shift from a proinflammatory to an anti-inflammatory response. Moreover, these observations may indicate changes in the RoI and the potential role of deficits in FPRs, which, as promiscuous receptors, can be activated by structurally diverse agonists that elicit proinflammatory or pro-resolving effects depending on chemical structure.

To date, only a few studies have shown the involvement of FPRs in behavioral changes. In fact, some findings indicated that FPRs can modulate anxiety and fear-elicited responses. It has been shown that *Fpr1*<sup>-/-</sup> knockout mice exhibit increased exploratory activity, reduced anxiety-like behavior, and impaired fear memory [161]. Furthermore, the role of glucocorticoids in the modulation of behavioral deficits has been confirmed, which suggests the potential for FPRs to influence the hypothalamic-pituitary-adrenal (HPA) axis. Other studies have shown that mice lacking *Fpr2/3* show increased exploratory behavior and reduced fear compared to those of WT mice, while these behavioral changes can be partially mimicked by the FPR2 antagonist Boc2 [162]. Interestingly, in both *Fpr2/3*<sup>-/-</sup> mice and Boc2-treated mice, reduced immunostaining for the phospho-p38/MAPK pathway, a key FPR downstream signaling pathway, has been found [162]. It has been demonstrated that, in mice, changes in anxiety-related behavior may result from the genetic deletion of *Fpr2/3*. Therefore, it appears that further studies aimed at discovering the mechanisms of ligand action on FPRs responses may be essential and provide better insight into the regulation of FPRs activation and RoI promotion.

However, only limited data are available so far. Various reports postulated that resolvin D1 exerts anti-inflammatory and pro-resolving actions in animal models of depression based on peripheral inflammation [34]. Resolvin D1 and D2 have been used as antidepressants in an animal stress model (the chronic unpredictable stress model), and their effectiveness has been proven [163]. Furthermore, in an LPS-induced depression model, RvD1 alleviates endotoxin-evoked depression-like behaviors *via* anti-inflammatory actions, including the inhibition of neutrophil chemotaxis and the expression of proinflammatory mediators [164].

Recently, it has also been shown that the pro-resolving lipids RvD1 and RvE1 downregulate LPS-induced proinflammatory cytokine (TNF- $\alpha$ , IL-6, and IL-1 $\beta$ ) gene expression in microglia [165]. These data clearly indicate that these lipids are involved in the RoI. It is crucial to note that the mechanisms of action of the two types of resolvins are

distinct, as RvE1 regulates the NF- $\kappa$ B signaling pathway, while RvD1 regulates miRNA expression [165].

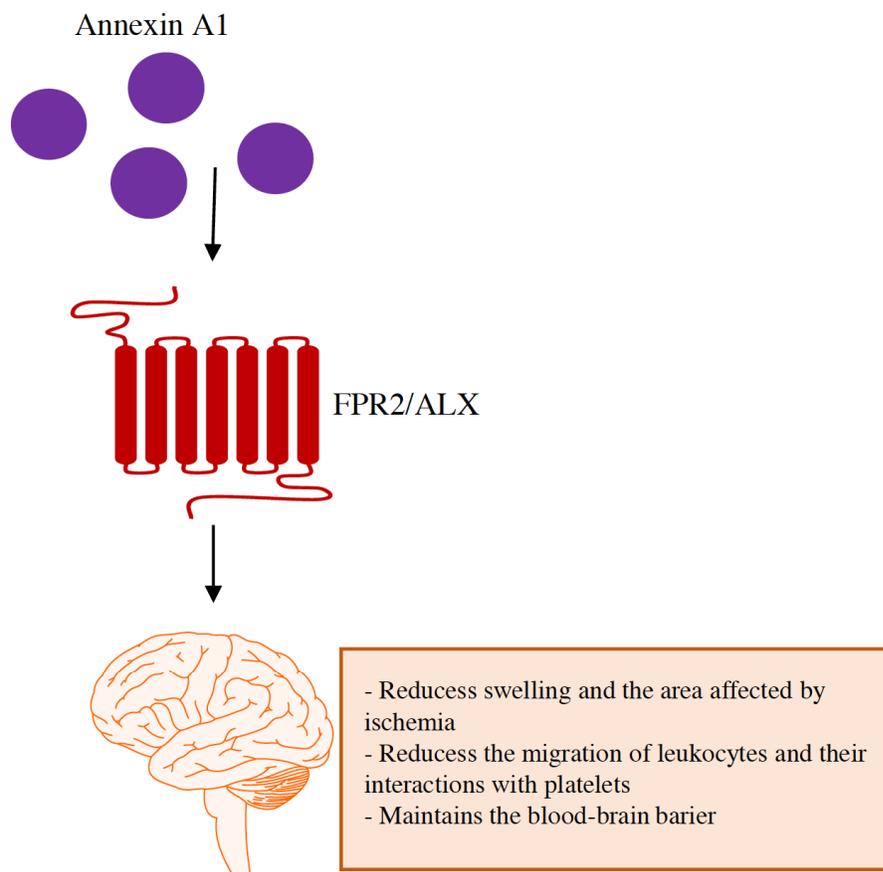
Therefore, it may be postulated that SPMs represent promising novel therapeutic agents that are able to potentiate the RoI in the brain; however, their role in mood disorders remains to be investigated. The anti-inflammatory effects of docosahexaenoic acids and their derivatives in microglial cells were recently analyzed in a special review [166].

