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Neuroinflammation and fibrosis in stroke: The good, the bad and the ugly

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Abstract

Stroke is the leading cause of death and the main cause of disability in surviving patients. The detrimental interaction between immune cells, glial cells, and matrix components in stroke pathology results in persistent inflammation that progresses to fibrosis. A substantial effort is being directed towards understanding the exact neuroinflammatory events that take place as a result of stroke. The initiation of a potent cytokine response, along with immune cell activation and infiltration in the ischemic core, has massive acute deleterious effects, generally exacerbated by comorbid inflammatory conditions. There is secondary neuroinflammation that promotes further injury, resulting in cell death, but conversely plays a beneficial role, by promoting recovery. This highlights the need for a better understanding of the neuroinflammatory and fibrotic processes, as well as the need to identify new mechanisms and potential modulators. In this review, we summarize several aspects of stroke-induced inflammation, fibrosis, and include a discussion of cytokine inhibitors/inducers, immune cells, and fibro-inflammation signaling inhibitors in order to identify new pharmacological means of intervention.

Keywords

Cytokines; Fibrosis; Immune cell infiltration; Neuroinflammation; Perlecan; Stroke

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1. Introduction

Stroke, a sudden interruption of blood supply to the brain or blood vessel in the brain, causes neurological deficits (Ojaghihaghghi et al., 2017). Stroke is the 3rd leading cause of death, and every year 15 million people are affected in the world (Jayaraj et al., 2019). In the United States, it is the 5th leading cause, and about 140,000 people die each year (<https://www.cdc.gov/>). The stroke risk varies with gender and age (Roy-O'Reilly and McCullough, 2014). Mortality rates are similar for both men and women below 45 years of age, but women aged 45–74 years are at less risk of stroke mortality than men of the same age (Reeves et al., 2008). Different types of stroke include ischemic stroke (clots), which accounts for 87% of total stroke cases, hemorrhagic stroke (bleeds), transient ischemic attack (TIA, also known as mini-strokes), cryptogenic (of unknown cause) stroke, and brain stem stroke (lack of body function) (<https://www.stroke.org/>). Ischemic stroke is further classified into two subgroups, thrombotic strokes and embolic strokes. Thrombotic strokes are caused by blood clots that develop in the brain blood vessels, whereas embolic strokes are caused by blood clots arising from the body via the bloodstream. Inflammation, initiated by blood supply to the brain is interrupted and is known to play a dual role in a stroke pathology and having both detrimental and beneficial effects (Iadecola and Anrather, 2011). The immune system consists of a wide variety of immune cells, which are important for brain development and function and contribute to several neurological diseases including stroke (Iadecola and Anrather, 2011). The circulating immune cells, such as neutrophils, lymphocytes, macrophages, and endothelial cells, are actively involved in the inflammatory process (Perera et al., 2006). Whereas microglia, astrocytes, and neurons are active contributors to inflammation in brain ischemia. Previous studies suggest that components of the immune system are involved in all stages of stroke-induced inflammation (Iadecola and Anrather, 2011; Urra et al., 2009). The inflammatory and immune responses to stroke results from activation of the innate and adaptive immune system, which may enhance tissue injury and also promote healing in stroke (McCombe and Read, 2008). In this review, we aim to summarize currently known information about how inflammation and, fibrosis underlies stroke pathology, and a better understanding immunology of stroke could be used to guide future research and intervention strategies.

2. Inflammatory response to ischemic stroke

Ischemic stroke causes decreased blood flow in the brain, loss of cellular integrity, and subsequent cellular damage resulting in inflammation (Rock et al., 2010). The onset of ischemic stroke leads to activation of stress signals via tissue hypoxia, glutamate excitation, oxidonitrosative stress, which causes activation of glial cells in the brain. There are two mechanisms of action in the immune system, adaptive immunity and innate immunity (Nakamura and Shichita, 2019). Microglial cells are resident myeloid cells derived from yolk-sac progenitors and are a key component of the innate immune response system (Gomez et al., 2015). At the same time, in response to the stroke injury, other bone marrow derived monocytes are recruited to the damaged tissue and exhibit morphology similar to that of microglia. Danger associated molecular patterns (DAMPs) such as high-mobility group box 1 protein (HMGB-1) derived from dying neurons (Agalave and Svensson, 2015), and further activates innate microglia and other immune cells, via mediating toll like

receptors (TLR), which in turn produce many proinflammatory cytokines, chemokines, and matrix metalloproteinases (MMPs). MMP 9 is upregulated by HMGB1 via TLR4 and its induced cytokines as TNF α or IL-1 β mediating cellular death after ischemic stroke (Qiu et al., 2010). Time-dependent inhibition of MMP-9 improves stroke outcomes via the degradation of DAMPS (Cauwe et al., 2009). Previous studies have demonstrated local and systematic inflammatory responses after stroke in humans (Zaremba and Losy, 2001; Pedersen et al., 2004). The studies with human brains by histology during autopsy showed that neutrophil recruitment starts from the first day and drastically increased within 2 to 3 days after stroke whereas macrophage infiltration begins after three days and persistent for several years in infarct regions (Chuaqui and Tapia, 1993; Mena et al., 2004).

3. Immune cells

Leukocytes, mainly neutrophils invade first to infiltrate the ischemic region and decreases rapidly with time (Jin et al., 2010). In humans, the circulating neutrophils constitute 50–70%, compared 75–90% in rodents (Mestas and Hughes, 2004). Infiltrating neutrophils contributes to inflammation and progression to injury by generation of pro-inflammatory signals, such as inducible nitric oxide synthase (iNOS), matrix metalloproteinases (MMPs), toll like receptor 2 (TLR2), antigen presenting proteins, chemokines and, immunoglobulins which are differentially expressed in mice and humans (Mestas and Hughes, 2004; Iadecola and Anrather, 2011). The use of antileukocyte strategies with antiadhesion molecule such as beta2-integrins are proven to be more effective in transient middle cerebral occlusion (tMCAO) model but not in permanent middle cerebral artery occlusion (pMCAO) (Yilmaz and Granger, 2008). Immune cells such as T lymphocytes, especially CD8+ T and CD4+ T cells, influx into the brain within 3 hr and 24 hr respectively, and are responsible for the damaging effect of this acute phase of stroke (Gill and Veltkamp, 2016; Rogove et al., 2002). Cytokines are key mediators in the inflammatory response following stroke (Doll et al., 2014). While the recent study reported the deleterious effect of interferon- γ in inflammation after stroke (Seifert et al., 2014). Increasing evidence suggests that immune cells are not only involved in the critical event of neuroinflammation as key contributors to stroke pathogenesis but are also important players in the maintenance of central nervous system (CNS) homeostasis (Greenwood et al., 2011). Inhibition of aberrant infiltration of immune cells and pro-inflammatory cytokine is vital to counter the deleterious effects of inflammation in stroke. Immune cells inhibitors/inducers for stroke therapy listed in table 1.

