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Jun Chen
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Non-Neuronal Mechanisms of Brain Damage and Repair After Stroke

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Non-Neuronal Mechanisms of Brain Damage and Repair After Stroke

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Introduction: Nonneuronal Mechanisms and Targets for Stroke

For several decades now, clinically effective neuroprotection has been an elusive goal. Although much progress has been made in terms of dissecting molecular pathways and cellular mechanisms, true translation has not succeeded for patients suffering from stroke, brain trauma, and neurodegeneration. Excitotoxicity, oxidative stress, and programmed cell death all represent logical targets for preventing neuronal demise. But it is now increasingly apparent that saving neurons alone may not be enough.

Based on these challenges, the neurovascular unit was proposed as a conceptual framework for reassessing neuroprotection, the fundamental premise being that central nervous system (CNS) function is not solely based on neuronal activity. The brain is more than just action potentials! For neurotransmission to work, release–reuptake kinetics must be coordinated between neurons and astrocytes. For myelinated signals to connect different brain networks, axons need to be in constant homeostatic communication with oligodendrocytes. For the blood–brain barrier to be manifested, crosstalk is required between glial endfeet and cerebral endothelium. Altogether, CNS function is based on cell–cell signaling between multiple cells. Therefore, neuroprotection requires one to do much more than just prevent neuronal death. Rescuing function and cell–cell crosstalk between all cell types in neuronal, glial, and vascular compartments should be required. It is in this context that this monograph *Nonneuronal Mechanisms of Brain Damage and Repair After Stroke* represents a significant addition to the literature and field.

This monograph is divided into five well-integrated sections. The first section focuses on microvascular integrity. Chapters here include analyses of the structural biology of tight junctions, the role of pericytes, glial regulation of barrier function, blood–brain barrier damage in neonatal stroke, and a reconsideration of angiogenesis after stroke. The second section covers the complex actions of glial cells and includes chapters on astrocyte protection, biphasic effects of microglia, and crosstalk between cerebral endothelium and oligodendrocyte precursor cells. The third section then goes on to examine the multifactorial pathways in stroke neuroinflammation, with analyses of peripheral immune activators, monocyte/macrophage responses, T cells, B cells, mast cells and neutrophils, and the web of cytokines that

all contribute to stroke pathophysiology. A critical part of stroke that is relatively less investigated comprises white matter response, and this is the focus of the fourth section of the monograph. In this section, chapters are devoted to assessing the age dependence of white matter injury and subsequently investigating the role of oligodendrogenesis for white matter plasticity. Finally, the last collection of chapters builds on the mechanistic themes explored thus far to develop potential therapeutic approaches. In this final section, chapters span a comprehensive range, including the targeting of leukocyte–endothelial interactions, methods to repair the entire neurovascular unit, immune-based treatments, and cell-based therapies that all seek to achieve neuroprotection by restoring crosstalk amongst the nonneuronal population of CNS cells.

Taken together, the chapters here represent the very best in cutting-edge hypotheses and translational ideas. The mechanisms dissected herein may eventually lead us to testable targets for stroke patients. Curated by editors and authors who are experts in their field, this is an impressive collection of stroke science.

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Part I
Microvascular Integrity in Stroke

Structural Alterations to the Endothelial Tight Junction Complex During Stroke

Anuska V. Andjelkovic and Richard F. Keep

1 Introduction

Cerebral endothelial cells and their linking tight junctions (TJs) form the blood–brain barrier (BBB) [1]. That blood/brain interface regulates the movement of compounds and cells into and out of brain. The BBB controls nutrient supply, aids in the removal of potential neurotoxic compounds from brain, and is an essential component regulating the composition of brain extracellular environment which is vital for normal neuronal function [1].

Stroke, including ischemic and hemorrhagic forms, causes BBB dysfunction [2–4]. Thus, stroke causes increased BBB permeability to blood-borne molecules, cerebral edema formation, and leukocyte infiltration. Such changes may enhance stroke-induced brain injury and worsen stroke outcome. This chapter describes the effects of stroke on the cerebral endothelium and the impact of those changes on brain injury. It examines the underlying mechanisms and potential therapeutic

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approaches to reduce stroke-induced BBB dysfunction. It particularly focuses on alterations in endothelial TJs, but it also addresses the potential role of enhanced endothelial transcytosis after stroke.

2 Normal BBB Structure and Function.

In contrast to systemic capillaries, brain capillary endothelial cells are linked by TJs and have a low basal rate of transcytosis (Fig. 1; [1, 5]). These characteristics limit the para- and transcellular pathways across the endothelium resulting in a very low permeability to many compounds. Thus, for example, the transendothelial electrical resistance, a measure of ionic impermeability, is orders of magnitude greater in cerebral compared to systemic capillaries [6, 7]. Exceptions to such low permeability are compounds which can diffuse across the endothelial cell membrane (e.g., O₂, CO₂, H₂O and molecules with high lipophilicity) or those with specific BBB influx transporters, such as D-glucose [1]. There are also an array of efflux transporters, such as *p*-glycoprotein, that enhance brain to blood transport [1]. Such transporters are involved in preventing potential neurotoxins from entering the brain and clearing metabolites from brain. They are a significant obstacle to the delivery of therapeutic agents for neurological disorders [1]. The BBB also has an array of enzymes that can degrade neuroactive compounds preventing their entry from blood into brain (i.e., it is also a metabolic barrier [1, 5]).

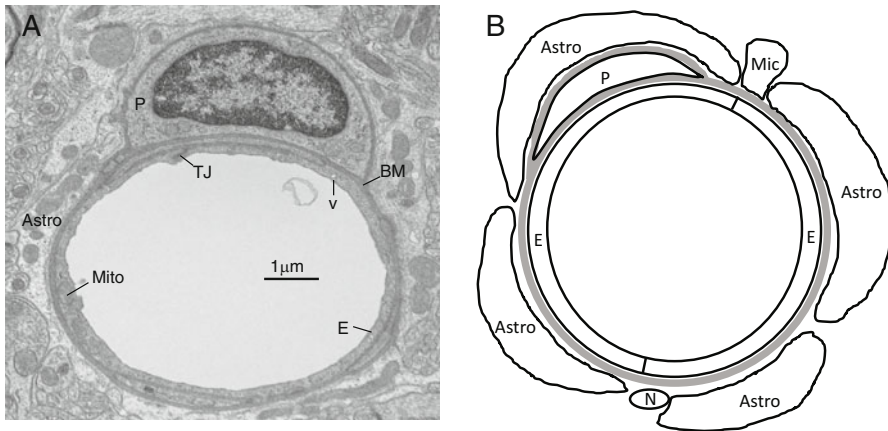


Fig. 1 (a) An electron micrograph showing the structure of a normal mouse cerebral capillary. (b) A schematic showing the relationship between the endothelium and other elements of the neurovascular unit. Brain capillary endothelial cells (E) are linked by tight junctions (TJ), show few vesicles (v) and have more mitochondria (mito) than systemic capillaries. Sharing the same basement membrane (bm) as the endothelium are pericytes (P). Capillaries are surrounded by astrocyte endfeet (astro), with occasional neuronal (N) and microglial processes (Mic). The endothelial cells and the ensheathing cells/basement membrane are called the neurovascular unit

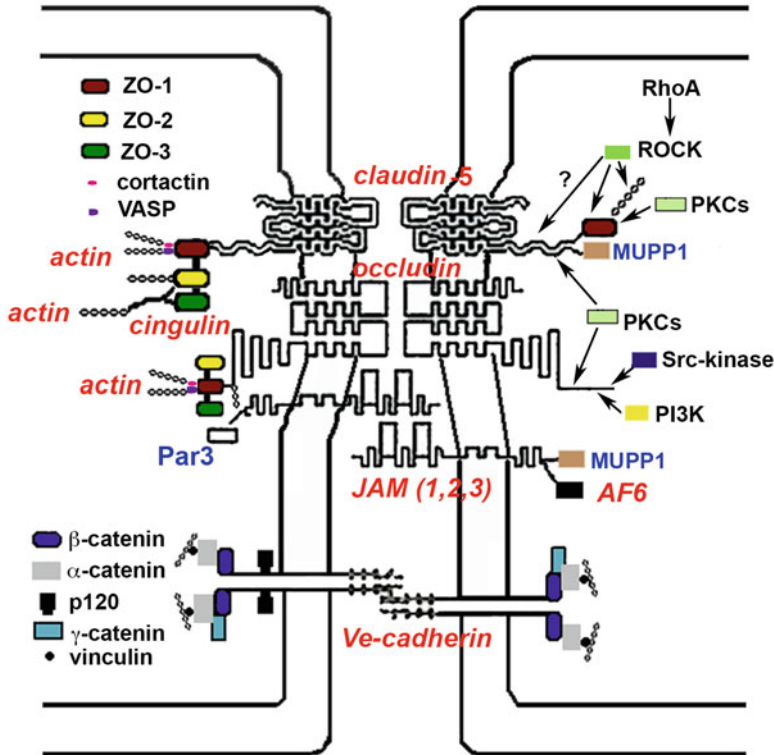


Fig. 2 Schematic representation of the basic components of the brain endothelial junctional complex. The *left* of the panel shows potential structural components of the TJ complex. Claudin-5, occludin, and the JAMs are transmembrane proteins that link adjacent cells. Claudin-5 is the major occlusive protein at the BBB while the cytoplasmic plaque protein, ZO-1, is important for clustering of claudin-5 and occludin and establishing connection with actin filaments. The roles of ZO-2, ZO-3, MUPP1, Par3, Af6 are less clear in brain endothelial cells. Brain endothelial cells also possess adherens junctions, containing the transmembrane protein, Ve-cadherin, and cytosolic accessory proteins (e.g., catenins). The *right* of the panel shows signaling molecules involved in modulating TJ complex assembly and disassembly

Endothelial TJs are integral for BBB function. They are comprised of transmembrane proteins, which form the links between adjacent endothelial cells, and cytoplasmic plaque proteins that form a physical scaffold for the TJs and regulate TJ function [1, 8–10]. In addition, there are links between TJs and the actin cytoskeleton and the adherens junctions that are important for TJ stability and formation (Fig. 2) [9, 10]. TJs are dynamic structures with, for example, claudin-5 and occludin having half-lives of 70–90 min and 6 h, respectively [8].

Transmembrane proteins: These include the claudins, occludin, and the junctional associated molecules (JAMs). In addition, tricellulin, a molecule with homology to occludin, is present at points of three cell contact. The transmembrane

proteins can form *trans*-interactions with proteins on other cells or *cis*-interactions within the same plasma membrane [8].

The claudins are a 27 gene family which are involved in closing the paracellular space between cells (e.g., claudin-5) and in forming ion pores (e.g., claudin-2) [8]. At the BBB, claudin-5 is by far the most predominant claudin although there is evidence of some other claudins (-3, -12, and possibly -1) [8]. Claudin-5 structure and function is regulated by phosphorylation (see below; [8]). It can undergo proteasomal degradation after ubiquitination as well as lysosomal degradation [11]. The claudin-5 knockout results in increased BBB permeability to small molecular weight compounds [12].

Although occludin is not directly involved limiting paracellular permeability, it is thought to be a central regulatory component of the TJ being involved in TJ formation and maintenance [8, 13]. It is a major link to cytoplasmic plaque proteins via binding to ZO-1. Like claudin-5, occludin structure and function is regulated by phosphorylation and it also undergoes ubiquitination and proteasomal degradation [8, 13].

Junctional adhesion molecules (JAM-A, -B, -C) are members of the immunoglobulin superfamily. Their role at the BBB has received much less attention than claudin-5 and occludin, but evidence indicates they are involved in regulating not only paracellular permeability but also leukocyte/endothelial interactions [14–16].

Cytoplasmic plaque proteins: As well as the transmembrane proteins, there are an array of cytosolic proteins associated with the TJs that form the cytoplasmic plaque (Fig. 2). These proteins can be divided on the basis of whether they contain PDZ binding domains. Those which do include members of the membrane-associated guanylate-kinase (MAGUK) superfamily, ZO-1, -2 and -3, as well as Par3, Par6, and AF6 [9]. As many of these proteins contain multiple PDZ domains (e.g., the ZO family members contain three [17]), they can act as scaffolding proteins linking different elements of the TJ (e.g., transmembrane proteins, cytoplasmic plaque proteins, and the actin cytoskeleton).

Cytoplasmic plaque proteins that do not contain PDZ domains include cingulin, 7H6, Rab13, ZONAB, AP-1, PKC ζ , and PKC λ , as well as several G proteins (G α i, G α s, G α 12, G α o). These proteins have multiple functions. Thus, for example, it is thought that cingulin acts as a cross link between TJ transmembrane proteins and the actin-myosin cytoskeleton and that the PKC isoforms and the G proteins regulate TJ assembly and maintenance [9].

Adherens junctions: Cerebral endothelial cells possess adherens junctions (AdJs) as well TJs. AdJs are also a complex of transmembrane (Ve-cadherin) and cytosolic accessory proteins (α , β catenin, p120) closely associated with actin filaments [9, 10]. Crosstalk between AdJs and TJs has been proposed to regulate TJ function [9, 10]. For example, Ve-cadherin upregulates claudin-5 expression in brain endothelial cells [18].

Cell cytoskeleton: The cytoskeleton comprises of actin microfilaments, intermediate filaments, and microtubules. The actin cytoskeleton is linked to the TJs via cytoplasmic plaque proteins such as ZO-1 and to AdJs via catenins (Fig. 2 [9]). Increasing evidence indicates that the actin cytoskeleton is a major regulator of TJ

function [19]. Changes in that cytoskeleton may impact TJ function by altering the physical support (scaffold) for the junction and by transmitting physical forces to the junction proteins.

Neurovascular unit (NVU): Although the cerebral endothelial cells and their linking TJs are the ultimate determinants of BBB permeability, perivascular cells (pericytes, astrocytes, neurons, microglia) have a large role in regulating that permeability. In particular, pericytes, which share the same basement membrane as the endothelium, and astrocytes, whose endfeet almost completely surrounds cerebral capillaries (Fig. 1), both regulate barrier permeability [20–22]. These cells, together with smooth muscle cells in larger vessels, act in concert to regulate cerebrovascular function (blood flow and barrier function), forming the NVU.

3 Alterations in BBB Function After Stroke

The hallmarks of ischemic and hemorrhagic stroke include an increased influx of compounds, such as plasma proteins (Fig. 3), into brain from blood, brain swelling due to a net movement of fluid from blood to brain (brain edema) and brain leukocyte infiltration [4, 23–25]. BBB dysfunction participates in all three of these processes.

Ischemic and hemorrhagic stroke cause increased BBB permeability to both small and large molecules [4, 23–25]. The degree and time course BBB disruption varies depending on the severity of the injury, the location of the tissue being sampled relative to the injury and whether or not there is reperfusion. The BBB disruption can be biphasic [24]. In animal models, BBB barrier disruption reaches a peak

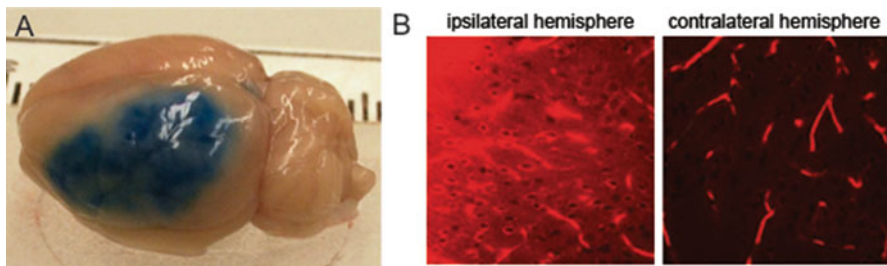


Fig. 3 (a) An example of BBB disruption after rat focal cerebral ischemia at the macroscopic level. A hyperglycemic rat underwent 2 h of middle cerebral artery (MCA) occlusion followed by 2 h of reperfusion. The rat was injected intravenously with Evans blue, which binds to albumin in the bloodstream and is excluded from normal brain by the BBB. In the ischemic MCA territory there was marked extravasation of the Evans blue. (b) An example of BBB disruption after mouse focal cerebral ischemia at the microscopic level. A mouse underwent 30 min of MCA occlusion followed by 24 h of reperfusion. Enhanced dextran-Texas red (40 kDa) leakage through the disrupted BBB was detected by confocal laser scanning micrographs of the brain ipsi- and contralateral to the occlusion. Note that the dextran was confined to the blood vessels in the contralateral tissue but had spread into the parenchyma in the ischemic tissue

~3–7 days after stroke and then gradually resolves [26]. However, there is evidence that there can be a chronic low level of BBB dysfunction after stroke [27–29]. BBB disruption may result in vasogenic edema and contribute to neuroinflammation. Thus, for example, an influx of prothrombin across the damaged BBB results in the production of thrombin in the brain which is pro-inflammatory and can cause brain edema [30]. Similarly, the entry of fibrinogen from blood to brain is a pro-inflammatory signal [31]. One potential beneficial effect of BBB dysfunction is that it may allow greater entry of therapeutics.