Based on these promising reports indicating the usefulness of SPMs, in our preliminary studies, we used ureido-propanamide FPR2 agonists to modulate the proinflammatory activation of glial cells observed in a prenatal stress model. The results obtained thus far indicate that, in primary microglia cultures, FPR2 agonists reduce LPS-induced cell mortality and lower LPS-induced proinflammatory cytokine levels, suggesting pro-resolving and anti-inflammatory properties. Moreover, the observed effects can be partly reversed by pretreatment with an FPR2 antagonist (unpublished data). We found that, in hippocampal organotypic cultures obtained from prenatally stressed animals, the expression of both FPR2 and proinflammatory cytokine genes is upregulated (unpublished data). However, these interesting findings undoubtedly require further confirmation in *in vivo* models, particularly in drug-resistant depression in which coexistent neuroinflammation has been described, for the development of a new supportive therapeutic strategy.

### 6.3. FPR as a Target for Brain Ischemia

Stroke remains the second leading cause of death and the main cause of long-lasting disability in adults despite enormous efforts in searching for efficient therapy [167, 168]. In general, two types of stroke, namely, hemorrhagic and ischemic stroke, have been identified; the latter accounts for 87% of all cases [169]. There are many causes of stroke, including cardiovascular disease, atherosclerosis of large vessels, atherosclerotic plaque rupture, and infarctions caused by the blockade of a deep artery [170].

Excitotoxicity, the distortion of the ionic balance, oxidative stress, and apoptosis are among the mechanisms that lead to cell death during brain ischemia. Their progression and intensity can slightly differ from case to case. On the other hand, it is known that these phenomena impair the function of neurons, glial cells, and blood vessels. Neurons, particularly those from the CA1 area of the hippocampus and from the cortex, cerebellar Purkinje cells, and oligodendrocytes appear to be more prone to injury and cell death than glial or endothelial cells [171]. Many studies demonstrated that inflammation plays an important role in the pathogenesis of ischemic stroke, while the extent of the affected area and ischemia duration are crucial factors related to outcomes. Since this inflammation is not induced by any specific immunogens, this inflammatory response is often termed sterile inflammation [172]. This process is characterized by the fast activation of microglial cells, the formation of proinflammatory mediators, including proinflammatory cytokines and chemokines, and the multistage infiltration of inflammatory cells, *i.e.*, microglial cells, leukocytes, neutrophils, lymphocytes T, monocytes/macrophages, to the damaged nervous tissue. These events contribute to the aggrava-



**Fig. (7).** Annexin A1 and its peptide mimetics are possible therapeutic strategies for the treatment of ischemia based on their ability to regulate inflammation to lead to the resolution of inflammation, decreased tissue injury, and the promotion of resolution. (A higher resolution / colour version of this figure is available in the electronic copy of the article).

tion of cell death [173, 174]. In tissue affected by ischemia, local inflammation involves astrocytes, activated microglia, and infiltrating monocytes and macrophages. Additional damage is generated by reactive oxygen species and the presence of other cells implicated in the inflammatory response (e.g., dendritic cells) [175].