3.1. Neutrophils

Neutrophils are the first and critical cells to invade into injured tissue after stroke, and the severity of pathogenesis depends on their influx in the ischemic region (Chen et al., 2016). After invading the CNS, in the diseased state, neutrophils transform into two different phenotypes, namely N1 phenotype (proinflammatory property) and N2 phenotype (anti-inflammatory property) (Easton, 2013; Kolaczowska and Kubes, 2013; Segel et al., 2011). Studies have shown the increased expression of N2 markers such as chitinase-like protein (Chil3 protein or YM1) YM1/Chil3, and Arginase 1+ proteins after stroke, whereas, N1 neutrophil markers like YM1⁻ were lesser in the ischemic brain following experimental stroke induced by middle cerebral artery occlusion (MCAO) in TLR4 knockout (KO) mice

(Garcia-Culebras et al., 2019). The shift in N2 polarization by peroxisome proliferator-activated receptor gamma (PPAR γ) activation with the agonist rosiglitazone is an essential event of inflammation that participates in neuroprotection after stroke in pMCAO mice model (Cuartero et al., 2013). Recent evidence indicates that neutrophil accumulation is downregulated with the administration of all-trans retinoic acid (atRA) in the MCAO mouse model (Cai et al., 2019). With the decrease in neutrophil accumulation, they transform toward the N2 phenotype in stroke lesions and reduce infarct volume. The atRA treatment suppressed STAT1 signaling through enhancing the expression of suppressor of cytokine signaling-1 protein (SOCS1) (Cai, Wang, 2019). Neutrophil migration into the stroke lesion can be determined by myosin1f levels in the brain after stroke and further study demonstrated that myosin1f KO mice subjected to MCAO had smaller infarcts than wildtype controls (Wang et al., 2019a). Given that neutrophil-mediated proinflammatory cytokines such as interleukin-1 beta (IL-1 β), tumor necrosis factor alpha (TNF- α), interferon-gamma (IFN- γ), and IL-6 are increased in ischemic stroke (Kostulas et al., 1999), targeting them is essential for therapeutic approaches to limiting inflammation after stroke (Doll et al., 2014; Jickling et al., 2015). Blockade of neutrophils by a recombinant neutrophil inhibitory factor (rNIF) such as UK-279, 276, which selective binding to CD11b integrin have not been effective in clinical trials (Del 2004). Similarly, the use of rNIF against neutrophil β 2 Integrin CD18 was failed to improve long-term outcomes after stroke (Smith et al., 2015).

3.2. Microglia

Microglia are the resident macrophages of the brain and possess an ability to maintain constant population at a homeostatic level in the CNS (Bruttger et al., 2015), whilst playing a key role in the immuno-surveillance of the CNS and maintaining a homeostatic environment by eliminating cellular debris (see (Thurgur and Pinteaux, 2019) for review). Microglial activation is considered to be the first step in the initiation of inflammation after stroke and is considered to be detrimental (Block et al., 2007). However, a recent study demonstrated that selective elimination of microglia leads to increased infarct volume and neuronal death with calcium overload, which is reversed by microglial repopulation, and the same study further found that microglia perform a critical role in the clearing of damaged neurons and promoting neuronal survival in the injured mice brain (Szalay et al., 2016). Microglial cells express macrophage like markers under normal physiological conditions, although several candidate markers like CD45 and CD39 can be used to differentiate microglia from the peripheral macrophage population (Dudvarski et al., 2016). Furthermore, several studies have identified markers such as Iba1^{high}, CD206⁻ CD45^{low}, CD163⁻ CD11b⁺, MHCII⁺, F480⁺, Cx3cr1^{high}, Ly6C⁻ being specifically expressed by activated microglia in the brain (Ginhoux et al., 2010; Kierdorf et al., 2013). Activation of microglia varies with the acute and delayed phase after injury and remains active from weeks to months after acute injury (Kabba et al., 2018).

Previously, microglia were referred to as M1/M2-like phenotype (activated macrophage 1 (M1) and alternatively activated macrophage 2 (M2)). However, several studies have recently demonstrated that microglial phenotypes are complex and cannot be classified strictly into just these two different classical phenotypes (Chiu et al., 2013; Geissmann et al., 2010; Kan et al., 2015). Regardless, the morphological and phenotypic changes of the

microglia after stroke are often accompanied by increased expression of IL-6, TNF- α , MMPs, as well as chemokines CCL2, CX3CL1, MIP-1, and free radicals (Kabba et al., 2018). A cell based therapy using preconditioned microglia by oxygen glucose deprivation (OGD) induces microglia to acquire an M2-like phenotype leading to secretion of neurotrophic factors vascular endothelial growth factor (VEGF), transforming growth factor- β (TGF- β), as well as MMP-9, rendering microglial modulation as a therapeutic strategy for ischemic stroke in rats (Kanazawa et al., 2017). One mechanism by which ischemic preconditioning triggers neuroprotection is via induction of low-grade inflammation. In line with this, our group previously reported that intra-arterial administration of low dose IL-1 α reduces microglial activation, improves functional outcomes and triggers neurogenesis and angiogenesis after experimental stroke, demonstrating that low levels of IL-1 α regulate microglial activation and has likely beneficial effects on neuroinflammation following stroke (Salmeron et al., 2019). A human study reported the intravenously administration of minocyclin with 10 mg/kg alone and in combination with tissue plasminogen activator is safe and was effective in multiple preclinical stroke models by reduction of microglial activation, reduced NO production, and inhibition of MMP activity in stroke patients (Fagan et al., 2010).

3.3. Macrophages

Macrophages are one of the blood-borne monocytes that infiltrate the brain parenchyma after stroke from perivascular spaces. They perform a critical role in neuroinflammation after ischemic stroke. They migrate through endothelial cells of the blood-brain barrier (BBB) to the stroke lesion under the action of cytokines, chemokines, and cell adhesion molecules (Jian et al., 2019). Macrophages are characterized as expressing different markers such as CD206⁺, CD163⁺, CD45^{high}, CD11b⁺, MHCII^{high}, Ly6C^{low}, F480⁺, Cx3cr1^{low}, Iba1^{low} under diseased state (Goldmann et al., 2016; Zeisel et al., 2015; Faraco et al., 2017; Faraco et al., 2016) and are commonly referred to as non-parenchymal macrophages, to differentiate them from microglia. According to Ritzel and colleagues, more recently activated, BrdU-positive (a marker of cell division) macrophages rather than microglia were observed in the ischemic brain of mice, and these macrophages exhibited increased IL-1 β production (Ritzel et al., 2015). The anti-inflammatory role of macrophages is characterized by expression of several cytokines including TGF- β , IL-4, IL-10, and IL-13 following stroke (Hu et al., 2015; Jian et al., 2019). A study revealed that half of the monocyte derived macrophages/mononuclear cells accumulate in the stroke-injured hemisphere (Wattananit et al., 2016). Another half of spontaneously recruited monocytes migrate to the injured site and contribute to long-term behavioral recovery via expression of TGF- β , Ym1, and CD163 in mice after MCAO (Wattananit et al., 2016).

3.4. Lymphocytes

Lymphocytes are essential subtypes of white blood cells in immune systems and include T cells and B cells. They are mainly involved in both pathogenesis and protective mechanisms in ischemic stroke (Liesz et al., 2015).

3.4.1. T lymphocytes (T cells)—T cells have multiple roles in ischemic stroke and cause inflammation after entering infarcted tissue by the release of pro-inflammatory

B (Breg) cells via the expression of IL-10 and were shown to strongly reduce the infarct volume in ischemic mice (Seifert et al., 2018). This observation demonstrated that the transfer of enriched Breg cell populations might be of important therapeutic value in stroke patients (Seifert et al., 2018). A study evaluated the effect of fingolimod, and it was administered orally (0.5 mg per day for 3 consecutive days) in 22 patients by comparing 11 control patients. The results showed that 11 fingolimod recipients had lower circulating lymphocyte and better neurological outcomes (Fu et al., 2014).