Brain edema is a major consequence of stroke which can result in increased intracranial pressure (ICP) and brain herniation. It is a major cause of morbidity and mortality in ischemic and hemorrhagic stroke [32, 33]. Brain edema is classically classified as cytotoxic or vasogenic dependent on whether the underlying cause is injury to parenchymal cells or the cerebrovasculature [33, 34]. It should be noted, however, that in stroke there is injury to both (mixed edema) and that in both cytotoxic and vasogenic there is a net influx of water from blood to brain across the BBB.

Stroke results in leukocyte (neutrophils, macrophages, and lymphocytes) infiltration into brain [35, 36]. That involves a stepwise process with leukocyte:endothelium interactions causing rolling, adhesion and then diapedesis of the leukocyte across the endothelium [35]. The expression of adhesion molecules on the endothelial luminal membrane plays a prominent role in that process. Leukocytes have been implicated in having a detrimental role in stroke, but they are also involved in tissue repair [35, 36].

The effects of stroke on other barrier functions, such as transport and enzyme activity, have received much less attention, although such changes might have important consequences (e.g., for edema formation and drug delivery) [37, 38]. In addition, although the cerebral and systemic vasculatures possess different properties, they share hemostatic properties that are very important in the occurrence and response to ischemia and hemorrhage [39].

4 Alterations to TJ Structure After Stroke

Alterations in TJ function may result from many different processes, including TJ protein modification (e.g., phosphorylation), altered TJ protein/protein interaction, TJ protein relocation, TJ protein degradation and alterations in transcription/translation (Figs. 4 and 5). There is evidence for each of these processes occurring in stroke and they are intimately linked (e.g., phosphorylation may lead to relocation and degradation). The relative importance of each of the processes likely depends on stroke severity and time point.

Protein modifications: Alterations in the phosphorylation state of TJs proteins are crucial factors influencing BBB permeability. Nevertheless, there is controversy due to the actions of different kinases on distinct residues on the same TJ proteins [40–43]. Most evidence on the effects of phosphorylation/dephosphorylation is on three TJ proteins, the transmembrane proteins occludin and claudin-5, and the scaffolding protein ZO-1.

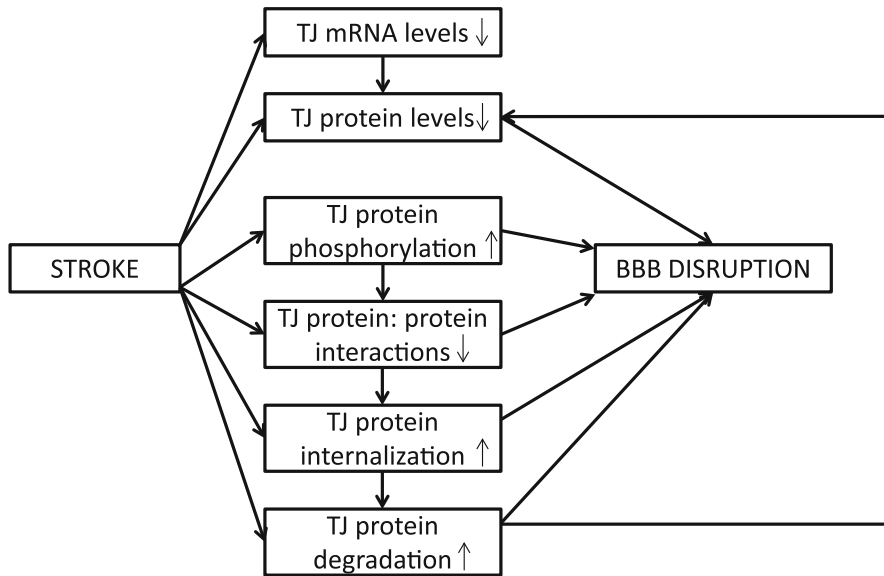


Fig. 4 Schematic showing changes occurring in TJ proteins after stroke that participate in BBB hyperpermeability. Although a potential link between stroke-induced TJ protein phosphorylation, loss of TJ protein interactions, TJ protein internalization and degradation is shown, stroke may affect those processes by several mechanisms, e.g., it may also affect TJ protein degradation by MMP activation

Occludin has several Ser, Thr, and Tyr phosphorylation sites on the C-terminus [40, 44, 45]. Elevated Ser/Thr phosphorylation of occludin is associated with BBB dysfunction during inflammation in the encephalitic human brain [46]. Particularly important findings on the role of Ser/Thr phosphorylation status and barrier function have come from the analysis of vascular endothelial growth factor (VEGF)-induced barrier disruption in retinal endothelial cells, a mechanism underlying diabetic retinopathy [47]. VEGF was shown to induce occludin phosphorylation at sites Thr-168, Thr-404, Sr-408, Ser-471, and Ser-490 via PKC β , with Ser-490 impacting occludin/ZO-1 interaction and leading to occludin ubiquitination and degradation [47, 48]. In stroke, direct evidence regarding occludin phosphorylation on these Ser/Thr residues is still lacking. There is though evidence that protein kinase C (PKC) isozymes (nPKC- θ and aPKC- ζ) are activated during hypoxia/reoxygenation injury and that cPKC α and Rho-kinase activation is induced by the chemokine CCL2, a major driver of leukocyte entry during stroke-induced brain injury [49, 50].

In epithelial cells, occludin Tyr residues are normally minimally phosphorylated and their phosphorylation is associated with barrier disruption [51]. Tyr-398 and Tyr-402 on occludin are involved in the interaction with ZO-1 and their phosphorylation destabilizes barrier function [44]. At the cerebral endothelium, focal cerebral ischemia, and glutamate treatment *in vitro* induces occludin Tyr phosphorylation. A Src-kinase inhibitor, PP2, reduces the occludin Tyr phosphorylation and BBB dysfunction found during ischemia/reperfusion (I/R) injury [52–54].

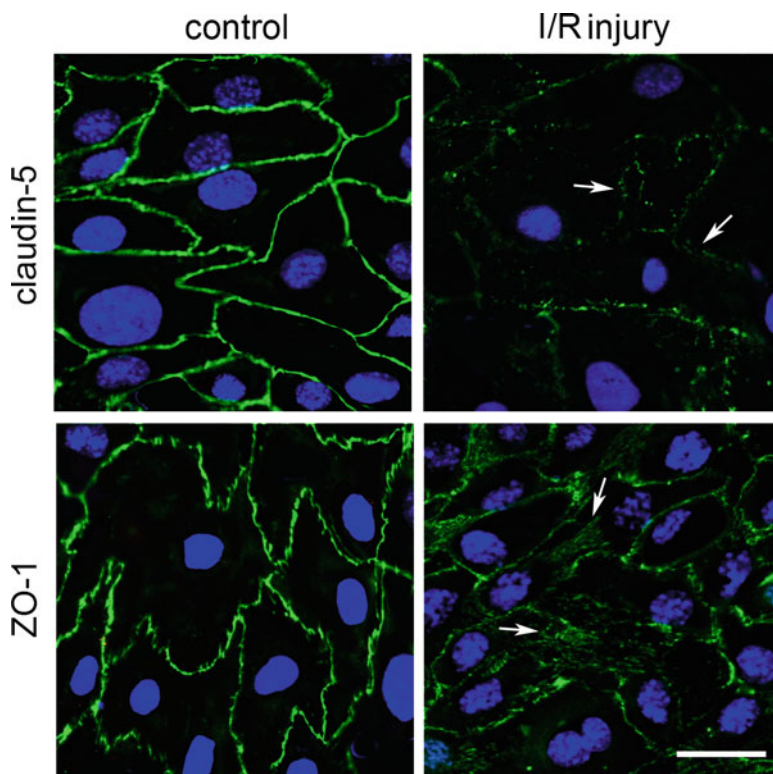


Fig. 5 Mouse brain endothelial cells were exposed to oxygen–glucose deprivation (OGD/ischemia) for 5 h followed by normal oxygen and glucose condition (reperfusion) for 1 h as a model of ischemia/reperfusion (I/R) injury. Control represents cells exposed to normal oxygen and glucose. Immunofluorescent staining was used to visualize the localization of claudin-5 and ZO-1 under the control and I/R conditions. Notice the alteration in claudin-5 and ZO-1 localization/expression on the lateral border of the brain endothelial cells after I/R injury (*arrows*). Scale bar = 20 μ m

Phosphorylation of claudin-5 on the Thr-207 residue in the C-terminal domain by PKA or Rho kinase generally affects TJ integrity in brain endothelial cells and increases barrier permeability [46, 55]. However, direct evidence on the pattern of claudin-5 phosphorylation in stroke is still lacking. The activation of nPKC- θ , aPKC- ζ , cPKC- α cPKC- β , PI3K γ , p38MAPK in different phases of stroke injury and correlation with alterations in claudin-5 localization and expression indirectly point to an essential role of Ser/Thr phosphorylation of claudin-5 in TJ complex disassembly [49, 50, 56, 57].

Evidence indicates that ZO-1 phosphorylation regulates barrier permeability in different systems [58–60]. VEGF increase pTyr-ZO-1 levels in retinal endothelial cells and causes barrier dysfunction [61]. In inflammation, ZO-1 phosphorylation by PKC- ϵ on threonine 770/772 causes endothelial barrier disruption [62]. Several recent studies also pinpoint that in brain endothelial cell models, cytokine-induced

injury (TNF- α , IL-6 or CCL2) significantly induces Tyr, Thr, and Ser phosphorylation of ZO-1 [43, 49].

The phosphorylation of TJ proteins is tightly associated with the “status” of protein/protein interactions in the junctional complex. Claudin-5/ZO-1, occludin/ZO-1, ZO-1/JAM-A, and ZO-1/actin cytoskeleton interactions are considered essential for the localization and stability of the TJ complex as well as for establishing the trans-interactions and adhesion properties of claudin-5 between adjacent cells [8, 9]. Thus, occludin phosphorylation on Thr 424/Thr438 by PKC ζ is required for TJ assembly in epithelial cells, while phosphorylation on Tyr (Tyr398 and Tyr402) and Ser residues (Ser490) attenuates interaction with ZO-1 and promotes dislocation from the lateral membrane in oxidative stress-induced barrier alterations [45, 46, 48, 63, 64]. A similar effect is also described for occludin phosphorylation on Ser408, with dissociation from ZO-1 and increased paracellular permeability [45]. In transient cerebral ischemia, occludin interaction with ZO-1 is diminished due to increased Tyr phosphorylation of occludin and inhibiting Src-kinase (inhibitor PP2) reduces BBB dysfunction [54].

The C-terminus of claudin-5 contains a PDZ binding motif for direct binding to ZO-1, ZO-2, and ZO-3 [65, 66]. Direct evidence regarding diminished interaction between claudin-5 and ZO-1 in the TJ complex during pathological conditions such as ischemia/reoxygenation is lacking although the disappearance of claudin-5 from the cell boundary and increased Ser and Thr phosphorylation suggest diminished interaction with ZO-1 in TJ complex disassembly [50, 56, 57, 67].

Relocation: Besides phosphorylation and diminished protein/protein interactions, disassembly of TJ complex is associated with redistribution of transmembrane TJ proteins. Both occludin and claudin-5 are internalized in inflammatory as well as ischemic conditions [68, 69]. Redistribution of claudin-5 in early brain ischemic conditions is mediated by caveolae [68]. CCL2, a chemokine increased in brain after I/R injury, also causes claudin-5 and occludin by caveolae-mediated internalization [69]. On the other hand, occludin may also be redistributed by clathrin-dependent pathways described in retinal endothelial cells [47]. In contrast, another transmembrane TJ protein, JAM-A, shows macropinocytotic-dependent redistribution under inflammatory conditions [16]. It is relocated from the lateral to the apical side of brain endothelial cells during I/R injury and BBB disruption, as well as during inflammation [15, 16]. JAM-A contains LFA binding sites in the C2 domain and has a role in leukocyte adhesion and diapedesis. Thus, the pattern of JAM-A relocation during TJ disassembly is closely associated with its role in leukocyte infiltration [15, 16].

Degradation: Many studies have described a loss of TJ immunostaining after stroke (e.g., [70–75]). This may result from in situ degradation or internalization of the proteins followed by degradation by the proteasome or lysosomes. The most extensively studied in situ degradation is via matrix metalloproteinases (MMPs). MMP-2 and -9 have been shown to degrade occludin and claudin-5 after cerebral ischemia [68, 75]. Intracellular proteasomal and lysosomal degradation of TJ proteins can occur after ubiquitination, with polyubiquitination generally favoring proteasomal degradation and monoubiquitination lysosomal [11]. With cerebral

ischemia there is evidence of ubiquitination of occludin at the BBB by an E3 ubiquitin ligase, Itch, followed by degradation [76]. Similarly, with retinal ischemia, occludin undergoes polyubiquitination after phosphorylation at the blood retinal barrier [77].

TJ mRNA expression: As well as degradation, loss of TJ proteins after stroke may result from decreased transcription or translation. Reductions in claudin-5, occludin, and ZO-1 mRNA levels have been described after ischemic stroke [67, 75, 78].

Oligomerization and the redox response of TJ proteins: Occludin and claudin-5 have cysteine residues in their cytosolic C-termini that are involved in oligomerization through disulfide bridge formation [40–42]. By building disulfide bridge formation occludin, for example, may become increasingly oligomerized [79]. Such bridges are redox-dependent with normoxic conditions supporting occludin oligomerization and TJ assembly, while hypoxia-reoxygenation results in TJ disruption [79].

5 Signaling Mechanisms Underlying Alterations in TJ Structure in Stroke

Direct and indirect effects of stroke on the endothelium. The effects of ischemic stroke on the endothelial TJs, and thus BBB disruption, may be directly on the endothelium or indirectly via effects on other cell types within the NVU or infiltrating leukocytes which then signal to the endothelium. It has been difficult to assess the relative importance of direct endothelial cell effects in vivo, but the generation of endothelial-specific knockout and transgenic mice will help to examine this question. That the endothelium can be directly affected by ischemia is demonstrated by in vitro experiments on brain endothelial monocultures using oxygen glucose deprivation (OGD) to partially mimic ischemia (Fig. 5). OGD with and without reoxygenation causes disruption of the barrier formed by brain endothelial cell monolayers [68, 80, 81] and this is associated with marked changes in endothelial cell TJs (e.g., relocation and/or loss of TJ proteins [68, 81]).

One of the major components of ischemic and hemorrhagic injury is inflammation with an infiltration of leukocytes into brain, the activation of resident microglia and the production of inflammatory mediators. Inflammation and inflammatory mediators have a major impact on the brain endothelial TJ complex. Thus, different cytokines (IL-1 β , TNF- α), chemokines (IL-8, CCL2), MMPs (MMP-2, MMP-9), adhesion molecules (ICAM-1), transcription factors (Nf κ B), Poly(ADP-ribose) polymerase-1 (PARP) affect TJ structure and function either directly or indirectly (by attracting leukocytes which bind to the endothelium and also produce inflammatory mediators) (e.g., [81–84]). Inflammation and inflammatory mediators are strong modulators of TJ complex disassembly and BBB disruption after cerebral ischemia (e.g., [81, 83, 85, 86]). Long-term changes in the TJ complex and BBB permeability after stroke may fuel alterations in the NVU towards development of chronic inflammatory foci which may lead to further disruption of BBB and progression of inflammation and injury [27–29].

Apart from direct effects of stroke on the cerebral endothelium and indirect effects due to inflammation, ischemic and hemorrhagic stroke may impact the endothelium by other effects on the NVU (e.g., altered trophic support). The reader is referred to other chapters in this book concerning the relationships between cells of the NVU.