It should be emphasized that, in contrast to the invariably deleterious effects of ischemia, cytokines and chemokines can play a dual role, and the final effect depends on which and how many specific mediators are produced, where they are released and at what time during ischemia or reperfusion they appear. In general, it is thought that  $\text{TNF-}\alpha$ , IL-6, and IL-1 $\beta$  have emerged as central players in the orchestration of mediator responses. Next, during reperfusion, while efficient ROI processes are preserved, anti-inflammatory markers, including CXCL13, Ym1, TGF $\beta$ , and CD163, are expressed [174].

The most recent studies suggested a significant role for FPR ligands in the mechanisms that maintain the balance between proinflammatory and anti-inflammatory pathways to retain homeostasis following ischemic insult. Ligands such as AnxA1 and its mimetic peptides, e.g., N-terminal-derived Ac2-26, are of particular interest [176, 177]. The prevailing view is that AnxA1 regulates cell apoptosis, proliferation, and differentiation [178] but also diminishes leu-

kocyte adhesion and migration, thus inhibiting proinflammatory cytokine release under ischemic conditions [175, 179]. Recent studies also demonstrated that the AnxA1/FPR axis plays a significant role in sealing the BBB in neonatal hypoxic-ischemic encephalopathy [180]. It is well known that the BBB is a critical gateway of communication between the periphery and the brain and that ischemia leads to BBB breakdown. Interestingly, neurodegenerative disorders, such as AD and amyotrophic lateral sclerosis (ALS), are associated with microvascular dysfunction and/or degeneration in the brain, neurovascular disintegration, defective BBB function, and vascular factors. BBB disruption has also been observed in *postmortem* studies of depressed patients. Therefore, the role of endogenous pro-resolving molecules, such as AnxA1, and their impact on FPRs may have a wider biological significance, which undoubtedly should be a subject of further research (Fig. 7).

Transgenic mouse models are particularly useful for studying the role of FPRs in brain injury. In fact, such studies confirmed the involvement of mouse FPR2 and/or FPR3 in circulating neutrophils in mediating AnxA1-induced protection [179, 181]. A study by Vital *et al.* indicated that *Fpr2/3*<sup>-/-</sup> KO mice show an exaggerated inflammatory reaction to brain ischemia that involves the increased infiltration of leukocytes and platelets. An exacerbated cerebral inflam-

matory response as a consequence of the lack of FPR2 and FPR3 has also been demonstrated by other studies [179]. These studies suggested that FPR activity is crucial for the brain ischemia-induced inflammatory response [182]. On the other hand, the use of endogenous FPR2/3 agonists, such as AnxA1, after nervous tissue damage produces a neuroprotective effect by reducing detrimental effects of brain ischemia, including infarct volume, leukocyte adhesion, and inflammatory marker expression in the middle cerebral artery occlusion model. It appears that microglial cells contribute to these effects because, in addition to blood vessels, they are the main cells exhibiting high FPR2 expression in mice.

Bearing in mind that most of the available literature data derive from the studies with endogenous agonists of FPRs, it should be mentioned that there are several reports in which synthetic ligands, characterized by different stability and bioavailability, have been recently published. As an example, synthetic FPR2 ligands have been studied in models of cardioprotection, *e.g.*, CGEN-855A [97]. They demonstrated that FPR1/FPR2 biased agonism can be crucial for the cardioprotective efficacy in the treatment of myocardial ischemia-reperfusion.

We performed preliminary experiments to assess the effects of FPR2 agonists in models of ischemia. In organotypic hippocampal cultures, oxygen-glucose deprivation (OGD), which is considered an *in vitro* model of ischemia, leads to the elevated expression of FPR2, while toxic OGD-induced effects can be modulated by FPR2 agonists (unpublished data). In light of the abovementioned data, it appears that there is still a need to support the RoI process to stimulate a shift between proinflammatory and anti-inflammatory pathways following an ischemic insult. Therefore, a better understanding of the basic mechanisms underlying FPR2 modulation may provide new potential therapeutic options for the treatment of ischemic diseases.