3.5. Dendritic cells

Dendritic cells/antigen-presenting cells (APCs) are immune cells responsible for the initiation of the adaptive immune response, and a recent study demonstrated that these cells are also found in the brain parenchyma after stroke (Kostulas et al., 2002). Conventional type 2 dendritic cells, IRF4+/CD172a+ infiltrate into ischemic brain while expressing IL-23 (Gelderblom et al., 2018). IL-23 receptor KO mice have been reported to be protected against ischemic stroke, albeit with defective IL-17 levels (Gelderblom et al., 2018). The study showed increased expression of dendritic cell-associated C-type lectin-1 (dectin-1) and spleen tyrosine kinase (Syk), which triggers neuroinflammation in a mouse model of cerebral focal ischemia. The dectin-1 antagonist laminarin-1 and Syk inhibitor piceatannol decrease TNF- α as well as inducible nitric oxide synthase (iNOS) expression, resulting in smaller infarct volume, and improved neurological score in a mouse model of cerebral focal ischemia (Ye et al., 2020). Hence, dectin-1/Syk mediated signaling is an important therapeutic target after ischemic stroke (Ye et al., 2020). Fisetin, a flavonoid administered before and after experimental cerebral ischemia, reduced the number of CD11c⁺ cells in the brain in a mouse tMCAO model (Gelderblom et al., 2012a). This study also suggests that fisetin mediated suppression of NF κ B activation and JNK/Jun phosphorylation is neuroprotective in cerebral ischemia as well (Gelderblom et al., 2012a).

4. Pro and anti-inflammatory cytokines role in neuroinflammation after stroke

Cytokines are immunoregulatory molecules released by immune cells in systemic circulation as well as in the CNS in response to various stimuli in order to re-establish homeostasis. An anti-inflammatory response opposes pro-inflammatory cytokine signals, and an imbalance between them leads to localized tissue and organ damage in stroke. Cytokine inhibitors/inducers for stroke therapy are listed in table2.

4.1. Pro-inflammatory cytokines

4.1.1. Interleukin-1 (IL-1)—IL-1 is a master regulator of inflammation and immunity and plays a pivotal role in most, if not all, inflammatory disease. The role of IL-1 in neuroinflammation induced by stroke and its associated risk factors that are linked with raised systemic inflammatory profile has been long established (Sobowale et al., 2016). Hence, current therapies aimed at targeting IL-1 actions in stroke are currently underway. The IL-1 family comprises 11 cytokines and a large family of IL-1 related receptors. These networks of cytokines regulate innate immune cells and play a key role in inflammation after stroke (Dinarello, 2011). IL-1 β is the main released isoform of the IL-1 family, expressed

primarily by immune cells, as a pro-IL-1 β released extracellularly via an NLR family pyrin domain containing 3 (NLRP3)/caspase-1 dependent mechanism (Weber et al., 2010). The role of IL-1 β in stroke has been long established with early studies demonstrating that IL-1 β is expressed in the brain after experimental stroke (Liu et al., 1993; Wiessner et al., 1993). Several published studies demonstrated that pharmacological inhibition of IL-1 β actions by using the IL-1 receptor antagonist (IL-1Ra) confer neuroprotection after experimental stroke (Garcia et al., 1995, Relton and Rothwell, 1992). Interestingly, studies using IL-1 KO mice showed that genetic deletion of IL-1 β failed to affect ischemic brain injury, whereas deletion of IL-1 β and IL-1 α (second main isoform that is intracellularly stored) induces a significant reduction in brain damage (Boutin et al., 2001), hence both IL-1 α and IL-1 β play an important compensatory effect in stroke. Further study found that both centrally- and peripherally-derived IL-1 α and IL-1 β contribute to stroke pathogenesis (Denes et al., 2013). IL-1 α is also expressed by microglia after stroke, although expression precedes that of IL-1 β expression (Luheshi et al., 2011), suggesting that both isoforms may exert specific non-overlapping actions in stroke. In light of this, our recent study found that IL-1 α , but not IL-1 β , induces brain cells to generate the laminin-like globular domain 3 (LG3) neuroprotective protein fragment of the extracellular matrix component perlecan, a prominent heparan sulfate proteoglycan extracellular matrix (ECM) component of the BBB (Saini et al., 2011), whereas IL-1 α is a key inducer of angiogenesis and neurogenesis after stroke (Salmeron et al., 2019). Therefore, alternative strategies aimed at selectively targeting IL-1 α or IL-1 β might prove more effective than complete IL-1 blockade by IL-1ra. In a randomized phase, II study reported that of intravenous (IV) administration of IL-1ra in patients with acute stroke showed reduced cerebral inflammation and greater reduction in National Institutes of Health Stroke Scale at three months (Emsley et al., 2005).

4.1.2. Interleukin-6 (IL-6)—IL-6 is a pro-inflammatory cytokine and is a critical biomarker used to predict stroke associated infection in elderly patients (Kwan et al., 2013). Studies proved that significantly higher serum IL-6 levels correlate with infarct volume and cerebral perfusion deficits in stroke-affected patients than healthy controls (Hotter et al., 2019; Jenny et al., 2019; Saroj et al., 2018). *Il6* gene polymorphisms also have a significant association with increased risk of ischemic stroke (Zhou et al., 2019). Recently, studies using pigs show IL-6 was increased after stroke and further elevated by wild type tissue-type plasminogen activator (tPA). Indeed, co-administration of LMT-28 with wild type tPA blocked JNK and endothelin 1 mediated increase of IL-6, reducing cerebrovascular autoregulation impairment, which in turn, would lead to improved outcomes of tPA treatment after stroke in pigs (Armstead et al., 2019). Although IL-6 has proinflammatory properties, it also has beneficial potential; IV administration of IL-6 to stroked mice results in reduced infarct volume and improved functional outcomes in IL-6 KO mice (Gronhoj et al., 2017).

4.1.3. Interleukin-18 (IL-18)—IL-18 is a cytokine of the IL-1 family and proinflammatory cytokine that plays a critical role in neuroinflammation and stroke (Kandikattu et al., 2019). A recent study demonstrated that serum IL-18 levels were higher in stroke patients, is increased with the severity of the stroke, and can be used as a diagnostic marker for stroke (Hao et al., 2019). *Il18* (137G/C and 607C/A) gene polymorphisms are

associated with an increased risk of ischemic stroke (Zhou et al., 2019). IL-18 level was reported to be increased in ischemic mice brain that showed depression-like behaviors, and the blockage of endogenous IL-18 by IL-18 binding protein rescued depressive phenotypes in spatial restraint-stressed mice. IL-18-mediated depressive behaviors are regulated by the interaction between the IL-18 receptor and NKCC1. NKCC1 antagonist bumetanide showed a therapeutic effect for post-stroke depression in IL-18-induced depressive mice (Wu et al., 2020). However, a study from our group showed that genetic deletion of IL-18 in mice had no effect on brain injury after stroke (Wheeler et al., 2003), indicating that IL-18 may be involved in mediating the depression occurring due to stroke, more than other aspects of stroke pathogenesis.