Protein kinase C: Hypoxic, ischemic, and I/R injury trigger a variety of signaling pathways involved in regulating TJs and barrier permeability. Among the most extensively studied TJ regulators is the PKC family of serine/threonine kinases. They regulate a variety of cell functions including proliferation, gene expression, cell cycle, differentiation, cytoskeletal organization cell migration, and apoptosis [87–89]. The PKC family includes many different isozymes that are involved in signal transduction from membrane receptors to the nucleus. The activation of cPKC- β II, nPKC- θ , PKC- ζ , PKC- α , and PKC- μ during hypoxia and post-hypoxic reoxygenation or under influence of endothelin-1 or CCL2 regulate TJ disassembly [50, 90–95].

Protein tyrosine kinases: The protein tyrosine kinases (PTKs) are another set of signaling molecules essential for regulating BBB integrity. There are two main classes of PTKs: receptor PTKs and cellular, or non-receptor, PTKs [96–98]. The receptor PTKs have extracellular domains comprised of one or more identifiable structural motifs. They are involved in the BBB effects of growth factors such as VEGF and PDGF [99, 100]. For example, Flt-1 activation and activation of signaling molecules associated with intracellular domain of receptor tyrosine kinases (phosphatidylinositol 3-kinase/Akt (PI3-K/Akt) nitric oxide synthases (NOS) and protein kinase G (PKG)) mediate hypoxia/VEGF-induced brain endothelial hyperpermeability and brain edema formation [101, 102].

The cellular PTKs (SRC, JAK, ABL, FAK, FPS, CSK, SYK, and BTK) are involved in some of critical events in TJ assembly and disassembly. For example, the Src-family kinases can modulate tyrosine phosphorylation on occludin attenuating its interaction with ZO-1, ZO-2, and ZO-3. They also regulate MMP activity and occludin degradation as well as caveolin-1 phosphorylation and activation and, thus, relocalization of TJ proteins, particularly during I/R-induced BBB disruption [52–54, 68, 103]. Some cellular PTKs, the JAKs, phosphorylate members of the signal transducer and activator of transcription family (the JAK/STAT pathway). That pathway is important in regulating claudin-5, ZO-1 and inflammation in the brain endothelial cells exposed to HIV [104]. PTKs are a major target of reactive oxygen species (ROS) [105]. Thus, for example, ROS generated during ischemia, brain injury, monocyte/neutrophil activation or alcohol exposure, can activate MMPs (-1, -2, and -9) and decrease tissue inhibitors of MMPs (TIMP-1 and -2) in a PTK-dependent manner [68, 106, 107].

MAP kinases. Besides tyrosine kinases, mediators of oxidative stress can also activate a complex system of MAP kinases (p38, Erk1/2) and regulate brain endothelial permeability [57, 108–110]. MAP kinases may be a nodal point in, for example, PTK signaling but they may also directly phosphorylate TJ proteins as occurs during exposure to 4-hydroxy-2-nonenal (4-HNE), one of the major biologically active aldehydes formed during inflammation and oxidative stress [110].

Rho GTPases: Another important regulator of TJs and paracellular permeability is the Rho family of GTPases. That family regulate TJs both indirectly, via effects on the actin cytoskeleton, and directly. Rho, GEF-H1 a guanine nucleotide exchange factor for Rho, Rac, and ROCK regulate cytoskeletal contractile responses via myosin ATPase activity [111]. ROCK, for example, promotes phosphorylation of the regulatory light-chain of myosin (MLC) on Ser19 and Thr18 through phosphorylation of the myosin light-chain phosphatase (MLCP) and, thus, blocks MLC dephosphorylation [112, 113]. This site-specific phosphorylation of MLC in turn elevates myosin ATPase activity, leading to actin-myosin contraction [111, 113]. Besides the specific actin filament polymerization (stress fiber formation), which generate in turn contractile intraendothelial forces impacting the TJs, all of these factors may also induce phosphorylation of TJ and AdJ proteins and their redistribution [112]. These data indicate that Rho/ROCK not only causes remodeling of actin cytoskeleton but also directly or indirectly via activation other signal pathways induces phosphorylation and remodeling of the TJ and AdJ complexes [70, 114]. In addition, ROCK is involved in regulating the expression of the proteolytic enzymes, such as MMP-9 and urokinase-type plasminogen activator (uPA), that impact BBB permeability [115]. Recent evidence has highlighted the beneficial effect of ROCK inhibition, with fasudil or Y2763, in treating BBB disruption in stroke and post-stroke conditions [70, 116–118].

Cerebral hemorrhage. In contrast to the signaling events occurring in the cerebral endothelium during cerebral ischemia, little is known about the events after cerebral hemorrhage [2]. A component of ICH-induced brain injury is related to clot-derived factors such as thrombin, hemoglobin, and iron [23]. The direct impact of such factors on brain endothelial cell signaling and TJ function in relation to ICH is generally understudied. However, there is evidence that thrombin can disassemble claudin-5 from endothelial TJs [119], that hemoglobin exposure can reduce claudin-5 and ZO-1 levels at the BBB [120] and that iron participates in the loss of BBB occludin and ZO-1 after cerebral ischemia [121]. Thus, it is very likely that clot-derived factors impact BBB TJs after hemorrhagic stroke.

6 Non-TJ Pathways Involved in BBB Hyperpermeability After Stroke

Although there is evidence for a role of TJ alterations in BBB disruption after stroke, two other pathways have also been implicated, increased transcytosis at the cerebral endothelium and endothelial cell death [122, 123] (Fig. 6). While cerebral endothelial cells have very few vesicles compared to systemic capillaries, there is some transcytosis under basal conditions. Such receptor-mediated transcytosis, via clathrin- and caveolin-1-coated vesicles, is important for the uptake of a number of macromolecules into brain (e.g., transferrin and leptin, [1, 124, 125]). In addition, some heavily cationized proteins undergo adsorptive-mediated transcytosis [1]. After ischemic stroke there is a marked increase in the number of vesicles in the

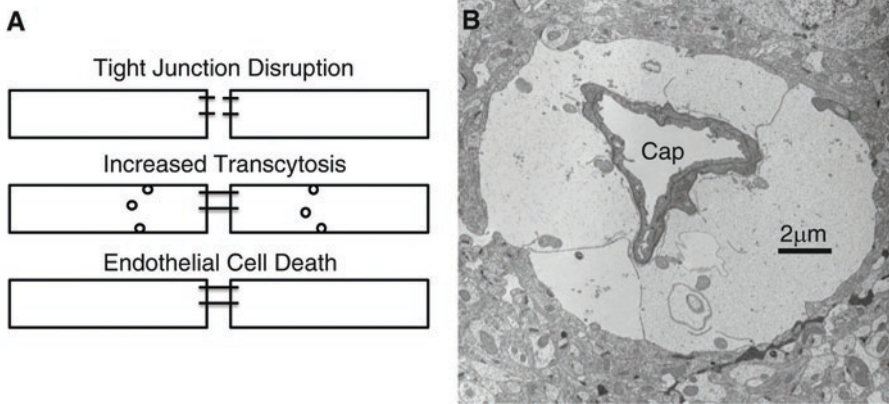


Fig. 6 (a) Routes proposed to participate in BBB disruption after stroke. Note that these routes are not mutually exclusive. (b) Electron micrograph of a cerebral capillary (cap) from the ischemic penumbra in a mouse that underwent 3 h of middle cerebral artery occlusion. Note the marked perivascular edema in the absence of marked changes in the rest of the parenchyma (compare to Fig. 1a)

cerebral endothelium [122, 126, 127]. If the increase in vesicles reflects increased transcytosis, this may contribute to increased BBB permeability after stroke. This may be particularly important for large molecular weight compounds as this pathway would be relatively size independent. Knowland et al. [122] recently proposed that such an increase in transcytosis is responsible for BBB disruption to albumin (but not a smaller tracer, Biocytin-TMR) observed 6 h after ischemic stroke, whereas profound TJ changes are responsible for later disruption at 2 days. Mice without caveolin-1, an essential component of caveolae, had reduced levels of endothelial- and parenchymal-associated albumin after stroke. It should be noted that caveolae are also involved in internalization of claudin-5 and occludin [69] and, therefore, the caveolin-1 knockout may have also affected TJ function after stroke although Knowland et al. did not identify such an effect [122].

One method to elucidate the relative importance of the transcellular (vesicular) and paracellular routes involved in BBB hyperpermeability is examining size selectivity. Preston and Webster [128] examined the relative permeability of two tracers (sucrose and inulin) of different molecular weights after cerebral ischemia and found that the absolute increase in permeability was greater for the smaller compound. They concluded that the results fitted with diffusion through a pore (with some steric hindrance) rather than vesicular transport. However, it should be noted that those experiments were 24–72 h after the ischemic event.

It has also recently been suggested that endothelial cell death is the primary cause of BBB disruption after cerebral ischemia [123]. Such endothelial cell death has been noted by other investigators after cerebral ischemia [129, 130]. A question that arises with a loss of endothelial cells being the route of BBB disruption is the extent to which such vessels continue to remain patent. Endothelial damage results in the formation of a platelet plug and activation of the coagulation cascade.

7 Therapeutic Interventions

Despite decades of work on developing neuroprotective agents for stroke [131] only reperfusion-based therapy has so far shown clinical efficacy. Tissue plasminogen activator [132] and now mechanical thrombectomy [133] improve outcomes in ischemic stroke. Those approaches do not directly target BBB changes after stroke. Indeed, tPA is associated with increased risk of cerebral hemorrhage after ischemic stroke [132]. There have been preclinical [134, 135] and now clinical trials (NCT02222714 and the European trial, Imatinib Treatment in Acute Ischemic Stroke) of agents that might protect the cerebral vasculature if given in combination with tPA. Reducing the occurrence of ICH would alleviate one impediment to its use in stroke patients. It might also increase the time window over which tPA can be given.

In preclinical models, many agents have been shown to decrease BBB hyperpermeability and alterations in TJ structure after ischemic stroke (e.g., [70–75]). A question arises as to whether those BBB effects are direct or indirect (e.g., via reducing parenchymal damage)? Endothelial-specific genetic manipulations should address this question and also whether BBB damage may contribute to parenchymal damage. It should be noted that if the BBB is a therapeutic target, drug delivery to the BBB avoids many of the problems with delivery to the brain parenchyma (e.g., brain penetration).

8 Conclusions

BBB disruption occurs in both ischemic and hemorrhagic stroke. It may contribute to brain injury by enhancing edema formation, leukocyte influx, and entry of neurotoxic compounds from blood to brain. While endothelial transcytosis and cell death have been proposed as being important in BBB dysfunction after stroke, considerable evidence indicates a major role for TJ modification. The extent to which the BBB is a primary therapeutic target for limiting brain injury, with and without reperfusion therapy, still needs to be fully clarified, but the tools necessary for such studies are becoming available.

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Role of Pericytes in Neurovascular Unit and Stroke

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Abbreviations

ACA	Anterior cerebral artery
BBB	Blood–brain barrier
CADASIL	Cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy
CNS	Central nervous system
MCA	Middle cerebral artery
NVU	Neurovascular unit
PA	Penetrating arteries

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PDGFR β	Platelet-derived growth factor receptor- β
PDGF β	Platelet-derived growth factor- β
RBCs	Erythrocytes = red blood cells
SMCs	Smooth muscle cells
tPA	Tissue plasminogen activator
α SMA	Smooth muscle α -actin

1 Neurovascular Unit and Stroke

The neurovascular unit (NVU) is composed of the endothelia, pericytes, basal membrane ensheathing them, astrocyte end-feet around the vessels and, neurons innervating the intraparenchymal vasculature (Fig. 1a). It plays an integrating role in matching the metabolic demand with the blood flow in addition to regulating the development and maintenance of the blood–brain barrier (BBB) [1–5], leukocyte trafficking across the BBB [6–9], and angiogenesis [10–12]. Ischemic injury to the NVU unfavorably impacts the stroke-induced damage and brain edema by

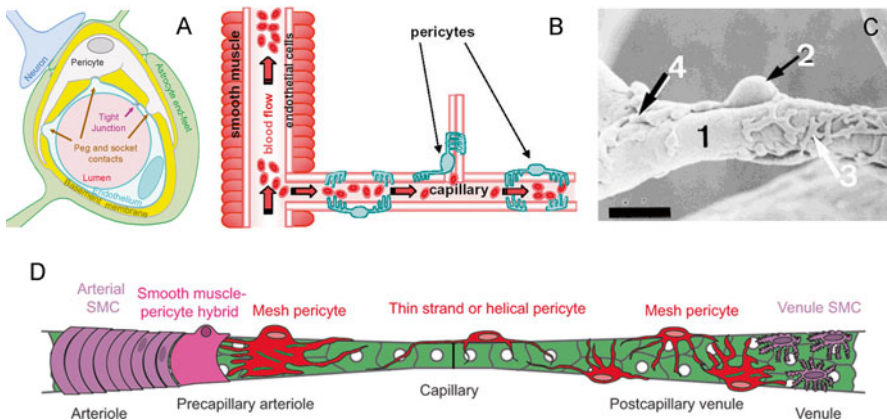


Fig. 1 Neurovascular unit and pericytes. (a) The neurovascular unit is composed of the endothelia and tight junctions between them, pericytes, the basal lamina ensheathing endothelia and pericytes, and astrocyte end-feet surrounding the microvessel. Note the peg and socket type contacts between endothelia and pericytes. (b) Potential blood flow control sites in cerebral vasculature: smooth muscles on arterioles and pericytes on capillaries (Reproduced from Peppiatt et al., 2006 with permission). (c) Scanning micrograph of a vascular cast of a cortical capillary (1) with a pericyte-like structure (2) having primary and secondary processes (3) distributed around the vascular cast and the capillary branching points (4). Scale bar = 11.5 μ m. (Reproduced from Rodriguez-Baeza et al., 1998 with permission). (d) Pericyte processes are highly varied with shapes ranging from thin singular strands that run parallel to the microvasculature to more complex mesh processes that enwrap the entire vessel lumen. Pericytes located closer to the arteriolar end of the microcirculation exhibit more circular processes that may be essential to their contractile function (Reproduced from Hartmann et al., 2015 with permission)

disrupting microvascular blood flow and BBB integrity, whereas ischemia-triggered signaling in the NVU of vasculature within the peri-infarct area positively impacts stroke outcome by promoting post-stroke angiogenesis and neurogenesis [13, 14].

2 Pericytes

First described by Rouget in 1873 [15], pericytes were later named by Zimmerman in 1923 based on their prominent location at the abluminal wall of microvessels [16]. However, their importance within the NVU has not been recognized until recently. Pericytes communicate with other cells of the NVU and contribute to the control of several microcirculatory functions such as neurovascular coupling, maintenance of the BBB and basal lamina, regulation of the angiogenesis, immune cell entry to the central nervous system (CNS) and scar formation [3, 11, 17–21]. They also function as pluripotent stem cells after injury to the NVU [22]. Although pericytes are also present on peripheral microvessels, the density of pericytes on microvessels is highest in the CNS and retina in accordance with their role in focal regulation of the microcirculatory blood flow and maintenance of the blood–brain/retina barrier [2, 21, 23, 24].

Pericytes are located on pre-capillary arterioles, capillaries, and post-capillary venules [2, 25, 26] (Fig. 1b–d). Pericytes are present on straight parts as well as branching points of the capillaries; 56 % of the pericyte somas are located at a capillary junction [27]. Unlike smooth muscle cells (SMCs), pericytes are embedded within two layers of basement membrane [24]. They form peg and socket type contacts and gap junctions with the endothelia through the basement membrane, facilitating the communication between pericytes and endothelia [28, 29] (Fig. 1a). Adjoining membranes of the neighboring pericytes are interconnected with gap junctions, which probably serve as a conduit for transmitting messages along the microvascular wall [19, 30]. Pericytes extend processes along and around the microvessels, which cover 30–90 % of the microvessel wall in the CNS [4, 23, 31, 32]. Processes are more circumferential at the arteriole side of the microvascular bed and at branching points, more longitudinal in the middle of the capillary bed, and have a stellate morphology at the venule end of the microcirculation (Fig. 1d). Junctional pericytes extend processes at least a cell width away from the junction [27]. It has long been recognized that pericyte morphology varies along the course of microvasculature, presumably to accommodate differing functions [25, 27, 33]. Not only their morphology, but also their protein expression varies [2, 34]. Several transitional forms are observed along the vascular bed at various developmental stages or after pathological stimuli [2, 25, 26, 35]. The transition from smooth muscle α -actin (α SMA) expressing SMCs to pericytes is also not sharp. Smooth muscle–pericyte “hybrid” cells precede the prearteriolar pericytes having mesh-like circular processes [26, 27]. Pericytes that give out more circumferential processes express more α SMA, when assessed either with immunohistochemistry of brain sections *ex vivo* [27, 34] or in mice cortex expressing reporter dyes under control of the

α SMA promoter in vivo [33]. It should be noted that immunohistochemistry directly detects the α SMA protein (mainly in the cytoplasm of the soma and processes), whereas reporter dyes expressed under the control of α SMA promoter are membrane-bound, therefore, basically label the pericyte membrane. Mid-capillary pericytes do also express α SMA [36]. However, the detection of the small pool of α SMA in their relatively short processes by immunohistochemistry requires rapid fixation before α SMA depolymerization, whereas low level of α SMA expression may be difficult to visualize due to dispersion of the limited amount of reporter fluorescent protein expressed over the large surface area of the pericyte membrane.