## CONCLUSION

Neuroinflammation plays a dual role in the brain. On the one hand, transient inflammation terminated by a proper resolution process is beneficial and usually elicits neuroprotective effects. On the other hand, the persistence of this process is the cause of chronic inflammation, which leads to long-lasting disturbances in homeostasis and to the development of permanent neurotoxic or neurodegenerative changes that become an underlying cause of CNS diseases. In such circumstances, classic anti-inflammatory therapies do not induce beneficial effects. In addition, classic anti-inflammatory drugs, by strongly suppressing the proinflammatory response, fail to balance proinflammatory and anti-inflammatory components. Thus, the response to future insults that require immune activation is hampered, and the body is rendered defenseless, a condition that may be life-threatening. Therefore, a novel and innovative approach to modulating the inflammatory response is needed. This perspective article illustrates how a resolution-based strategy can offer new therapeutic opportunities, especially for CNS disorders characterized by prolonged neuroinflammation. Based on the data included in this review, FPRs, and FPR2, in particular, are potential novel targets for such therapeutic

strategies. The characterization of the pharmacological properties of FPR2 agonists in both preclinical and clinical trials strongly suggests a prominent anti-inflammatory role in the context of the innate immune response and supports the possibility of developing a resolution-based approach. Lipid SPMs have been proven to possess particularly beneficial properties because they are able to exert both pro-resolving and anti-inflammatory actions. Hence, the search for ligands characterized by an adequate pharmacological profile and bioavailability, which may become widely used to promote endogenous the RoI through FPR activation, appears justified.

We hope that this narrative review has shed more light on the current knowledge of the role of FPRs in the RoI in brain disorders that are different in terms of clinical manifestation but are similar with respect to the involvement of neuroinflammation.

## LIST OF ABBREVIATIONS

7-TM	=	7-transmembrane receptors
AA	=	Arachidonic acid
AD	=	Alzheimer's disease
AnxA1	=	Annexin A1
APP	=	$\beta$ -amyloid precursor protein
AT-LXA4	=	Aspirin-triggered lipoxins
A $\beta$	=	$\beta$ -amyloid plaques
BBB	=	Blood-brain barrier
bFGF	=	Basic fibroblast growth factor
BGF	=	Brain growth factor
cAMP	=	Cyclic adenosine monophosphate
CDK5	=	Cyclin dependent kinase 5
ChemR23	=	Chemokine-like receptor 1
CNS	=	Central nervous system
COX-2	=	Cyclooxygenase 2
CSF	=	Cerebrospinal fluid
DHA	=	Docosahexaenoic acid
EPA	=	3-eicosapentaenoic acid
ERK 1/2	=	Extracellular signal-regulated kinases
fMLP	=	<i>N</i> -formyl-methionyl-leucylphenylalanine
FPR2/ALX	=	Formyl peptide receptor/lipoxin A4 receptor
GPCRs	=	G-protein-coupled receptors
HPA	=	Hypothalamic-pituitary-adrenal
IAV	=	Influenza A viruses
IFN	=	Interferon
IL	=	Interleukin
KO	=	Knock-out mice

LDH	=	Lactate dehydrogenase
LOX	=	Lipooxygenase
LPS	=	Lipopolysaccharide
LTB4R or BLT1	=	Leukotriene B4 receptor
LXA4	=	Lipoxin A4
MARCO	=	Macrophage scavenger receptor
MHC	=	Major histocompatibility complex
NF-κB	=	Nuclear factor-κB
NGF	=	Neural growth factor
NO	=	Nitric oxide
OGD	=	Oxygen-glucose deprivation
PAMP	=	Pathogen-associated molecular patterns
PRR	=	Pattern-recognition receptors
RoI	=	Resolution of inflammation
ROS	=	Reactive oxygen species
RvD1	=	Resolvin D1
SAA	=	Serum amyloid A
SAMP8	=	Senescence-accelerated mice prone 8
SPMs	=	Specialized pro-resolving mediators
TGF	=	Transforming growth factor
TNF	=	Tumor necrosis factor
WT	=	Wild-type mice

### CONSENT FOR PUBLICATION

Not applicable.

### FUNDING

This work was supported by a grant from the Alzheimer's Association (AARG-NTF-18-565227) and by the Polish National Science Centre, grant no. 2017/26/M/NZ7/01048. Natalia Bryniarska is a holder of a scholarship from POWER no. POWR.03.02.00-00-1013/16.

### CONFLICT OF INTEREST

The authors declare no conflict of interest, financial or otherwise.

### ACKNOWLEDGEMENTS

Declared none.

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