4.2. Anti-inflammatory cytokines

4.2.1. Interleukin-4 (IL-4)—IL-4 triggers a pleiotropic phenotype in both microglia and macrophages and is involved in diverse immune responses of M2 microglia polarization (Francos et al., 2016). Results from a previous study showed that IL-4 induces PPAR- γ mediated activation of M2 polarization. Protein expression of translocator protein (TSPO) antagonist PK11195 treatment modulates IL-4 expression that subsequently promotes increased expression of CD206, Arg-1, YM-1, and FIZZ-1 under hypoxic ischemia (Zhou et al., 2020). Intraperitoneal administration of dimethylxalylglycine, a HIF-1 α activator, reduced the infarct size by promoting IL-4 and IL-10 levels (Yang et al., 2018). Elevated HIF-1 α activity had a synergistic effect with limb remote ischemic preconditioning (RIPC) on reducing infarction volume after stroke in rats by inhibiting prolyl hydroxylase (PHD) enzyme inactivation (Yang et al., 2018). In accordance with this notion, mice that lacked IL-4 were associated with worse neurological score outcome, along with increased immune cell infiltration and Th1/Th2 ratio in the infarct region (Xiong et al., 2011). Together, these findings indicate that IL-4 signaling has a beneficial role of reducing inflammation in the ischemic core.

4.2.2. Interleukin-10 (IL-10)—IL-10 mediated neuroprotection is well studied in brain injury. As an anti-inflammatory cytokine, it promotes neuronal cell survival via various signaling pathways such as suppression of tumorigenicity 2 (ST2)/IL-33 (Liu et al., 2020; Mills, 2001). Previous study using IL-10 KO mice showed that increased infarct volume and neurologic deficits were observed in ischemic mice (Perez et al., 2013). IL-10 deficiency in mice was reported to promote the expression of CTLA-4, a T-cell inhibitory molecule in the ischemic tissue (Perez et al., 2013). A recent study explained that transplanted mesenchymal stem cell (MSC)/IL-10 intravenously injected through the catheter at 0 or 3 hr after ischemia-reperfusion significantly reduced infarct volume and enhanced motor functional recovery at 72 hr and 7 days after MCAO in rats as an acute phase of ischemic stroke (Nakajima et al., 2017). A recent meta-analysis revealed *IL10* gene polymorphism (1082A/G) to be associated with ischemic stroke (Liu et al., 2017). In another study focused on different promoter regions of IL-10, functional polymorphisms at -1082 promoter region of IL-10 was found to be rare in the Chinese Han population compared to American/European people, and it was concluded that it might be a protective factor for ischemic stroke (Tong et al., 2018).

4.2.3 Interleukin-13 (IL-13)—IL-13 is a key anti-inflammatory cytokine secreted by activated T cells. Transplanting IL13-expressing MSCs leads to a M2 microglia phenotypic switch with a significant increase of Arg-1 and decreased expression of major histocompatibility complex II (MHC-II) in a mouse model of MCAO (Hamzei et al., 2018). A recent study revealed that peripherally administered IL-13 after pMCAO in mouse significantly reduced the infarct volume, increased levels of Arg1 and Ym1, and improved neurologic deficit functions (Kolosowska et al., 2019). A previous study has shown TREM2 as neuroprotective in the ischemic penumbra of a mouse model of MCAO. TREM2 expression also increased in IL-4/IL-13-treated microglia under OGD (Zhai et al., 2017).

4.2.4. Interleukin-33 (IL-33)—IL-33 is a cytokine of the IL-1 family and a crucial mediator of the immune response (Chen et al., 2018; Liew et al., 2016). IL-33 is neuroprotective after ischemic stroke via regulating the inflammatory response (Luo et al., 2015). A recent study demonstrates the role of IL-33/suppression of tumorigenicity 2 signaling in microglial activation and neuroprotection after stroke in mice (Yang et al., 2017). The activated microglia, in turn, release IL-10 which helps in neuronal survival under *in vitro* OGD conditions (Yang et al., 2017). Previous findings demonstrate that the administration of recombinant mouse IL-33 protein to mice before MCAO helps to attenuate brain damage and CNS inflammation (Luo et al., 2015). The neuroprotective effect of IL-33 might be associated with inhibition of neuroinflammation via turning on the T helper 1 and 2 (Th1/Th2) response while suppressing Th17 immune response (Luo et al., 2015). IL-33 acts as a neuroprotectant via mediating regulatory T cell (Tregs) response in experimental ischemic stroke (Xiao et al., 2019). However, another study showed that IL-33 treatment increased the number of Tregs in the MCAO model and suggested that IL-33-ST2 signaling-mediated neuroprotection in stroke (Liu et al., 2020).

5. Mechanisms of neuroinflammation and the inflammatory pathway

Fig. 1 depicts a schematic of stroke-induced neuroinflammation. Stroke induces activation of stress signals, hypoxia, and oxidonitrosative stress that leads to activation of NLRP3 and NF- κ B (nuclear factor kappa-light-chain-enhancer of activated B cells) signaling, infiltration of immune cells and cytokines, and neuroinflammation. Several reports indicate that these signals are activated via mitochondrial damage (Gong et al., 2018; Kandikattu et al., 2017a), lysosomal degradation (Aftabizadeh et al., 2019), reactive oxygen species (ROS) induction (Xu et al., 2018), Ca²⁺ release mediated ER stress (Guo et al., 2018a; Kandikattu et al., 2020; Nakka et al., 2010), and autophagy (Kandikattu et al., 2017b; Wang et al., 2019b). Understanding the stress inducers that activate inflammatory pathways and the development of drugs that inhibit inflammation is key in the treatment of stroke. Other pharmacological therapies with anti-inflammatory activity for stroke therapy are listed in table 3.

5.1. DAMPs and NLRP3 Inflammasome pathway

Damaged-associated molecular patterns (DAMPs) are released from injured tissue after stroke (Umahara et al., 2018). DAMPS are a large family of factors which constitute pathogen-associated molecular patterns (PAMPS) and alarmins (Tang et al., 2012). Alarmins are further divided into protein alarmins such as HMGB1 or heat shock proteins (HSPs), and

non-protein alarmins such as adenosine triphosphate (ATP) (Gulke et al., 2018; Lucchese et al., 2019). HMGB1 is a cytokine-like nuclear protein and mediator of neuroinflammation and pathogenesis after ischemic stroke (Ye et al., 2019). HMGB1 is released into the bloodstream and induces inflammation via its receptors TLR2, TLR4, and receptor for the advanced glycation end product (RAGE) (Ye et al., 2019). HMGB1 is highly expressed in blood under both acute phase and for about 2 weeks after stroke in rats (Kim et al., 2006). Previous experiments have demonstrated that lipopolysaccharides administration induced the release of HMGB1 in a rat model of MCAO (Kim et al., 2018). In this study, blocking HMGB1 function by treatment with HPep1 reduces infarct volume and is considered as a therapeutic target for preventing lipopolysaccharides induced post-stroke infection (Kim et al., 2018).

NLRP3 inflammasomes are known as one of the novel inflammatory pathways discovered in ischemic stroke. They are responsible for mediating cellular damage and death after stroke (Abulafia et al., 2009). Under stroke-induced stimuli, NLRP3 assembles by an apoptosis-associated speck-like protein containing a caspase (ASC), and caspase-1 leading to release and maturation of IL-1 β . NLRP3 inflammasomes are first expressed in microglial cells, followed by microvascular endothelial cells and neurons under oxygen-glucose deprivation/reoxygenation (OGD/R) (Gong et al., 2018). Mitochondrial dysfunction leads to NLRP3 activation in microglia *in vitro*. Hence, the mitochondrial protector diazoxide could inhibit NLRP3 mediated inflammation after stroke in the rat model of tMCAO (Gong et al., 2018). NLRP3-inflammasome inhibitor, MCC950 treatment, reduced infarction via decreased expression of TNF- α , Poly (ADP-ribose) polymerase (PARP), Caspase-3, and I κ B α levels and showed protection in mouse model of tMCAO (Ismael et al., 2018). However, contradicting results from another study suggest, through the use of NLRP3 KO mice or targeting NLRP3 with a pharmacological inhibitor, MCC950 showed that NLRP3 is not a critical mediator in ischemic brain damage (Lemarchand et al., 2019).