Pericytes can be identified by a number of proteins that they express. Unfortunately, the antibodies against to majority of them either detect only a subpopulation of pericytes or do not have the desired specificity to unambiguously distinguish pericytes from other cell types [2, 25]. Recent studies with transgenic mice expressing fluorescent reporter proteins under the control of the growth factor platelet-derived growth factor receptor- β (PDGFR β) and of the proteoglycan NG2, suggest that PDGFR β detects a larger population of pericytes compared to NG2 and, interestingly, CD13 immunohistochemistry is reportedly to match the performance of fluorescent reporters expressed under the control of PDGFR β promoter [27, 33, 37].

2.1 Regulation of Microcirculatory Blood Flow and BBB by Pericytes

Pericytes are considered contractile cells since their discovery [15, 16]. However, this view was challenged in the past few decades based on the premise that the blood flow increase evoked by neuronal activity was mediated by relaxation of SMCs around arterioles as well as by failure of detection of α SMA in capillary pericytes in some studies. As reviewed in detail by Díaz-Flores et al. [12], the pericyte contractility is supported by several lines of evidence including their characteristic morphology with processes that envelop the microvessels as well as ultrastructural and immunohistochemical demonstration of contractile proteins [24, 34, 36, 38–45] in addition to the presence of receptors for vasoactive mediators on their surface [19, 30, 46]. In vitro studies on cerebellar, cerebral, and retinal slices or on isolated microvessels and in vivo studies have clearly disclosed that pericytes contract or dilate in response to vasoactive mediators applied [30, 46, 47]. A recent in vivo study showed that cortical capillaries dilated before arterioles during sensory stimulation, supporting the view that microvascular blood flow in the CNS is regulated by pericytes in response to the very focal demand originating from a small group of nearby cells as a final step of flow regulation after the arterioles, which serve a larger cohort of cells [18]. This flow regulation with fine spatial resolution may be essential for tissues with high functional specialization such as the brain and retina. However, it should be noted that all microvascular pericytes are not contractile and proportion of the contractile ones may vary with the tissue, species, and developmental stage as well as along the arteriovenous axis as noted above [20, 33, 48].

It has also been shown that a close interaction between the endothelia and pericytes as well as astrocytes is required for development and functioning of the NVU and BBB [4, 49]. Pericytes promote the formation of tight junctions and inhibit transendothelial vesicular transport [4, 49, 50]. The number of pericytes per endothelial cell and the surface area of the vascular wall covered by pericytes determine the relative permeability of capillaries. Accordingly, pericyte dysfunction or deficiency causes increased BBB permeability [2, 4, 21, 49].

2.2 Pericytes Are Vulnerable to Ischemic Injury

The pericyte contractility is regulated by intracellular Ca^{2+} concentration as in vascular SMCs surrounding the upstream vessels [19, 46, 51]. Accordingly, these highly dynamic cells bear the risk of Ca^{2+} overload when they are unable to maintain low intracellular Ca^{2+} concentrations. The energy insufficiency is a well-characterized cause of loss of the intracellular Ca^{2+} equilibrium, which relies on several energy demanding processes. In addition, factors such as reactive oxygen species (ROS) may also contribute to an uncontrolled rise in intracellular Ca^{2+} [52, 53]. Pericytes have large elongated mitochondria, which follow the central core of the pericyte longitudinally, and may be a significant source of ROS under pathological conditions [31]. NADPH oxidase, a major superoxide-producing enzyme is highly expressed in brain pericytes [54, 55]. Indeed, ROS has been shown to cause a sustained increase in Ca^{2+} in cultured human brain microvascular pericytes [52, 53]. Reactive oxygen and nitrogen species, especially peroxynitrite, have been reported to induce pericyte contraction during focal ischemia/reperfusion in the intact mouse brain [56]. In addition to unregulated Ca^{2+} rise, several other processes such as ATP and thromboxane A₂ released from the ischemic brain or platelets, which are potent constrictors of pericytes, may also contribute to pericyte contraction [48].

3 Changes in Pial and Penetrating Arteries Shortly After Stroke

On occlusion of the middle cerebral artery (MCA), collaterals between the anterior cerebral artery (ACA) and MCA are opened, large arterial branches but especially surface pial network and penetrating arteries (PA) dilate while flow directions are reorganized to sustain the flux rates in PAs as high as possible [57–59]. While these compensatory changes can preserve cell viability at the periphery of the MCA area, creating an opportunity for recovery if MCA recanalization can be attained within a couple of hours, the severe decrease in blood flow velocity, volume, and distal capillary perfusion in the core ischemic area makes infarction unavoidable [58, 60–62] (Fig. 2). In vivo imaging of cerebral circulation in intact mice under anesthesia has unequivocally illustrated that the PAs, especially those with smaller luminal

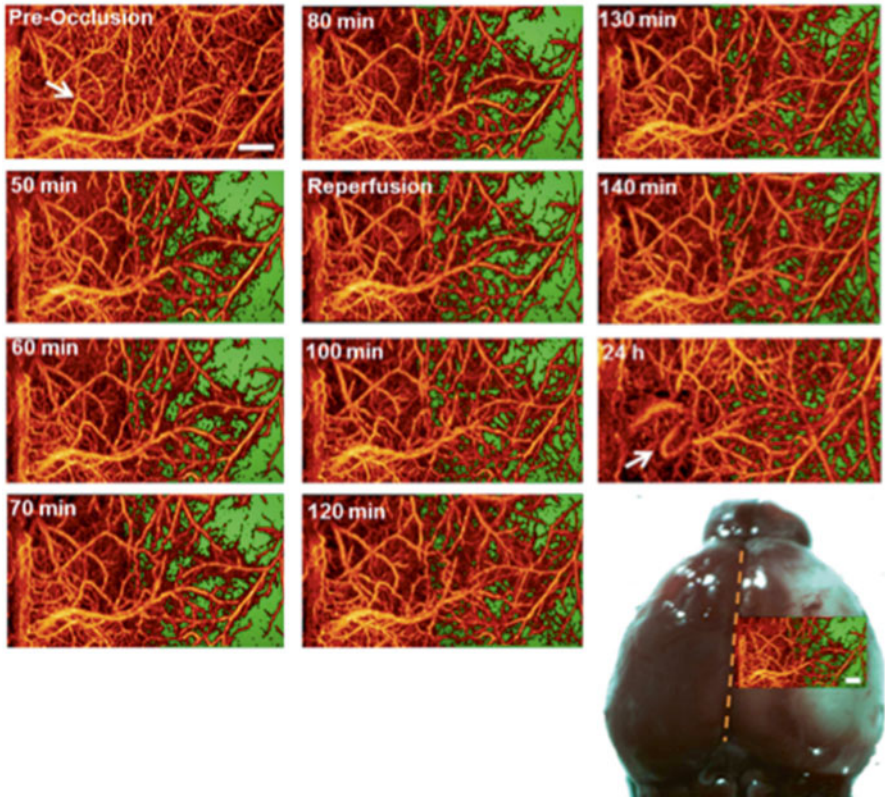


Fig. 2 Incomplete microcirculatory reflow after recanalization. Dynamic imaging of cortical blood flow using optical microangiography during 90-min proximal MCA occlusion followed by recanalization illustrates the lack of microcirculatory blood flow in the MCA territory (the *green area*) during occlusion and its partial recovery after recanalization (incomplete microcirculatory reperfusion) in the mouse. Consecutive images are shown at 10-min intervals. Image size is $2.2 \times 4.4 \text{ mm}^2$. The image in the *lower right* is the optical microangiography image taken at 50 min overlaid on the 24 h infarct analysis by histological staining as the area of pallor. Scale bar = $500 \mu\text{m}$ (Reproduced from Dziennis et al., 2015 with permission)

diameter dilated to compensate for the low perfusion pressure [58, 61]. The magnitude of dilation decreased with the distance from the pial arteriolo-arteriolar anastomoses with sufficient collateral flow and was replaced by constriction in areas further away [58]. Majority of these changes are reversible if recanalization is achieved within a short time. However, when perfusion deficit is prolonged some of these changes are not reversible and may negatively impact the recovery after stroke. For example, in the mouse brain, part of the microcirculatory flow cannot be reinstated after MCA occlusion lasting more than an hour despite complete reopening of the MCA [33, 56].

4 Incomplete Microcirculatory Reflow After Recanalization

An impaired tissue reperfusion due to microvascular constrictions (no-reflow phenomenon) was first noted after global and focal cerebral ischemia more than half a century ago [63, 64]. The emergence of “no-reflow” depends on the duration and severity of ischemia as well as the brain region studied although these variables have not been systematically compared in the setting of focal cerebral ischemia. In the mouse, MCA occlusion induces nodal microvascular constrictions that generally do not recover after recanalization starting 1 h after ischemia and affecting more than half of the microvessels within 2 h [33, 56] (Fig. 2). Narrowed microvessel lumina are filled with entrapped erythrocytes (RBCs), leukocytes, and fibrin-platelet deposits [65–69]. RBCs are the predominant cell types in aggregates as they are the most prevalent cells in circulation. In addition to the constricted segments observed at the arteriolar end of microcirculation and capillaries, leukocytes adhered to post-capillary venules for entering to the parenchyma also induce luminal aggregates together with fibrin and platelets [65, 69–71].

Experimental data strongly suggest that incomplete restoration of the microcirculatory blood flow negatively impacts tissue recovery even if reopening of the occluded artery is achieved when there is still salvageable penumbral tissue. Pharmacological agents and genetic manipulations reducing microvascular clogging by inhibiting leukocyte adherence, platelet activation or fibrin–platelet interactions have been shown to restore microcirculation and improve stroke outcome in animal models [65, 72–75]. Importantly, neuroprotection obtained with some BBB-impermeable agents strongly support the idea that restoring microvascular patency can improve stroke outcome independently of parenchymal mechanisms [56, 76]. Consequently, restitution of the microcirculatory reperfusion emerges as an exciting target to improve the success rate of recanalization therapies.

In the past, microvessel constrictions were thought to be caused by swollen astrocyte end-feet encircling microvessels [66, 67]. Recently, pericytes on microvessels were proposed to play an important role in incomplete microcirculatory reperfusion because they contracted during ischemia and remained contracted despite reopening of the occluded artery [18, 56, 77] (Fig. 3). Although it has been claimed that α SMA expressing microvascular cells with contractile capability should be defined as SMCs [33], the mural cells with a bump-on-a-log morphology located on the abluminal wall of microvessels downstream to arterioles, including their transitional forms to SMCs, are named as pericytes since their original description by Zimmerman [16, 78]. The fact that pericytes are a heterogeneous group of cells sharing some transitional features with SMCs and that some but not all express α SMA have always been a matter of confusion and a source of debate. Nomenclature disagreements notwithstanding the important point for the stroke pathophysiology is that contractile cells on brain microvessels impede reperfusion after ischemia and unfavorably impact the outcome of recanalization. It should be noted that even small decreases in capillary radius caused by subtle pericyte contractions can lead to erythrocyte entrapments because capillary luminal size hardly allows passage of

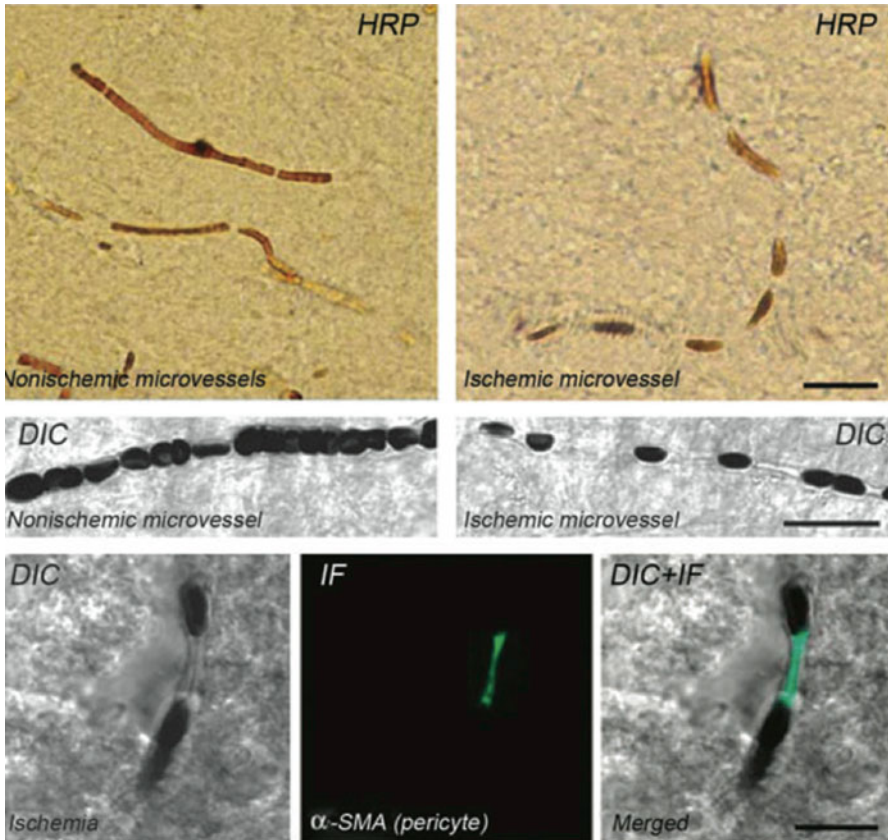


Fig. 3 Ischemia causes persistent pericyte contraction, which is not restored after complete recanalization of the occluded artery. Mice were subjected to 2 h of proximal MCA occlusion and intravenously injected with horseradish peroxidase (HRP) before decapitation, 6 h after reopening of the MCA. HRP-filled microvessels exhibited sausage-like segmental constrictions in ischemic areas on brain sections (*upper row*). The differential interference contrast (DIC) microscopy images illustrate frequent interruptions in the erythrocyte column in an ischemic capillary contrary to a continuous row of erythrocytes flowing through an intact capillary (*middle row*). The constricted segments colocalized with α -smooth muscle actin (α -SMA) immunoreactive pericytes (*bottom row*). IF denotes immunofluorescence. Scale bar for *upper* and *middle row*, 20 μ m; *bottom row* 10 μ m (Reproduced from Yemisci et al., 2009 with permission)

RBCs [19, 56] (Fig. 3 middle row). Entrapped erythrocytes trigger platelet and fibrin aggregation by impeding passage of blood cells [69, 79]. The failure of erythrocyte circulation within some of the microvessels and increased heterogeneity of RBC transit times through patent capillaries (due to varying degrees of capillary resistances) can catastrophically reduce O_2 delivery to the tissue struggling to recover from ischemia-induced perturbations [80]. Since the plasma flow in constricted capillaries is relatively less restricted compared to RBC flux, glucose supply to some parts of the tissue may exceed O_2 supply and stimulate anaerobic glycolysis,

hence, lactic acidosis ([56, 77], please also see supplementary movies 5–7 in [33]). Therefore, ischemia-induced pericyte contractions emerge as a viable target for restoring impaired microcirculatory reperfusion. Indeed, sustained release of adenosine within circulation from nano-assemblies (NA) (adenosine itself has only a few minutes of plasma residence time) has recently been shown to reduce ischemia-induced erythrocyte entrapments and improve microcirculatory reflow by relaxing contracted pericytes after 2 h of MCA occlusion [76] (Fig. 4). Unlike adenosine infusion or synthetic adenosine agonists, slow release from squalenoyl-adenosine NAs did not cause cardiotoxicity or hypotension in the mouse model used.