6. Fibrosis

Inflammation is a prime driver for the induction of fibrosis in various diseases, including stroke. Immune cell infiltration and cytokine response in stroke induce profibrotic proteins like TGF- β , collagens, MMPs accumulation, ECM deposition, and epithelial to mesenchymal transmission in infarct regions of ischemic stroke.

6.1. Extracellular matrix deposition in ischemic stroke

ECM proteins are rapidly increased in tissue and cells under pathological stress conditions. The ECM consists of a group of proteins that bind to cell surface receptors and regulates many genes involved in cellular behavior. ECM proteins have both harmful as well as protective roles in stroke (Kawakita et al., 2019). In the stroke environment, the composition of ECM proteins is altered, which is implicated in BBB disruption that causes brain damage (Baeten and Akassoglou, 2011).

6.1.1. ECM proteins of the BBB in stroke—ECM proteins participate in the multicellular activity and support systems for many cells when in complex with integrins (heterodimeric receptors containing a transmembrane α and β protein subunit), (Edwards

and Bix, 2019). ECM proteins, mainly composed of proteoglycans, glycoproteins, and collagens, are primarily present in the basement membrane (BM) of brain microvessels between endothelial cells (Summers et al., 2013) and astrocyte end-feet (Baeten and Akassoglou, 2011). ECM proteins, fibronectin, laminins and collagen type I and type IV play a critical role in fibrosis and are induced in the infarct region after stroke (Baeten and Akassoglou, 2011; Summers et al., 2013). Platelet-derived growth factor receptor β (PDGFR β) is an essential molecule for pericyte proliferation and survival and is a mediator of fibrosis induced in stroke. Research from our group demonstrated that a protein portion of perlecan (an ECM proteoglycan) called domain V (DV) plays a crucial role in modulating PDGF responses in angiogenesis (Bix et al., 2007; Bix and Iozzo, 2008). Administration of SU11652, an inhibitor of PDGFR β , reduces fibrosis through decreased desmin and α -smooth muscle actin (α -SMA) levels in vascular cells (Makihara et al., 2015). However, further studies reported that fibrotic scar development after stroke does not primarily occur by PDGFR β + pericytes, and that it is not a contributor to the fibrotic ECM in pMCAO mice (Roth et al., 2020). Previous studies showed that the fibronectin-splicing variant containing extra domain A (Fn-EDA) was elevated in the plasma of diabetes mellitus and hypercholesterolemia comorbid patients (Dhanesha et al., 2015). Further studies showed smaller infarcts and lesser expression of phospho-NF- κ B p65, IL-1 β , and TNF- α levels in the MCAO model of apolipoprotein E-deficient mice expressing Fn-EDA (Fn-EDA+:Apoe KO mice) (Dhanesha et al., 2015). Still further studies investigated the role of plasma versus endothelial Fn-EDA in stroke exacerbation in the comorbid condition of hyperlipidemia. These observations suggest that plasma Fn-EDA KO:Apoe KO mice displayed improved stroke outcomes compared with endothelial Fn-EDA KO (Fn-EDAfl/flTie2Cre) mice, and endothelial-specific KO mice did not contribute to stroke outcome. Hence it suggests that plasma Fn-EDA exacerbates stroke outcome by promoting post-ischemic secondary thrombosis. Therefore, targeting plasma Fn-EDA may help to reduce brain damage after reperfusion (Dhanesha et al., 2019).

6.1.2. ECM receptors of the BBB in stroke—Ubiquitously expressed cellular ECM receptors are primarily dystroglycan and integrins, (Edwards and Bix, 2019). Amongst many different types of integrins, endothelial cells express fibronectin receptor α 5 β 1, α 4 β 1 and α v β 3 integrins (Guell and Bix, 2014; Roberts et al., 2017). We have demonstrated that α 5 integrin, (an obligate pair to the β 1 integrin subunit) endothelial cell-specific knockout mice (α 5-EC-KO) were resistant to ischemic infarction after tMCAO in mice. We further demonstrated that α 5 integrin destabilizes the BBB via decreased expression of claudin-5 after stroke, suggesting that this integrin could be a therapeutic target for stroke (Guell and Bix, 2014; Roberts et al., 2017). This was further confirmed by post-stroke administration of the α 5 β 1 inhibitor, ATN-161, which significantly reduced the expression of α 5 β 1 in the infarct region, stabilized the BBB, and increased collagen IV expression, whilst further reducing CXCL12, MMP-9, IL-1 β , and CD45 + cells in the brain. These results suggest that ATN-161 is a promising novel stroke therapeutic (Edwards et al., 2019).

α 4 β 1 integrin, also known as VLA-4 or CD49d/CD29, is primarily localized in lymphocytes, monocytes, and macrophages. Some researchers in both rodent and preclinical studies reported that inhibition of α 4 showed reduced infarct volumes and improved

functional deficits (Becker, 2002; Llovera et al., 2015) in stroke patients. A recent study by using hydrogels precisely controlled $\alpha_3/\alpha_5\beta_1$ integrin binding and promoted endothelial sprout clumping under *in vitro* (Li et al., 2017). Further, these hydrogels (containing nV and fragments) were injected directly into the stroke cavity promoted non-tortuous blood vessel formation and non-leaky blood vessels by 10 days after stroke. Hence precisely controlled integrin activation from a biomaterial can be used to direct therapeutic vessel regeneration and reduce VEGF-induced vascular permeability *in vivo* (Li et al., 2017). In preclinical studies targeting $\alpha_M\beta_2$, by rNIF (UK279276) and humanized Hu23F2G (Leukarrest) were failure to target $\alpha_M\beta_2$ integrin might be due to not increase of $\alpha_M\beta_2$ expression in human stroke patients compared to rodents (Caimi et al., 2001). $\alpha_6\beta_4$ integrin is expressed on both astrocytes and endothelial cells (Milner and Campbell, 2006). The expression was decreases within 2–4 hrs after MCAO and increases from day 4 to day 14 (Wagner et al., 1997). A recent study on myeloid-specific integrin $\alpha_9\beta_1$ KO mice improved stroke outcome by inhibiting post-ischemia/reperfusion inflammation and reduced fibrin, platelet thrombi, neutrophil, NETosis, and decreased phospho-NF- κ B, tumor necrosis factor- α , and IL- 1β levels in tMCAO mice brain. (Dhanesha et al., 2020).