Since pericytes also play an important role in maintenance of the BBB integrity [4, 21, 49], the ischemia/reperfusion-induced pericyte dysfunction may contribute to BBB leakiness as well. This can be further aggravated by the death of capillary pericytes within 24 h after MCA occlusion as shown in mice and rats [18, 81]. Increased BBB permeability predisposes to intraparenchymal hemorrhage and brain swelling in about 6% of patients receiving i.v. tissue plasminogen activator (tPA) [82]. Diabetic patients are more prone to hemorrhage perhaps due to dysfunctional microvascular pericytes, a well-known cause of diabetic retinopathy [83–85]. Interestingly, pericyte loss is increasingly reported for conditions that are risk factors for stroke, such as ageing, hypertension, and diabetes, the impact of which on stroke outcome needs to be clarified with future research [86]. Among many complex mechanisms, overproduction of oxygen and nitrogen radicals on the microvascular wall appears to contribute to both BBB leakiness and incomplete reflow during cerebral ischemia/reperfusion [56, 87]. Altogether, these findings bring about the exciting possibility that effective suppression of oxidative/nitrative stress during reopening of the occluded artery may improve the outcome of recanalization therapies by promoting microcirculatory reperfusion as well as by preventing hemorrhagic conversion and vasogenic edema [87]. Despite failure of an antioxidant agent in clinical trials, the experimental evidence still warrants pursuit of this goal [88, 89].

5 Clinical Evidence for “No-Reflow” After Recanalization Therapies for Stroke

A short therapeutic time window limits the use of recanalization therapies for majority of stroke patients [90, 91]. This brief therapeutic time window is attributed to rapid loss of neuronal viability in the ischemic penumbra [82, 92]. However, increasing clinical evidence suggests that an incomplete reperfusion plays a critical role in determining tissue survival after successful recanalization [93, 94]. Several recent imaging studies serially analyzing recanalization and reperfusion in ischemic stroke patients report that, on average, 26% of recanalized patients with thrombolytics do not show reperfusion [95]. This incomplete reperfusion is observed after pharmacological (intravenous or intraarterial) as well as interventional recanalization [93, 96–98]. Clinical trials have repeatedly demonstrated that a good outcome was better correlated with reperfusion than recanalization in stroke patients treated

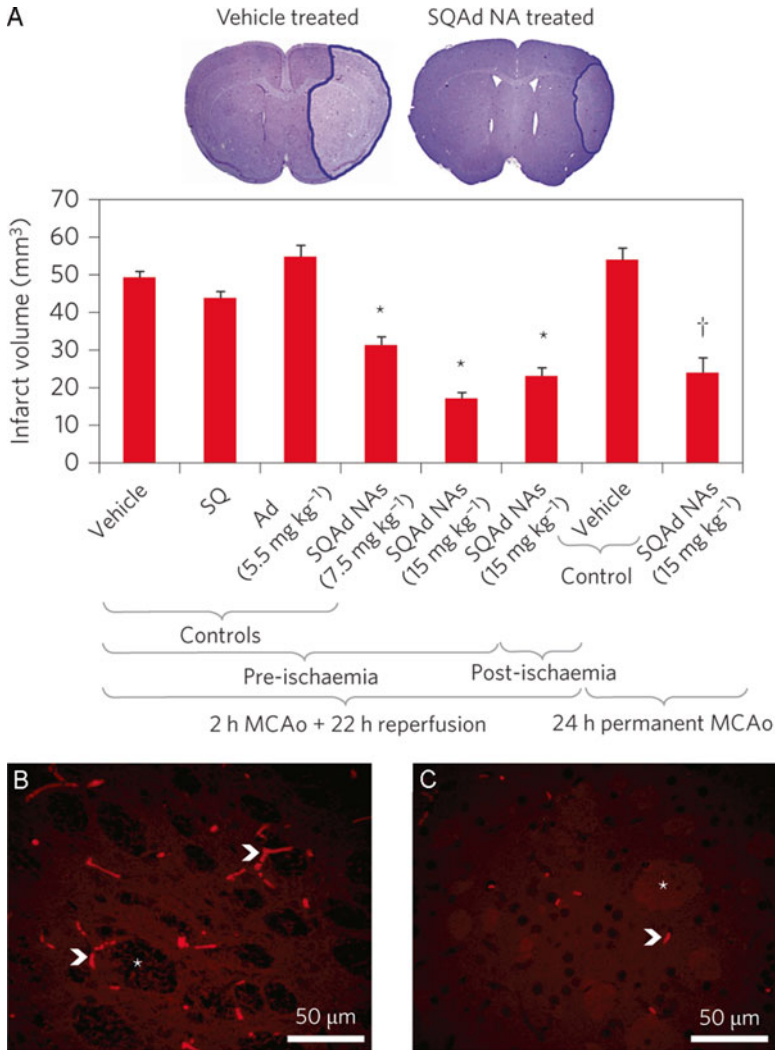


Fig. 4 Systemic administration of squalenoyl-adenosine (SQAd) nano-assemblies (NAs) provides significant neuroprotection in a mouse model of focal cerebral ischaemia. **(A)** Infarct areas in control and treated mice subjected to transient (2 h MCAo and 22 h reperfusion) and permanent (24 h MCAo) focal cerebral ischemia were identified by reduced Nissl staining under a light microscope (magnification $\times 10$, insets) (data are presented as mean (mm³) \pm S.D., $N=6$ animals per group; † and * indicate $P < 0.05$ compared to respective controls). Intravenous administration of 7.5 or 15 mg kg⁻¹ SQAd NAs just before ischemia or 2 h post-ischemia significantly decreased the infarct volume compared with control groups that received vehicle (dextrose 5%), adenosine-unconjugated SQ NAs (9.45 mg kg⁻¹) or free adenosine equivalent to the amount in NAs (5.5 mg kg⁻¹). A significant therapeutic effect was also observed when SQAd NAs were administered 2 h post-ischemia in the permanent MCAo model. **(B, C)** In untreated mice, capillaries in the ischemic brain were filled with trapped erythrocytes, whose hemoglobin was rendered fluorescent by treating brain sections with NaBH₄ **(B, red, arrowheads)** 6 h after reopening of the MCA following 2 h of occlusion, whereas the majority of capillaries were not clogged in SQAd NAs-treated mice **(C)**

with tPA or interventional methods [93, 94, 96, 97, 99, 100]. Recent imaging studies show that increased capillary transit time heterogeneity (a measure of incomplete/impaired reperfusion) is a good predictor of the tissue destined to infarct [101].

6 Role of Pericytes in CADASIL

Cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy (CADASIL) is caused by mutations in the *NOTCH3* gene [102]. The protein encoded by the *NOTCH3* gene is expressed in pericytes as well as vascular SMCs. Recent studies in Notch3 transgenic mice expressing one of the human mutations have disclosed that Notch3 aggregated around microvascular pericytes, leading to pericyte loss or reduced coverage of capillaries by pericyte processes [103, 104]. These changes were associated with a leaky BBB, reduction in communication with endothelial cells and neurovascular dysfunction. Confirming the clinical significance of these findings, pericyte loss was also observed in skin and muscle biopsies of CADASIL patients [105].

7 Post-stroke Angiogenesis and Pericytes

Pericytes are essential especially for the early phase of neovascularization (angiogenic sprouting) [10, 11] (Fig. 5). Pericytes and endothelial cells communicate with each other for regulation of angiogenesis [10–12]. Platelet-derived growth factor- β (PDGF β), transforming growth factor- β , notch, angiopoietin, and sphingosine-1-phosphate signaling mediate this crosstalk [2, 106]. Increasing evidence suggests that pericytes play an important role in post-stroke angiogenesis as well [107–111]. Typically, endothelial cells start to proliferate and give off vessel sprouts 12–24 h after brain ischemia, leading to formation of new vessels in the peri-infarct region 3 days after ischemic injury [107, 112, 113]. Following a similar time course, the PDGFR β expression is upregulated in pericytes, which increase in number and start migrating from the microvessel wall to the newly formed vessel sprouts to foster their maturation after ischemic injury [114–117]. Renner et al. found that PDGFR β increased in pericytes 48 h after permanent ischemia [117]. Similarly, NG2+ or PDGFR β + pericytes reportedly increase in peri-infarct areas 1–3 weeks after transient MCA occlusion [81, 118]. A proportion of locally proliferating pericytes give rise to microglial cells [119]. Interestingly, chronic administration of cilostazol, an antiplatelet drug, has been claimed to promote pericyte proliferation, which might decrease the final infarct size by promoting new vessel formation after naturally occurring stroke in spontaneously hypertensive rats [120]. Corroborating these studies, conditional knockout of PDGF β /PDGFR β signaling in adult mice that have normally developed brain vasculature, led to larger infarcts than controls when subjected to focal cerebral ischemia [121]. Similarly, Zechariah et al. showed that

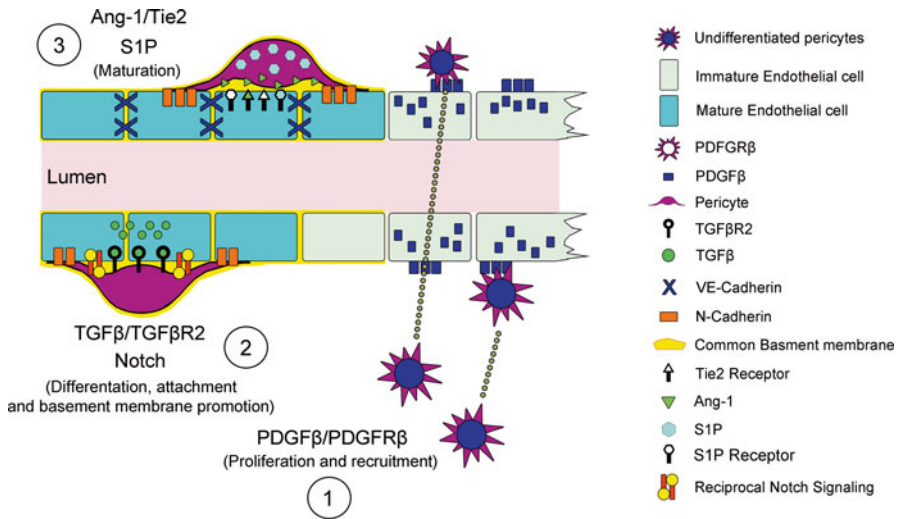


Fig. 5 Role of pericytes in angiogenesis. The interaction between PDGFβ secreted by the endothelium and its receptor localized on pericytes (PDGFRβ) is essential for recruitment of undifferentiated mesenchymal cells/pericytes to newly formed vessels. Once pericytes are at the vascular wall, reciprocal Notch signaling between the endothelia and pericytes as well as interactions between TGFβ secreted by endothelial cells and its receptor TGFβR2 located at pericytes differentiate mural cells and attach them to the newly formed vessels. The TGFβ/TGFβR2 interaction also promotes formation of the common basement membrane and stabilizes newly formed vessels by inhibiting endothelial proliferation. Ang-1, which is secreted by pericytes, activates its endothelial receptor Tie2 and promotes blood–brain barrier formation. Finally, S1P, whose receptor is abundantly expressed on pericytes downregulates genes related to vascular permeability and promotes both endothelial–endothelial (VE-cadherin) and pericyte–endothelial cell (N-cadherin) interconnections

pericytes did not appropriately cover the brain capillaries in hyperlipidemic mice exposed to ischemia and, this was associated with attenuation of post-stroke angiogenesis [111].

Albeit indirectly, further supporting a role for pericytes in post-stroke angiogenesis, intravenous injection of a combination of smooth muscle progenitor cells and endothelial progenitor cells 1 day after MCA occlusion enhanced the angiogenesis and vessel maturation in the peri-infarct areas [122]. Since the adult bone marrow is considered to be a rich reservoir of pericyte progenitor cells, bone marrow-derived pericytes may be involved in post-ischemic angiogenesis [109, 123, 124]. Indeed, Kokovay et al. showed that, following brain ischemia, bone marrow-derived cells with a pericytic phenotype and expressing angiogenic factors were recruited to cerebral capillaries [109].

Angiogenesis is also essential to promote neurogenesis after stroke [122, 125, 126]. In fact, newly formed neurons have been found located near to the remodeled vessels [127], probably because vascular cells recruit and form a niche for neural stem cells [126, 128]. Since pericytes are essential in post-stroke angiogenesis and express factors that can induce neurogenesis as well as angiogenesis, pericytes may

also be involved in post-stroke neurogenesis [25, 129]. In vitro studies have clearly shown that the brain-derived pericytes have a potential to differentiate into neurons in response to trophic factors such as basic fibroblast growth factor, Sox2 and Mash1 [22, 130–132]. Pericytes obtained from ischemic MCA tissue of adult animals or pericytes cultured under ischemic conditions also showed capability to differentiate to cells of neural as well as vascular lineage [133].

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Glial Support of Blood–Brain Barrier Integrity: Molecular Targets for Novel Therapeutic Strategies in Stroke

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Abbreviations

ABC	ATP-binding cassette
ALK	Activin receptor-like kinase
APC	Activated protein C
AQP	Aquaporin
BBB	Blood–brain barrier
BCRP	Breast cancer resistance protein
CNS	Central nervous system
COX	Cyclooxygenase
EAAT	Excitatory amino-acid transporter
EPCR	Endothelial protein C receptor
GSH	Glutathione
GSSG	Glutathione disulfide
H/R	Hypoxia/reperfusion
HMG-CoA	3-Hydroxy-3-methylglutaryl coenzyme A
JAM	Junctional adhesion molecule
Keap1	Kelch-like ECH-associated protein 1
MAGUK	Membrane-associated guanylate kinase
MAPK	Mitogen-activated protein kinase
MCAO	Middle cerebral artery occlusion
MCP	Monocyte chemoattractant protein

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MDR	Multidrug resistance
MMP	Matrix metalloproteinase
MRP	Multidrug resistance protein
NO	Nitric oxide
NOS	Nitric oxide synthase
Nrf2	Nuclear factor E2-related factor-2
NVU	Neurovascular unit
OATP	Organic anion transporting polypeptide
PAR	Proteinase-activated receptor
P-gp	P-glycoprotein
RANTES	Regulated upon activation normal T-cell expressed and secreted
ROS	Reactive oxygen species
r-tPA	Recombinant tissue plasminogen activator
SGLT	Sodium–glucose cotransporter
SLC	Solute carrier
SOD	Superoxide dismutase
TEMPOL	4-Hydroxy-2,2,6,6-tetramethylpiperidine- <i>N</i> -oxyl
TGF	Transforming growth factor
TLR	Toll-like receptor
VEGF	Vascular endothelial growth factor
ZO	Zonula occluden

1 Introduction

The blood–brain barrier (BBB) is an essential barrier system that separates the central nervous system (CNS) from the circulation. It is formed by a monolayer of endothelial cells that limit brain uptake of xenobiotics in an effort to maintain cerebral homeostasis. The BBB is also a significant obstacle to CNS drug delivery. In fact, several currently marketed drugs have limited or no efficacy in the brain due to an inability to cross the BBB and attain therapeutic CNS concentrations [1]. Brain microvascular endothelial cells are not intrinsically capable of forming a “barrier.” Formation of the BBB requires coordinated cell–cell interactions and signaling from adjacent glial cells (i.e., astrocytes, microglia), pericytes, neurons, and extracellular matrix [2]. Such an intricate relationship implies existence of a “neurovascular unit (NVU).” During ischemic stroke, various NVU cell types are triggered by pathological stimuli. Understanding endothelial and glial cell responses that are involved in modifying the NVU/BBB in the context of stroke provides an opportunity not only to protect BBB integrity but also to target these mechanisms for effective CNS drug delivery. In this chapter, we summarize BBB physiology including interactions with associated cell types/structures (i.e., glial cells, pericytes, neurons, extracellular matrix) of the NVU. Additionally, we highlight endothelial and glial cell mechanisms that contribute to BBB/NVU dysfunction. Finally, we provide insights on how such mechanisms can be targeted in an effort to protect BBB integrity and/or optimize CNS drug delivery for stroke pharmacotherapy.