6.1.3. Role of perlecan in BBB after stroke—ECM proteins generated and proteolytically processed with BBB disruption such as perlecan, play important roles in pathology after stroke (Lee et al., 2011). Research from our group demonstrated that one such proteolytic protein component of perlecan, DV, plays a crucial role in modulating PDGF responses in angiogenesis (Bix et al., 2007; Bix and Iozzo, 2008). Perlecan DV, a C-terminal perlecan fragment that can be generated by an unknown protease(s) *in vivo* after experimental stroke (Lee et al., 2011), can be further processed into the protein fragment LG3 via proteases cathepsins B and L (Bix et al. 2004; Gonzales et al. 2005; Cailhier et al. 2008 and Saini and Bix 2012), and BMP-1 (Gonzalez et al., 2005). LG3 has been shown to be present in the blood, cerebrospinal fluid and the urine of human patients with end-stage kidney disease, thus demonstrating that it is a proteolytic fragment *in vivo*, especially in humans (Adkins et al., 2002; Cartier et al., 2004; Pieper et al., 2004). Increased levels of LG3 were observed in primary fetal cortical neurons (FCN) under OGD/reperfusion condition, which was not blocked by the cathepsin L specific inhibitor, Z-FY-CHO. However, the cathepsin B inhibitor CA074 could inhibit LG3 production under OGD (Saini et al., 2011). Furthermore, IL- 1α treatment increases LG3 levels via cathepsin L and B mediated expression in FCN (Saini and Bix, 2012), and promotes angiogenesis in brain endothelial cell cultures, and induces angiogenesis derived VEGF and CXCL1 expression and therefore, IL- 1α is neuroprotective against stroke (Salmeron et al., 2016; Salmeron et al., 2019). Increased expression of DV and LG3 in the ischemic core of both mice and rats after stroke was shown to be beneficial (Bix, 2013; Saini and Bix, 2012). In seeming agreement with the importance of endogenous DV after stroke, perlecan hypomorph mice that express 10% of total protein (Pln $^{-/-}$ mice) showed larger ischemic stroke lesions following stroke (Lee et al., 2011; Yanagihara et al., 1984). Further, perlecan DV expression is reported to be upregulated in the brains of stroke patients (Trout et al., 2020). Studies using perlecan deficient mice showed significantly fewer NPCs at the subventricular zone (SVZ) following MCAO and leads to larger infarcts and decreases in neurogenesis (Trout et al., 2020). Administration of DV to mice 7 days after stroke enhanced neurogenesis and

improved neurological deficit function with less infarct volume (Trout et al., 2020). Collectively, these studies suggest that perlecan DV could be a novel therapeutic for ischemic stroke (Bix, 2013; Marcelo and Bix, 2014; Parham et al., 2014; Trout et al., 2020).

6.1.4. Role of other basement membrane components in the BBB after stroke

Laminin is a heterotrimeric protein that occurs in 15 different isoforms (including α , β and γ subunits). The increased expression of laminin in both endothelial cells and astrocytes of ischemic penumbra are observed within 24 hrs after rat MCAO (Kang and Yao, 2020). Endothelial laminin-10 is essential for BBB integrity after *in vitro* OGD by regulating occludin and ZO-1 expression and localization to the extracellular cell wall, decreasing paracellular resistance through the endothelial cells (Kangwantas et al., 2016). Type IV collagen are a major component of all basement membranes (Mao et al., 2015). A recent study with small stroke patient populations revealed the variation in *COL4A1* and *COL4A2* genes causes of weakness of the basement vascular membranes causes perinatal arterial ischemic stroke (Kocak et al., 2020). However, collagen IV-deficient mice (null allele of the *Col4a1/2* locus in mice) die at E10.5– E11.5 due to vascular bleeding in the heart and arteries (Poschl et al., 2004). The proteolytic fragment of type IV collagen, tumstatin are generated by MMP-9 proteolysis and are known to suppress angiogenesis via $\alpha V\beta 3$ integrin (Hamano et al., 2003). Studies on collagen IV expression need more investigation and improvement before any determination of collagen IV's impact on stroke severity, or poststroke recovery can be made (Edwards and Bix, 2019).

6.2. Matrix metalloproteinases in ischemic stroke

MMPs are a large family of proteolytic enzymes with crucial roles in ECM remodeling and BBB disruption (Chang et al., 2016). In turn, they affect leukocyte infiltration and subsequent inflammation, in addition to cerebral edema. Neutrophils express different types of MMPs at the injury site after ischemic stroke (Jickling et al., 2015). A number of studies have highlighted the increase in levels of MMPs 1, 2, 3, 8, 9, 10 and 13 in response to stroke (Chelluboina et al., 2015a; Chelluboina et al., 2015b; Cuadrado et al., 2009; Hafez et al., 2016; Han et al., 2016; Hirono et al., 2018; Ma et al., 2016; Orbe et al., 2011; Roncal et al., 2017). In addition, recent meta-analyses have revealed *Mmp* gene polymorphisms that are implicated in predispositions to ischemic stroke (Wang et al., 2018; Zhang et al., 2018). Although most of the MMPs act as a pro-inflammatory factor, MMP-9 has a vital role in neuronal proliferation and apoptosis (Morancho et al., 2010; Vandooren et al., 2014). Time-dependent inhibition of MMP-9 improves stroke outcomes via the degradation of DAMPS (Cauwe et al., 2009). Indeed, the timing of inhibition of MMPs is thought to be critical in terms of therapeutic effect, because MMPs have been shown to play different roles based on timing relative to stroke onset. In the early stages after stroke, MMPs have been reported to contribute to the injury process (Asahi et al., 2001; Lee and Lo, 2004; Lo et al., 2003), while later on, they contribute to the repair and recovery process (Zhao et al., 2006). Previous results demonstrated that MMP-12 upregulation leads to increased expression of other proteases such as MMP-9 and MMP-2 that could contribute to disruption of tight junction proteins in ischemic tissue after MCAO (Chelluboina et al., 2015a).

Finally, the activity of MMPs is tightly regulated via interactions with Tissue Inhibitors of Metalloproteinases (TIMPs) (Cuadrado et al., 2009; Hirono et al., 2018) and ADAMs (a disintegrin and metalloproteinases) (Montaner et al., 2019), as well as the interactions with other MMPs. Recent studies have shown that MMP-12 KO in rats and genetic deletion of MMP-12 in mice causes major alterations in the expression of other MMPs and that nature of this alteration is different in rats from mice (Chelluboina et al., 2015b; Nalamolu et al., 2018). TIMP-1 has been shown to inhibit MMP-9 and is implicated in post-stroke BBB preservation (Fujimoto et al., 2008). Therefore, it is possible to target MMPs directly, as well as indirectly via TIMP activity modulation for therapeutic intervention.

7. Conclusion and future therapeutic perspective

Stroke is a cerebrovascular disease that affects millions of people every year. Immuno-inflammatory mechanisms leading to ischemic damage are still not fully understood, but accumulating evidence suggests that inflammation is a key element in stroke pathogenesis. Activation of immune cells releases many inflammatory cytokines and critical inflammatory mediators that are not only harmful but also exhibits beneficial effect during inflammation, and fibrosis after stroke. How the temporal expression patterns and polarization into different immune cell subsets affects their function after stroke should be taken into special consideration. The role of DAMPS and NLRP3 signals and recognition of extracellular matrix proteins and the intracellular molecular switches could be promising therapeutic approaches for post-stroke inflammation and progression to fibrosis.

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Highlights:

- Stroke is a disorder of blood vessels accompanied with the detrimental interaction between glia, neurons, vascular cells, and matrix components.
- Illustrates immune cell infiltration, cytokines response in stroke
- Discusses fibrosis ECM deposition, MMPs in stroke
- Elucidates the neuroprotective role of perlecan domain V in stroke via its angiogenesis property
- Summarizes cytokine inhibitors/inducers, immune cell, and inflammation signaling inhibitors, and other pharmacological therapies for stroke.