2 The Neurovascular Unit

The NVU consists of multiple CNS cell types/structures including brain microvascular endothelial cells, astrocytes, microglia, pericytes, neurons, and extracellular matrix [3]. The concept of the NVU emphasizes that the brain response to ischemic stroke occurs via coordinated interactions between these cell types [4, 5]. Decreased BBB functional integrity occurs prior to neuronal injury, which suggests that NVU dysfunction is directly linked to the BBB disruption observed in the setting of stroke [6, 7].

2.1 *Components of the Neurovascular Unit*

2.1.1 Endothelial Cells and the Blood–Brain Barrier

CNS function requires precise regulation of the brain extracellular space. Additionally, metabolic demands of brain tissue are considerable and account for approximately 20% of total oxygen consumption [8]. The interface between CNS and systemic circulation must possess mechanisms that can facilitate nutrient transport, exactly regulate ion balance, and provide a barrier to potential toxins. This emphasizes a critical need for both physical and biochemical mechanisms that contribute to BBB barrier properties.

The current understanding of BBB structure is based upon studies by Reese, Karnovsky, and Brightman in the late 1960s [9, 10]. Anatomically, BBB endothelial cells are demarcated by a lack of fenestrations, minimal pinocytotic activity, and presence of tight junctions [11]. Cerebral endothelial cells have elevated mitochondrial content that is required for transport of solutes into and out of the brain thereby contributing to maintenance of CNS homeostasis [12]. Several receptors, ion channels, and influx/efflux transport proteins are expressed in brain microvascular endothelial cells. Transporters are a critical BBB biochemical mechanism that are prominently involved in permitting brain entry of some substances while excluding accumulation of other substances in brain parenchyma [13].

2.2 *Molecular Characteristics of the BBB*

2.2.1 Tight Junction Protein Complexes

BBB endothelial cells are interconnected by tight junctions (Fig. 1), which are large multi-protein complexes maintained by direct contact with astrocytes [14]. Physiologically, tight junctions form a continuous, almost impermeable barrier that limits paracellular diffusion of blood-borne substances with the exception of small, lipid soluble molecules [11]. The high BBB transendothelial resistance (1500–2000 Ω cm²) further restricts free flow of water and solutes [15]. BBB tight

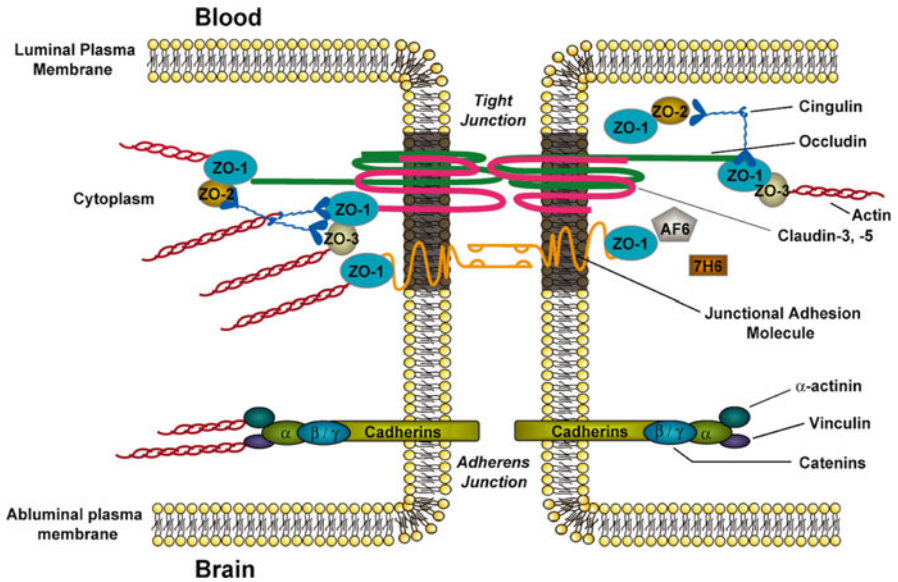


Fig. 1 Basic molecular organization of tight junction protein complexes at the blood–brain barrier. Adapted from Ronaldson & Davis. *Curr Pharm Des.* 18(25): 3624–3644 (2012)

junction formation primarily involves specific transmembrane proteins (i.e., junction adhesion molecules (JAMs), occludin, and claudins (i.e., claudin-1, -3, and -5)) that are linked to cytoskeletal filaments by interactions with accessory proteins (i.e., zonula occluden (ZO)-1, -2, and -3) [16].

Several JAM isoforms have been identified at the mammalian BBB including JAM-1, JAM-2, and JAM-3 [3, 16]. JAMs regulate transendothelial migration of neutrophils and monocytes/macrophages [17, 18]. Loss of JAM protein expression is directly correlated with BBB disruption and injury [19]. Additionally, studies in an immortalized human brain endothelial cell line (hCMEC/d3) showed that inflammatory stimulation led to increased JAM movement away from the tight junction and increased paracellular solute diffusion, which further suggests a central role for JAMs in maintaining BBB integrity [20].

Monomeric occludin is a 60- to 65-kDa protein that is highly expressed along endothelial cell margins in brain vasculature [21, 22]. Our group has shown that occludin is a critical regulator of BBB permeability in vivo [21, 22]. Occludin assembles into dimers and higher-order oligomers, a characteristic that is required for restriction of paracellular permeability. Altered occludin expression is associated with BBB dysfunction in several pathologies including hypoxia/aglycemia [23], hypoxia/reoxygenation stress [22], and focal cerebral ischemia [24].

Claudins are 20- to 24-kDa proteins that contribute to the physiological “seal” of the tight junction [25]. In cerebral microvascular endothelial cells, various claudin isoforms have been detected including claudin-1, -3, and -5 [24, 26–28]. In experimental stroke models [19, 29], altered expression of claudin-5 and an associated

enhancement in paracellular permeability has been reported. Claudin-1 expression was shown to decrease at the *in vivo* BBB following exposure to human amyloid- β (A β_{40}) peptide [30]. A β deposition at the BBB is characteristic of cerebral amyloid angiopathy, a pathological condition that contributes to microvascular injury including hemorrhages and ischemia.

Proper physiological functioning of the BBB, particularly restriction of paracellular solute transport, requires association of JAMs, occludin, and claudins with cytoplasmic accessory proteins. In brain microvascular endothelial cells, membrane-associated guanylate kinase-like (MAGUK) proteins are involved in clustering of tight junction protein complexes to the cell membrane [31]. Three MAGUK proteins have been identified at the tight junction: ZO-1, -2, and -3. ZO-1 links transmembrane tight junction proteins (i.e., JAMs, occludin, claudins) to the actin cytoskeleton [32]. Previous studies have shown that dissociation of ZO-1 from the junction complex is associated with increased permeability, suggesting that the ZO-1-transmembrane protein interaction is critical to tight junction stability and function [33–35]. ZO-1 has been shown to localize to the endothelial cell nucleus under conditions of proliferation or injury [36], following Ca²⁺ depletion [37], and in response to nicotine [27]. Similarly, ZO-2 binds tight junction constituents, signaling molecules and transcription factors [38]. In fact, ZO-2 may act redundantly with ZO-1 as it has been shown to facilitate formation of morphologically intact tight junctions in cultured cells lacking ZO-1 [39]. ZO-3 is expressed at the BBB but its exact role in the formation of tight junctions and/or maintenance of tight junction integrity has not been elucidated [40]. CNS pathologies including hypoxia, cerebral ischemia/reperfusion injury, and focal cerebral ischemia are associated with reduced ZO-1 expression at the BBB [41–43].

2.2.2 Adherens Junctions

Adherens junctions are specialized cell–cell interactions, which are formed by cadherins and associated proteins that are directly linked to actin filaments [44]. Cadherins regulate endothelial function by activation of phosphoinositide 3-kinase signaling, an intracellular pathway that organizes the cytoskeleton and enables complex formation with vascular endothelial growth factor receptor 2. Therefore, cadherin-mediated signaling is essential for endothelial cell layer integrity and for the spatial organization of new microvessels [45]. Barrier-forming endothelium has been shown to express higher levels of cadherin-10 relative to VE-cadherin [46]. In contrast, circumventricular organs and choroid plexus capillaries (i.e., brain microvasculature that is devoid of BBB properties) primarily express VE-cadherin [46]. Optimal cadherin function requires direct association with catenins. At least four catenin isoforms (i.e., β , α , χ , and p120) are expressed at the BBB, with β -catenin linking cadherin to α -catenin, which binds this protein complex to the actin cytoskeleton [47]. *In vitro* studies by Steiner and colleagues demonstrated that the heparin sulfate proteoglycan agrin also contributes to barrier properties of adherens junction by promoting localization of VE-cadherin and β -catenin to endothelial cell

junctions [48]. Paracellular expression of VE-cadherin and/or β -catenin is associated with BBB repair following focal astrocyte loss in vivo [49]. VE-cadherin expression was also decreased at the BBB in mice subjected to focal cerebral ischemia, suggesting that adherens junction disruption contributes to BBB dysfunction in the context of stroke [50].

2.3 Transporters

For many endogenous and exogenous substances, proteins expressed at the BBB determine their ability to traverse biological membranes. Such transport proteins include ATP-binding cassette (ABC) transporters and solute carrier (SLC) transporters [13]. In order to target transporters for optimization of pharmacotherapy, it is critical to understand localization (i.e., luminal versus abluminal), expression, and activity of endogenous transporters at the BBB endothelium (Fig. 2).

ABC Transporters. ABC transporters require biological energy via ATP hydrolysis to transport drugs and their metabolites against their concentration gradient. ABC drug transporters, specifically P-glycoprotein (P-gp), Multidrug Resistance Proteins (MRPs in humans; Mrps in rodents) and Breast Cancer Resistance Protein (BCRP in

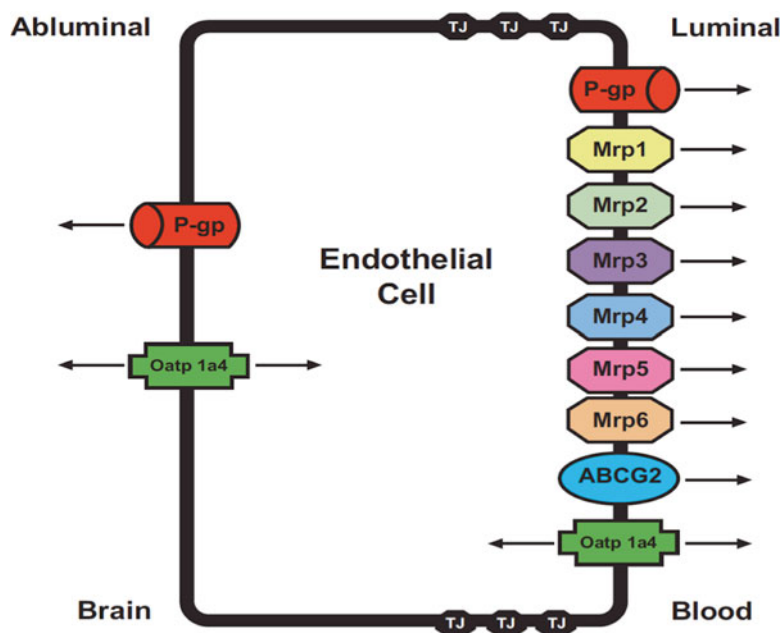


Fig. 2 Endothelial localization of drug transporters known to be involved in transport of therapeutic agents at the blood–brain barrier. Adapted from Ronaldson & Davis. *Curr Pharm Des.* **18**(25): 3624–3644 (2012)

humans; Bcrp in rodents) are involved in cellular extrusion of drugs and greatly contribute to limiting effective therapeutic delivery to the brain. In general, P-gp transports cationic or basic and neutral compounds, whereas MRPs/Mrps are involved in cellular efflux of anionic drugs as well as their glucuronidated, sulfated, and glutathione-conjugated metabolites [13, 51]. BCRP/Bcrp has significant overlap in substrate specificity profile with P-gp and has been shown to recognize a vast array of sulfo-conjugated organic anions, hydrophobic, and amphiphilic compounds [52].

P-gp is a 170-kDa ATP-dependent transporter that primarily functions as a defense mechanism against potentially toxic xenobiotics [53]. P-gp orthologues from different species have greater than 70 % sequence identity [53] and are encoded by closely related genes (i.e., multidrug resistance (MDR) genes), which have two isoforms in humans (MDR1, MDR2) and three isoforms in both mice (i.e., *mdr1*, *mdr2*, *mdr3*) and rats (i.e., *mdr1a*, *mdr1b*, *mdr2*). The human MDR2 gene and the murine/rodent *mdr2* gene products are exclusively involved in hepatic transport of phosphatidylcholine. In contrast, human MDR1, murine *mdr1/mdr3*, and rodent *mdr1a/mdr1b* are involved in drug transport at the BBB endothelium. P-gp has been localized to both the luminal and abluminal membrane of brain microvascular endothelial cells [54]. Abluminal localization of P-gp has also been identified on perivascular astrocyte foot processes [54–56]. Increased expression and/or activity of P-gp has been reported in hippocampal microvessels isolated from stroke-prone spontaneously hypertensive rats [57], which suggests that alterations in P-gp expression and/or activity may be a component of the BBB response to pathological stressors.

The mammalian MRP family belongs to the ABCC group of proteins [58]. Many of the functionally characterized MRP isoforms (i.e., MRP1/Mrp1, MRP2/Mrp2, MRP4/Mrp4, Mrp5, Mrp6) that have the potential to determine CNS drug delivery have been localized to the mammalian BBB [13, 16]. Additionally, the ability of Mrp isoforms to actively efflux the endogenous antioxidant glutathione (GSH) may have significant implications in stroke. GSH is responsible for maintenance of cellular redox balance and antioxidant defense in the brain. It has been demonstrated that functional expression of Mrps is upregulated in response to oxidative stress, a primary component of stroke pathophysiology, which leads to increased cellular efflux of GSH [59]. Enhanced functional expression of Mrp isoforms at the BBB can cause reduced brain and/or endothelial cell concentrations of GSH, an alteration in cellular redox status, and increased potential for cell injury and death.

Several recent studies have demonstrated localization of BCRP at the brain microvasculature, particularly along the luminal side of the BBB [60, 61]. In terms of transport activity, data from recent *in vitro* and *in vivo* studies are controversial. Although some studies have suggested that BCRP is not functional at the BBB [61, 62] or plays a minimal role in xenobiotic efflux from the brain [63], more detailed analyses have confirmed that BCRP is a critical determinant of drug permeation across the BBB [64, 65]. More recently, oxygen/glucose deprivation was shown to increase the endothelial expression of Bcrp at the mRNA level in an *in vitro* BBB model [66]. *In vivo*, Bcrp mRNA expression was observed to be upregulated in the peri-infarct region following reversible middle cerebral artery occlusion (MCAO) [67]. The functional implications of increased Bcrp molecular expression at the BBB/NVU have yet to be determined in the context of stroke.

Solute Carrier (SLC) Transporters: In contrast to ABC transporters, membrane transport of circulating solutes by SLC family members is governed by either an electrochemical gradient utilizing an inorganic or organic solute as a driving force or the transmembrane concentration gradient of the substance actually being transported. Perhaps the most viable SLC candidates for transporter targeting are members of the SLC21A/SLCO family, which includes organic anion transporting polypeptides (OATPs in humans; Oatps in rodents). OATPs/Oatps have distinct substrate preferences for amphipathic solutes [68]. OATPs/Oatps are well known to transport 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase inhibitors (i.e., statins), which have been shown to exhibit neuroprotective and antioxidant properties [69, 70]. Studies in Oatp1a4(−/−) mice demonstrated reduced blood-to-brain transport of pitavastatin and rosuvastatin as compared to wild-type controls, which suggests that Oatp1a4 is involved in statin transport across the BBB [71]. Although OATP isoforms are expressed in several tissues, not all exist at the BBB. Immunofluorescence staining of human brain frontal cortex demonstrated OATP1A2 localization at the level of the brain microvascular endothelium [72]. Expression of Oatp1a4, Oatp1c1, and Oatp2a1 has been reported at the rodent BBB [73–78]. Oatp1c1 is selectively expressed at the BBB [74], has relatively narrow substrate specificity, and primarily transports thyroxine and conjugated sterols [75, 76]. Oatp2a1 regulates BBB transport of prostaglandins [73]. It has been proposed that Oatp1a4, a rodent homologue of OATP1A2, is the primary drug transporting Oatp isoform expressed at the rat BBB [79, 80]. Recently, our laboratory demonstrated that Oatp1a4 is a BBB transporter that can be effectively targeted for facilitation of effective CNS drug delivery [77, 78].

2.4 Astrocytes

Astrocytes are the most abundant cell type in the brain and cover over 99% of cerebral capillaries with their end-feet [3, 81]. Astrocytes are critical in development and/or maintenance of BBB characteristics [14, 82, 83]. Astrocytes may be involved in transient regulation of cerebral microvascular permeability [84], in particular via dynamic Ca^{2+} signaling between astrocytes and the endothelium via gap junctions and purinergic transmission [85, 86]. Additionally, astrocytes play an essential role in regulating water and ion exchange across the brain microvascular endothelium [87]. Astrocytes possess two high-affinity transporters for uptake of glutamate, termed excitatory amino-acid transporter 1 and 2 (i.e., EAAT1 (i.e., GLAST) and EAAT2 (i.e., GLT-1)) that remove excess glutamate from the synapse [88]. Elevated brain levels of glutamate may lead to a pathological condition known as excitotoxicity, which has been implicated in neuronal damage in ischemic stroke [89]. Additionally, astrocytes are known to express volume-regulated anion channels. These channels are involved in Ca^{2+} -independent release of anionic amino acids (i.e., glutamate, aspartate, taurine) during conditions that cause astrocyte swelling such as cerebral hypoxia [90]. Astrocytes are also known to express transport

proteins including P-gp [91], MRP/Mrp isoforms [13, 59], and Bcrp [92]. The expression of multiple drug transporters in astrocytes suggests that these glial cells may act as a secondary barrier to CNS drug permeation. That is, the balance of transporters in astrocytes may either sequester drugs within the astrocyte cytoplasm, thereby preventing these compounds from reaching their site of action in the brain, or concentrate drugs in brain extracellular fluid. Pharmacological agents within brain extracellular space can be effluxed by active transport mechanisms at brain barrier sites or via “sink” effects of the CSF [51].

2.5 Microglia

Microglia are derived from the monocyte lineage that represents approximately 20% of the total glial cell population within the CNS [93]. Under normal physiological conditions, microglia exist in a quiescent state lacking endocytotic and phagocytotic activity. These microglia possess a ramified morphology characterized by a small (5–10 μm) cell body and multiple radial cell processes extending from the cell body. Ramified microglia are thought to contribute to CNS homeostasis by participating in extracellular fluid cleansing and neurotransmitter deactivation [51]. During disease or trauma, microglia may become activated and the degree of this activation is directly correlated to the type and severity of brain injury [94]. Activated microglia are identified by their larger cell body and short cytoplasmic processes as well as upregulation of cell surface receptors such as CD14 and Toll-like receptors (TLRs) [95, 96]. During an immune response, activated microglia may be further converted into a reactive state, which is characterized by a spheroid or rod-like morphology and the presence of phagocytotic activity. Microglia activation and proliferation has been implicated in the development of neuronal death in ischemic stroke and cerebral hypoxia [97–99]. Furthermore, activation of microglia is associated with dysfunction of the BBB characterized by changes in TJ protein expression and enhanced paracellular permeability [100]. When activated, microglia produce high levels of neurotoxic mediators such as nitric oxide and peroxide as well as inflammatory cytokines (i.e., TNF- α), proteases, and complement components [94]. Excessive production of these substances may further lead to cell injury in the CNS characterized by astrocyte activation, further microglia activation, and neuronal cell death.

Microglia express several ion channels including multiple potassium, calcium, sodium, and chloride channels [101]. The expression patterns of these ion channels depend on the microglial functional state and are involved in a variety of physiological functions including proliferation, ramification, and maintenance of membrane potential, intracellular pH regulation, and cell volume regulation [102]. Glutamate receptors [103] and nutrient carrier systems such as GLUT-1 [104] are expressed in microglia. These cells also express membrane proteins involved in drug transport. Studies in a continuous rat microglia cell line (i.e., MLS-9) demonstrated functional expression of P-gp [91], Mrp1 [105], and Mrp4/Mrp5 [106].

2.6 *Pericytes*

In addition to glia, pericytes also play a crucial role in the maintenance of BBB homeostasis [107]. Pericytes are flat, undifferentiated, contractile cells that attach at irregular intervals along the capillary walls and communicate with the other cell types of the NVU [108]. These cells, via secretion of pericyte-derived angiopoetin, induce expression of occludin at the BBB, which suggests that pericytes are directly involved in induction and/or maintenance of barrier properties [109]. Involvement of pericytes in induction of BBB properties is also exemplified by the observation that proper localization of endothelial proteins (i.e., P-gp, utrophin) requires co-culture with pericytes [110]. Studies using adult-viable pericyte-deficient mouse mutants demonstrated that pericytes are critical in maintaining expression of BBB-specific genes in endothelial cells (i.e., transferrin receptor) and by inducing polarization of astrocyte end-feet adjacent to the cerebral microvasculature [111]. Additionally, MRP isoforms (MRP1, MRP4, MRP5) have been identified in pericytes in vitro, which implies that pericytes may contribute to regulation of BBB xenobiotic permeability and to CNS drug delivery [112].

2.7 *Neurons*

Noradrenergic [113], serotonergic [114], cholinergic [115], and GABAergic [116] neurons have been shown to make distinct connections with other cell types of the NVU. The need for direct innervation of brain microvasculature comes from the dynamic nature of neural activity and the metabolic requirements of nervous tissue, implying that the cerebral microcirculation must be highly responsive to the needs of CNS tissue. Interestingly, disruption of BBB integrity induced by pathophysiological factors (i.e., inflammation, hypertension, ischemia) often accompanies changes in cerebral blood flow and perfusion pressure [117–119] and there is evidence that such BBB opening may be a selective, compensatory event rather than a simple anatomical disruption. This implies that communication between neurons and the brain microvasculature may not simply regulate blood flow, but BBB permeability as well.

2.8 *Extracellular Matrix*

In addition to cellular components of the NVU, the extracellular matrix of the basal lamina also interacts with the BBB endothelium. Disruption of extracellular matrix is strongly associated with increased BBB permeability in pathological states including stroke [120, 121]. The extracellular matrix serves as an anchor for the endothelium via interaction of laminin and other matrix proteins with endothelial integrin receptors [122]. Matrix proteins can also influence expression of TJ proteins [123, 124]. Proteolysis of extracellular matrix proteins is well known to occur

in response to stroke, an effect that greatly contributes to BBB disruption [125]. Additionally, protein fragments generated from extracellular matrix proteolysis (i.e., perlecan domain V) may have beneficial effects in neuroprotection and post-stroke brain repair [125, 126].

3 Ischemic Stroke

3.1 Overview of Ischemic Stroke

Stroke is a leading cause of death and long-term disability in the United States. Every year, more than 795,000 Americans suffer from either a new or recurrent stroke, which averages one incidence of stroke every 40 s [127]. Of all strokes, 87 % are ischemic [127]. The annual cost of stroke rehabilitation in the United States is an estimated \$34 billion [127]. Several factors have been identified that increase risk of stroke including history of transient ischemic attacks, hypertension, impaired glucose tolerance and diabetes mellitus, atrial fibrillation, cigarette smoking, and low serum concentrations of HDL cholesterol [127].

Stroke is characterized by a heterogeneous spectrum of conditions caused by interruption of blood flow supplying the brain [128]. Such a deficit in cerebral blood flow causes an irreversibly damaged ischemic core and salvageable surrounding neural tissue known as the penumbra [129]. CNS energy requirements are met by brain uptake of oxygen and glucose, which are metabolized to enable phosphorylation of ADP to ATP. When blood flow to the brain is interrupted during stroke, the ischemic core is quickly deprived of oxygen and glucose. Inability to provide sufficient quantities of ATP causes collapse of ion gradients and subsequent release of neurotransmitters (i.e., dopamine, glutamate), an event that causes neuronal cell death and development of an infarction [130]. Excess release of glutamate is particularly deleterious to the CNS due to overstimulation of glutamate receptors, activation of phospholipases/sphingomyelinases, phospholipid hydrolysis, release of arachidonic acid and ceramide, and disruption of CNS calcium homeostasis [4]. Oxidative stress is also observed in the CNS at early time points following ischemic injury and is well known to contribute to cell death in the ischemic core [131]. As neuronal cell damage extends to the ischemic penumbra, neuroinflammation and apoptosis become more prevalent and dramatically affect viability of salvageable brain tissue within the penumbra [131].

Cell death processes in the ischemic core occur extremely rapidly (i.e., within minutes) thereby rendering this region difficult to protect using pharmacological approaches [4]. In contrast, cells within the ischemic penumbra die more slowly by active cell death mechanisms [4]. The primary goal of drug therapy for acute ischemic stroke is to salvage the penumbra as much as possible and as early as possible [129]. Currently, there is only one therapeutic agent approved by the FDA for acute ischemic stroke treatment, recombinant tissue plasminogen activator (r-tPA) [132]. The primary goal of r-tPA therapy is to restore the blood flow and oxygen supply to

ischemic brain tissue; however, most cellular damage to the brain occurs when cerebral perfusion is re-established (i.e., reoxygenation) [132]. Such hypoxia/reoxygenation (H/R) stress/injury is associated with neuronal apoptosis characterized by cytochrome *c* release, caspase-3 activation, and internucleosomal DNA fragmentation [80]. Pathophysiological mechanisms that can cause neuronal apoptosis during H/R include increased production of reactive oxygen species (ROS) [16, 80]. ROS contribute to brain injury by interacting with proteins, lipids, and nucleic acids as well as via activation of redox-sensitive signaling pathways. Such responses are characterized by increased CNS production of hydrogen peroxide, upregulation of the cellular stress marker heat shock protein-70, and increased nuclear expression of hypoxia-sensitive transcription factors such as hypoxia-inducible factor-1 and nuclear factor- κ B [80]. The H/R component of stroke is also associated with decreased brain concentrations of GSH [133], an effect that is further indicative of oxidative stress. These molecular events associated with H/R injury emphasize a critical need in stroke therapy for discovery of new drugs that can be administered alone or in conjunction with r-tPA for “rescue” of neural tissue.

3.2 The Neurovascular Unit in Ischemic Stroke

3.2.1 Disruption of the Blood–Brain Barrier

BBB homeostasis is dependent on discrete interactions between NVU components [134, 135]. Perturbation of extracellular matrix (i.e., type IV collagen, heparan sulfate proteoglycan, laminin, fibronectin, perlecan) disrupts cell-matrix and cell–cell signaling mechanisms critical to NVU function [121, 125]. Proteinases such as matrix metalloproteinases (MMPs) contribute to breakdown of the extracellular matrix and BBB disruption in stroke [136]. This includes MMPs that are activated by HIF-1 α -dependent mechanisms (i.e., MMP2) and MMPs whose activation is triggered by pro-inflammatory cytokines (i.e., TNF- α , IL-1 β) such as MMP3 and MMP9 [135]. Involvement of MMPs in BBB disruption following ischemic stroke has been reported in experimental stroke models [137, 138]. Furthermore, a recent clinical study demonstrated that MMP9 levels were elevated in stroke patients [139]. MMPs directly compromise the BBB by degrading tight junction constituent proteins such as claudin-5 and occludin [135]. MMP-mediated opening of the BBB in ischemic stroke may be regulated by nitric oxide (NO) signaling [140]. Overall, damage and subsequent opening of the BBB is a key event in development of intracerebral hemorrhage and brain edema following ischemic stroke.

Experimental models of stroke have provided considerable information on solute leak across the BBB. Using the transient MCAO rodent model, Pfefferkorn and Rosenberg demonstrated increased leak of sucrose, a vascular marker that does not typically cross the BBB, in the ischemic hemisphere [141]. However, BBB disruption following an ischemic insult is much more profound than to allow leak of small molecules only. Recently, it was shown that BBB disruption following focal cerebral

ischemia was sufficient to allow blood-to-brain leak of Evan's blue dye [41]. Evan's blue dye, when unconjugated to plasma proteins, is a relatively small molecule with a molecular weight of 960.8 Da. It is well established that Evan's blue dye irreversibly binds to serum albumin *in vivo*. This leads to the formation of a very large, solute-protein complex (i.e., in excess of 60,000 Da) that can only traverse the BBB under considerable pathological stress such as that observed during an ischemic stroke [142]. Jiao and colleagues observed redistribution of various tight junction proteins following 2 h of focal cerebral ischemia, an event that correlated with increased blood-to-brain flux of Evan's blue-albumin [41]. Reorganization of tight junction proteins following focal cerebral ischemia is also mediated by vascular endothelial growth factor (VEGF) [143] and NO [144].

BBB dysfunction and subsequent leak across the microvascular endothelium following focal ischemia enables considerable movement of vascular fluid across the microvascular endothelium and development of vasogenic edema [145]. Recent studies using the MCAO model have shown that water movement across the BBB is exacerbated by enhanced blood-to-brain movement of sodium. Alterations in sodium gradients across the microvascular endothelium dramatically alter oncotic pressure and are facilitated by increased functional expression of Na–K–Cl cotransporter [146], as well as Na–H exchangers NHE1 and/or NHE2 [147]. MCAO studies in spontaneously hypertensive rats demonstrated that the NHE1 transporter is a critical regulator of ischemic-induced infarct volume [148]. Disruption of sodium gradients across the BBB during ischemic stroke can also involve upregulation of sodium-dependent glucose transporters such as sodium–glucose cotransporters (SGLTs). Specifically, pharmacological inhibition of SGLT in MCAO rats significantly reduced infarct and edema ratios, which implies that this transporter may be a critical determinant of stroke outcome [149].

Functional BBB integrity is disrupted by production of ROS and subsequent oxidative stress (Fig. 3). Production of superoxide anion, a potent ROS generated when molecular oxygen is reduced by only one electron, is a known mediator of cellular damage following stroke [150, 151]. Superoxide dismutase (SOD) enzymes tightly control biological activity of superoxide anion, a by-product of normal physiological processes. Under oxidative stress conditions, superoxide is produced at high levels that overwhelm the metabolic capacity of SOD. This phenomenon is supported by the observation that infarct size and cerebral edema were markedly reduced in mice engineered to overexpress SOD as compared to wild-type controls [150]. Increased levels of superoxide also contribute to BBB endothelial dysfunction [152, 153]. BBB damage can be intensified by conjugation of superoxide and NO to form peroxynitrite, a cytotoxic and pro-inflammatory molecule. Peroxynitrite causes injury to cerebral microvessels through lipid peroxidation, consumption of endogenous antioxidants (i.e., reduced GSH), and induction of mitochondrial failure [80, 154]. Peroxynitrite is known to induce endothelial damage by its ability to nitrosylate tyrosine, leading to functional modifications of critical proteins [155]. Peroxynitrite formation at the BBB becomes more likely with activation of endothelial nitric oxide synthase (eNOS) and inducible NOS (iNOS) because NO rapidly diffuses through membranes and reacts with superoxide anion [154].