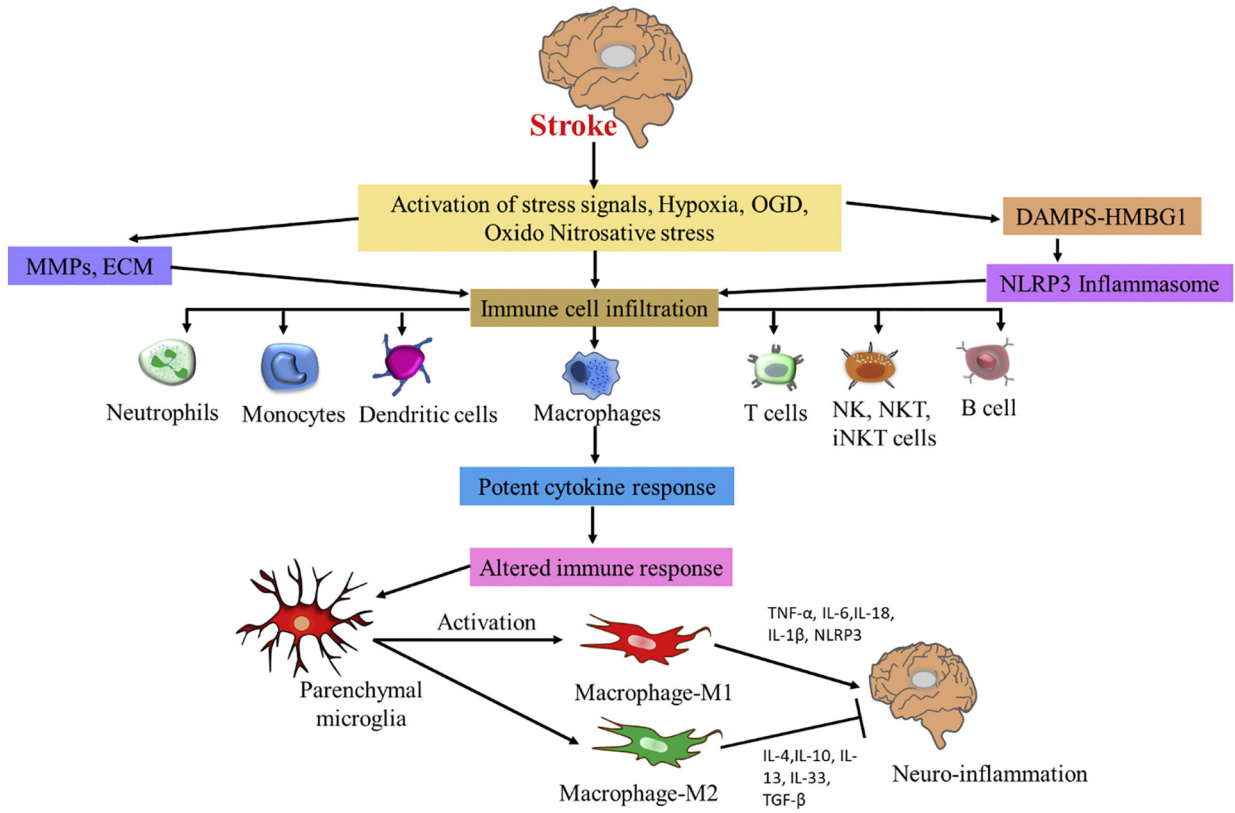


Fig.1. Schematic of stroke-induced neuroinflammation. Stroke induces activation of stress signals, hypoxia, OGD, oxido-nitrosative stress, DAMPS such as high-mobility group box 1 protein (HMGB-1) derived from dying neurons, and it further activates innate microglia and other immune cells, via mediating toll like receptors (TLR), which in turn produce many pro-inflammatory cytokines, and matrix metalloproteinases (MMPs), which leads to activation of NLRP3 signaling and infiltration of immune cells and potent cytokine response and neuroinflammation. Following stroke, activated parenchymal microglial cells transforms to M1 and M2 macrophage polarization. M1 macrophages release proinflammatory cytokines whereas M2 macrophages release anti-inflammatory cytokines and inhibit neuroinflammation.

Table 1

Immune cell inhibitors/inducers for stroke therapy.

Immune cells	Immune cell inhibitors/inducers	Study organism	Experimental study	Biological effects	Reference
Microglia	Pexidartinib (PLX3397, 290 mg in 1 kg chow)	CS7BL/6 J and Cx3Cr1GFP/+ mice	fMCAO (60 min)	↑ in Cx3Cr1 ^{GFP/+} , – (no change) in BBB injury and ↑ in brain injury with ↑ in TNFα, IL-6, MCP-1, KC (CXCL1), 1L-4	(Szalay et al., 2016)
Macrophages	Anti-CCR2 monoclonal antibody MC-21 (IP)	Male C57BL/6 J and B6SJL (CD45.1), CX3CR1-EGFP (CD45.2), (β-actin-GFP + C57BL/C mice	fMCAO	↑ in MDMs, TGFβ, CD163, Yml,	(Wattananit et al., 2016)
	DHA sodium salt (10 mg/kg, IP)	CS7BL/6 mice (both male and female, 8–12 wk)	tMCAO (60 min)	↓ infiltration of peripheral macrophages (CD11b + CD45 ^{high} Ly6G ⁻), neutrophils (CD11b + CD45 ^{high} Ly6G ⁺), B lymphocytes (CD19 + CD3 ⁻), T lymphocytes (CD3 + CD19 ⁻), and ↑ in CD206 + Ibal ⁺ , 1L-10, Arginase-1, TGFβ	(Cai et al., 2018)
T cells (Treg)	Monoclonal antibody CD28SA (300 μg, IP)	Male C57BL/6 J mice (8–12wk)	pMCAO, fMCAO (60 min)	↑ in IL-10, Treg cells and CD45, and ↓ CD11b ⁺ , MHC on macrophages/dendritic cells.	(Na et al., 2015)
B cells	Rituximab (Micromedex, 100 μg IP)	Transgenic human CD20 ⁺ expressing (hCD20 +/–) mice	tMCAO	↓ infarct volumes and didn't alter neurogenesis	(Ortega et al., 2020)
Dendritic cells	Transgenic GFP + overexpress sTNFR1 cells (2 × 10 ⁶ , IV)	Adult male SD	tMCAO	↓ TNF-α and infarct volumes	(Works et al., 2013)

Table Abbreviations, ↑: Increase; ↓: Decrease; –: No change; fMCAO: Filamentous Middle Cerebral Artery Occlusion; BBB: Blood Brain Barrier; TNFα: Tumor Necrosis Factor Alpha; IL: Interleukin; MCP: Mast Cell Protease; MDMs: Monocyte-derived macrophages; TGF-β: Transforming Growth Factor Beta; CD: Cluster Of Differentiation; tMCAO: Transient Middle Cerebral Artery Occlusion; Treg: Infiltrating Regulatory T Cells; IP: Intraperitoneally; pMCAO: Permanent Middle Cerebral Artery Occlusion; MHC: major histocompatibility complex; GFP: Green Fluorescent Protein; sTNFR1: Soluble Tumor Necrosis Factor Receptor-1; IV: Intravenously; wk.: Week; SD: Sprague-Dawley; hCD-20: Human Anti-CD20; mo: Months; CXCL: C-X-C Motif Chemokine Ligand.

Table 2

Cytokine inhibitors/inducers for stroke therapy.

Cytokine	Cytokine Inhibitor/inducer	Study organism	Experimental study	Biological effects	Reference
IL-1 α	IL-1 α (50 pg/ μ L LA)	Male C57BL/6 mice	CCA/M CAO (60 min) followed by reperfusion for 7 days	IL-1 α reduced infarct volumes via \downarrow IL-1 β , IL-6, or CXCL-1 levels and \uparrow expression of PECAM, ICAM-1, and VEGFR2	(Salmeron et al., 2019)
IL-1Ra	IL-1Ra overexpressing mice	CS7BL/6-Tg (UBC-GFP)30Scha/J (Stock No. 004353) (GFP-TG) breeding pairs	tMCAO (40 min)	\uparrow IL-1Ra and \downarrow expression of IL-1 β produced by microglia mediating MAPK signaling in the ischemic cortex	(Clausen et al., 2016)
IL-1 β	Monoclonal anti-IL-1 β antibody (10 μ g/g IV)	JunD siRNA (siJunD)-treated mice with CS7BL/6 J WT (wild type) background mice	tMCAO (45 min)	\downarrow in infarct volume. – (no change) in IL-6, TNF- α , and 4-hydroxynonenal levels	(Diaz-Canestro et al., 2019)
IL-4	IL-4 KO	CS7BL/6 J WT,	tMCAO (60 min)	\downarrow in smi32/MBP ratio	(Zhang et al., 2019)
IL-10	Recombinant IL-10	SD rats (8 wk)	tMCAO (90 min)	\downarrow in TNF- α , IL-1 β , IL-6	(Nakajima et al., 2017)
	ICV IL-10	CS7BL/6 J (10–12 wk)	pMCAO	\downarrow in PD-L1, CXCL9 RE	(Liesz et al., 2014)
IL-13	Recombinant IL-13	Male (4-mo) BALB/cOlaHsd mice	MCAO	– (no change) in GFAP, Ibal, \downarrow in C.D45 $^{+}$, \uparrow in Ibal $^{+}$ /Arg1 $^{+}$	(Kolosowska et al., 2019)
IL-33	Recombinant IL-33	Male C57BL/6 mice (8–10 wk)	CCA/MCA (30 min)	\downarrow in IFN- γ + T cells, \uparrow in Foxp3 $^{+}$ T cells, IL-4, IL-10, TGF- β in spleen tissues	(Xiao et al., 2019)

Table Abbreviations, \uparrow : Increase; \downarrow : Decrease; –: No change; IL-1 α : Interleukin 1 alpha; CCA/MCAO: Central Carotid Artery/ Middle Cerebral Artery Occlusion; Intracerebroventricular; I A: Intra Arterial; IP, Intraperitoneally; IV: Intravenously; min: Minutes; hr Hours; mo: Month; wk.: Week; Yr: year, IL: Interleukin; CXCL: C-X-C Motif Chemokine Ligand; PECAM-1: Platelet/endothelial cell adhesion molecule-1; ICAM-1: Intercellular Adhesion Molecule 1; VEGFR-2: Vascular endothelial growth factor receptor 2; IL-1Ra: Interleukin-1 Receptor Antagonist; MAPK: Mitogen-activated protein kinase; TNF α : Tumor Necrosis Factor Alpha; WT: Wild Type; GFP: Green Fluorescent Protein; SD: Sprague-Dawley; KO: Knock Out; tMCAO: Transient Middle Cerebral Artery Occlusion; IL-1 β : Interleukin 1 Beta; siRNA: Small Interfering RNA; SMI32: a marker of demyelinated axons; MBP: major myelin protein; ICV: Intracerebroventricular. pMCAO: Permanent Middle Cerebral Artery Occlusion; GFAP: Glial fibrillary acidic protein; Interferon gamma; FOXP3:Forkhad Box P3; TGF- β : Transforming Growth Factor Beta.

Table 3

Other pharmacological therapies with anti-inflammatory activity for stroke therapy.

Cell types	inhibitors	Study organism	Experimental study	Biological effects	Reference
Microglia	Melatonin (20 mg/kg. IP)	Male C57BL/6 J mice (8–10 wk)	dMCAO	STAT3 mediated pathway with ↓ In pSTAT3. CD11b. CD86, iNOS, TNF- α . IL-6. and IL-1 β and ↑ in CD206. Arg-1. Ym1/2, TGF- β . IL-10	(Liu et al., 2019)
	Berberine hydrochloride. BP1108 (50 mg/kg/day)	Male C57BL/6 mice	tMCAO (45 min)	AMPK mediated signaling with ↓ in IL-1 β . CD32. TNF- α . and iNOS, p-AMPK and ↑ in CD206. Arg-1. IL-10, Ym1/2	(Zhu et al., 2019)
	α -lipoid acid (50 mg/kg)	Male SD	MCAO (30 min)	↓ in Iba-1-TNF- α -specific cells	(Wu et al., 2016)
Neurons	Fluoxetine (40 mg/kg. IP)	C57B1V6 J mice (3 mo.)	t.MCAO (1 h)/24hR. Drug administered 1 h and 12 h after tMCAO	Fluoxetine was inhibiting IL-10. Bax. and p53 and ↑anti-apoptotic protein Bcl-2 level.	(Shan et al., 2016)
	JNK inhibitor SP 600125 and the ET-1 antagonist BQ 123	Yorkshire pigs (1.1–1.6 kg; 2–7 days old)	Photothrombotic stroke	↓ in IL-6 levels	(Armstead et al., 2019)
Neutrophils	rtPA (10 mg/kg. femoral vein) and NLRP3 shRNA	SD rats	Thromboembolic focal cerebral ischemia model	PPARY/SIRT6/FoxO3a mediated pathway with ↑ in PPAR γ . SIRT6 and ↓ in p-FoxO3a. IL-6, IL-1 β . TNF- α ; inhibited the microglia activation and neutrophils infiltration	(Guo et al., 2018b)

Table Abbreviations, ↑: Increase; ↓: Decrease; -: No change; IL-1 α : Interleukin 1 alpha; Intracerebroventricular. IA: Intra Arterial; IP: Intraperitoneally; IV: Intravenously; min: Minutes; hr: Hours; mo: Month; wk.: Week; Yr: year; IL Interleukin; dMCAO: Distal middle cerebral artery occlusion; STAT3: Signal transducer and activator of transcription 3; pSTAT3: Phospho- signal transducer and activator of transcription 3; iNOS: Inducible nitric oxide synthase; CD: Cluster Of Differentiation; IL-1 β : Interleukin 1 Beta; TNF: tumor necrosis factor, TGF- β : Transforming Growth Factor Beta; p-AMPK: Phospho-AMPK;PPARs: Peroxisome proliferator- activated receptors; SIRT6: Sirtuin 6; FOXO: Forkhead box class O; Arg1: Arginase-1; tMCAO: Transient Middle Cerebral Artery Occlusion; JNK: c-Jun N-terminal kinase; ET-1: Endothelin 1; ET-1: Endothelin 1; PPAR γ : Peroxisome Proliferator-Activated Receptor Gamma; NLRP3:NLR Family Pyrin Domain Containing 3; FOXP3:Forkhead Box P3; Sirt6:Sirtuin 6; Iba1: Ionized calcium binding adaptor molecule 1.