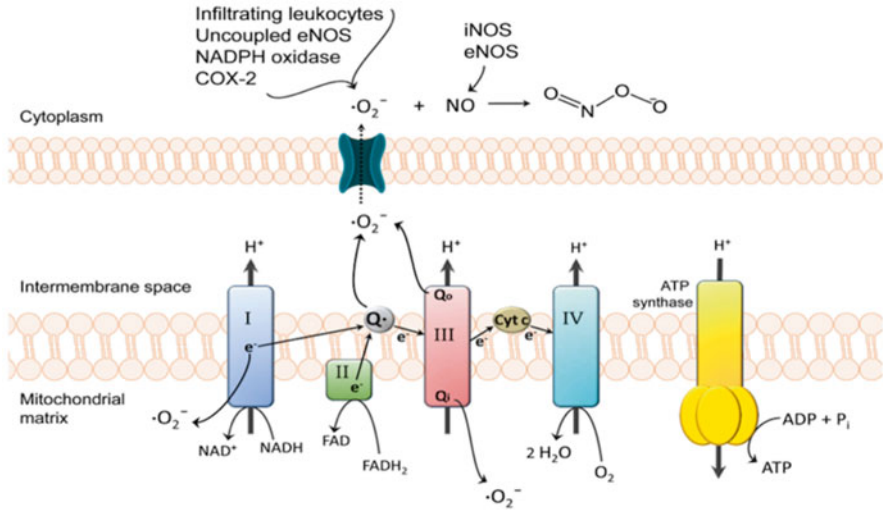


Fig. 3 Generation of reactive oxygen species (ROS) in brain microvascular endothelial cells. During disease, mitochondrial superoxide levels increase via NO inhibition of cytochrome complexes and oxidation of reducing equivalents in the electron transport chain. Complex I as well as both sides of complex III (i.e., Q_i and Q_o sites) are the most common sources of mitochondrial superoxide. Superoxide generated within the intermembrane space of mitochondria can reach the cytosol through voltage-dependent mitochondrial anion channels. Superoxide levels further increase via cyclooxygenase-2, NADPH oxidase, uncoupled eNOS, and infiltrating leukocytes. The resulting high levels of superoxide coupled with the activation of NO-producing eNOS and iNOS increases the probability of peroxynitrite formation. Peroxynitrite-induced cellular damage includes protein oxidation, tyrosine nitration, DNA damage, poly(ADP-ribose) polymerase activation, lipid peroxidation, and mitochondrial dysfunction. Adapted from Thompson and Ronaldson (2014). *Adv Pharmacol.* **71**: 165–209

Inflammatory stimuli are critical mediators of BBB dysfunction in the setting of stroke. Previous research has demonstrated that inflammatory mechanisms in focal cerebral ischemia are mediated through pro-inflammatory cytokines TNF- α and IL-1 β , which appear within 2–6 h following ischemic insult [156]. Pro-inflammatory signaling induces adhesion molecules and subsequent transmigration of activated neutrophils, lymphocytes, or monocytes into brain parenchyma [157]. Physiologically, expression of vascular adhesion molecules such as ICAM-1 and VCAM-1 are minimally detectable at the BBB; however, their expression is dramatically increased in response to diseases such as stroke [157, 158]. Pro-inflammatory mediators also alter functional expression of endogenous BBB transporters and tight junction proteins. For example, TNF- α increased expression of P-gp but decreased BCRP expression in hCMEC/d3 cells [159]. Production and secretion of TNF- α and IL-1 β has been observed to alter the expression of occludin and ZO-1 [160], suggesting involvement of inflammation in exacerbating paracellular leak during stroke.

3.3 *Ischemic Stroke and Glial Support of the BBB*

It is well established that the glial response to an ischemic insult is highly complex and multifaceted; however, it is known that injury and/or activation of astrocytes at the NVU leads to compromise of the BBB. In the inferior colliculus, focal astrocyte loss demarcated by reduced GFAP immunoreactivity directly corresponded with decreased paracellular localization of critical tight junction proteins claudin-5, occludin, and ZO-1 [83]. At the same time points that tight junction proteins were downregulated, an increase in leak of dextran (10 kDa) and fibrinogen was observed [83], which suggests a significant disruption of the BBB due to astrocyte cell death. The results of this study and others [161, 162] illustrate the requirement of intercellular communication between astrocytes and endothelial cells for maintenance of BBB integrity. Additionally, astrocytes have many other features that contribute to BBB physiology. For example, astrocytes are well known to express the water channel aquaporin 4 (AQP4) at end-feet localized adjacent to brain microvascular endothelium, which contributes to endothelial cell polarity and brain water volume [163]. Astrocytes secrete VEGF and fibroblast growth factor-2 (FGF-2), which promote angiogenesis and regulate biological transport at the BBB [164].

Astrocytes are essential contributors to the brain immunological response during ischemic stroke. Astrocytes maintain focal contacts with neighboring microglia and maintain these cells in a dormant, ramified state [94]. Regulation of microglia by astrocytes is prevented by inflammatory signaling, thus enabling microglia to elicit an immune response [94]. Astrocytes can directly contribute to brain immunological responses via upregulation of adhesion molecules (i.e., ICAM-1, VCAM) at their cell surface, an event that contributes to CNS targeting of leukocytes [165]. This enhancement is also facilitated by astrocytic secretion of chemokines such as macrophage inflammatory protein-1 α/β (MIP-1 α/β), monocyte chemoattractant protein-1 (MCP-1), and regulated upon activation normal T-cell expressed and secreted (RANTES). Production and secretion of these chemokines by astrocytes is well known to occur in response to ischemic stroke [166, 167] and to increased brain parenchymal concentrations of tPA [168]. Of particular note, MCP-1 secretion by astrocytes was shown to coincide with a significant increase in FITC-albumin leak in an *in vitro* co-culture of endothelial cells and astrocytes subjected to 5 h oxygen–glucose deprivation, suggesting that MCP-1 may be a critical factor involved in BBB opening following an ischemic stroke [166]. Astrocytes also synthesize several pro- and anti-inflammatory cytokines including interleukins (i.e., IL-1 α , IL-1 β , IL-4-8, IL-10), TNF- α , and interferon- γ [169] as well as transforming growth factor- β (TGF- β) [170]. Cytokines such as TNF- α , IL-1 α , IL-1 β , and interferon- γ can trigger the endothelium and activate processes involved in BBB disruption. Although secretion of TGF- β 1 may play a neuroprotective role in brain parenchyma [170], its effects on the endothelium are much more deleterious. Specifically, excessive endothelial stimulation by TGF- β 1 affects the angiogenic response to ischemic stroke by causing formation of capillaries that lack pericytes, contain fewer endothelial cells and are shorter in length [84]. Indeed, pharmacological

targeting of TGF- β signaling at the level of the brain microvascular endothelium may be an efficacious approach for the protection of BBB integrity and/or preservation of angiogenic responses to stroke.

Inflammatory signaling by astrocytes is a critical event in exacerbation of CNS oxidative stress during ischemic stroke. Previous studies have shown that astrocytic production of pro-inflammatory cytokines can induce deleterious processes in astrocytes themselves via upregulation of iNOS [171]. Upregulation of iNOS leads to a significant enhancement in NO production, which can react with superoxide to produce peroxynitrite. Increased exposure of astrocytes to peroxynitrite can lead to rapid astrocyte proliferation and hypertrophy (i.e., reactive astrocytosis) and astrocyte apoptosis [172, 173]. Reactive astrocytosis is associated with disruption of tight junctions between adjacent brain microvessel endothelial cells and increased BBB permeability [174].

Although astrocyte injury is a critical determinant of BBB dysfunction in the setting of ischemic stroke, endothelial damage can also be induced via immune stimulated microglia [175]. This is supported by the observation that minocycline, a pharmacological inhibitor of activated microglia, dramatically reduced cell death in cultures of murine endothelial cells exposed to activated microglia in vitro [175]. Activated microglia produce pro-inflammatory cytokines in response to cerebral ischemia [176], all of which can trigger BBB disruption. In the setting of ischemic stroke, cytokine production in microglia is mediated by NF- κ B signaling [177]. Inflammatory signaling in microglia may also involve cyclooxygenase-2 (COX2), which is inducible in response to ischemic injury and contributes to opening of the BBB [135]. In neuroinflammation, COX2 activates sphingomyelinases leading to release of ceramides, an event that leads to activation of p38 mitogen-activated protein kinase (MAPK) and subsequent secretion of pro-inflammatory cytokines [178–180]. Taken together, these observations point to a critical role for microglia in the inflammatory response to ischemic stroke.

3.4 Targeting the Neurovascular Unit in Ischemic Stroke

3.4.1 Targeting the Tight Junction

The BBB and associated glial support network of the NVU are clearly compromised in response to ischemic stroke. A critical “component” of ischemic stroke is cerebral hypoxia and subsequent brain injury resulting from reoxygenation/reperfusion (i.e., H/R stress). Over the past several years, our laboratory has studied BBB changes associated with H/R stress in an in vivo rodent model [21, 22, 26, 78, 148, 181]. Changes in BBB integrity under H/R conditions were demarcated by enhanced brain accumulation of 14 C-sucrose [22, 26, 181], a vascular marker that does not typically cross the brain microvascular endothelium. Additionally, H/R stress also increased vascular leak to dextrans (molecular weight range 4–10 kDa) in hippocampal and cortical microvessels [182]. These alterations in BBB permeability in animals subjected to H/R stress were directly associated with an increase in the

expression of HIF-1 α and NF- κ B in nuclear fractions isolated from intact microvessels [183]. In our studies, changes in brain solute uptake is not likely attributed to altered cerebral blood flow because we have previously shown that blood flow changes are negligible in our in vivo H/R model [26]. Changes in BBB permeability to 14 C-sucrose and dextrans were directly correlated with modified organization and/or expression of constituent TJ proteins including occludin, claudin-5, and ZO-1 [22, 26, 182]. Of paramount significance was the observation that H/R stress disrupted disulfide-bonded occludin oligomeric assemblies, thereby preventing monomeric occludin from forming an impermeable physical barrier to paracellular transport [21]. These changes in tight junction organization and BBB solute leak also correlated with a significant increase in brain water content following H/R, providing further evidence that disruption of the BBB under conditions of cerebral ischemia contributes to vasogenic edema [181].

Production of ROS and subsequent oxidative stress has been shown to alter the BBB expression of claudin-5 and occludin leading to increased paracellular solute leak [184]. Therefore, we hypothesized that oxidative stress associated changes in BBB permeability and occludin expression could be attenuated with the use of an antioxidant drug. In order to conduct these studies, we utilized 4-hydroxy-2,2,6,6-tetramethylpiperidine-*N*-oxyl (TEMPOL), a stable, membrane-permeable, water-soluble nitroxide antioxidant. TEMPOL shows SOD-like activity towards the superoxide anion as well as reactivity with hydroxyl radicals, nitrogen dioxide, and the carbonate radical. TEMPOL readily crosses the BBB and has been previously shown to provide neuroprotection as a free radical scavenger in several models of brain injury and ischemia [185, 186]. Using the dual artery in situ brain perfusion technique, we demonstrated that administration of TEMPOL 10 min before H/R treatment significantly attenuated CNS uptake of 14 C-sucrose as compared to animals subjected to H/R only [22]. This reduction in 14 C-sucrose leak was associated with a preservation of occludin localization and occludin oligomerization at the TJ [22]. Specifically, TEMPOL inhibits breakage of disulfide bonds on occludin monomers and thus prevents breakdown of occludin oligomeric assemblies and subsequent blood-to-brain leak of circulating solutes (Fig. 4). Restoration of BBB functional integrity coincided with a decrease in nuclear translocation of HIF-1 α and a decrease in microvascular expression of the cellular stress marker heat shock protein 70 (hsp70) in rats subjected to H/R stress and administered TEMPOL [22]. Taken together, these observations provide evidence that the tight junction can be targeted pharmacologically during ischemic stroke for the purpose of reducing both oxidative stress associated injury to the brain microvascular endothelium and blood-to-brain solute leak (i.e., vascular protection).

3.5 Targeting Endogenous BBB Transporters

The ability of a drug to elicit a pharmacological effect at the level of the BBB requires achievement of efficacious concentrations within CNS. This therapeutic objective is dependent upon multiple mechanisms of transport that may include uptake into the brain by an influx transporter and/or extrusion by an efflux

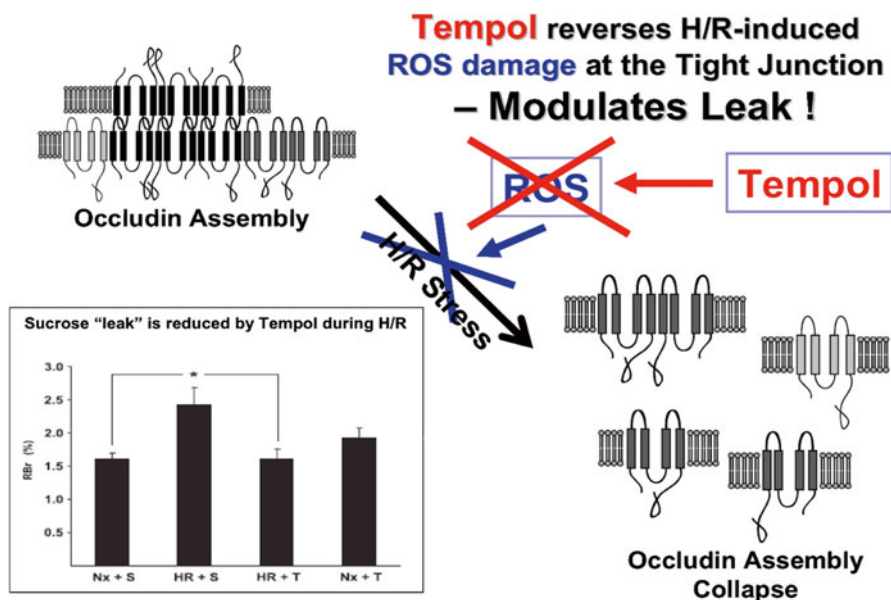


Fig. 4 Effect of TEMPOL on H/R-mediated disruption of the tight junction. ROS and subsequent oxidative stress are known to disrupt assembly of critical TJ proteins such as occludin. Our results show that administration of TEMPOL, by scavenging ROS, prevents disruption of occludin oligomeric assemblies. Furthermore, TEMPOL attenuates the increase in sucrose leak across the BBB observed in animals subjected to H/R stress. Taken together, our studies with TEMPOL demonstrate that the TJ can be targeted pharmacologically in an effort to preserve BBB functional integrity during ischemic stroke. Adapted from Ronaldson & Davis. *Curr Pharm Des.* **18(25)**: 3624–3644 (2012)

transporter. For many drugs, it is this balance between influx and efflux that determines if a drug will elicit a therapeutic effect in the brain or at the BBB. The complexity of drug transporter biology is further underscored by the observation that functional expression of transporters can be dramatically altered by oxidative stress [59, 187, 188]. A thorough understanding of regulation and functional expression of endogenous BBB transporters in both health and disease is essential for effective pharmacotherapy. Furthermore, such information will enable effective targeting of transporters and/or transporter regulatory mechanisms, thus allowing endogenous BBB transport systems to be exploited for purposes of improving CNS drug delivery and/or conferring BBB protection.

Considerable research has focused on studying mechanisms that limit endothelial membrane transport by describing the role of P-gp in restricting drug uptake from the systemic circulation [1, 189–191]; however, clinical trials targeting P-gp with small molecule inhibitors have been unsuccessful in improving pharmacotherapy due to inhibitor toxicity and/or enhanced tissue penetration of drugs [192, 193]. An alternative approach for optimizing delivery of drugs is to focus on BBB transporters that are involved in blood-to-brain transport. One intriguing candidate is Oatp1a4 (Fig. 5), which is known to transport HMG-CoA reductase inhibitors

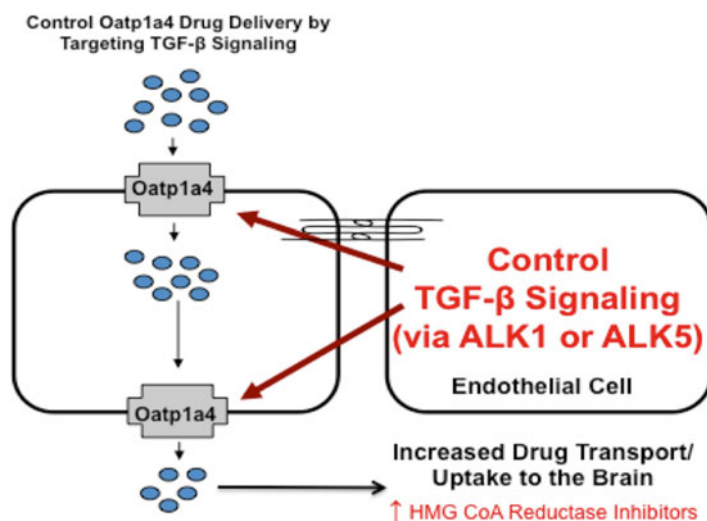


Fig. 5 Targeting Oatp transporters at the BBB for optimization of CNS drug delivery. Results from our studies demonstrate that targeting Oatp transporters during pathophysiological stress can modify CNS drug delivery. Oatp1a4 facilitates brain delivery of drugs that may exhibit efficacy in treatment of peripheral inflammatory pain or cerebral hypoxia such as statins and opioid peptide analgesics. The TGF- β signaling pathway enables control of Oatp isoforms by targeting TGF- β receptors (i.e., ALK1, ALK5) with small molecule therapeutics

(i.e., statins). Recent evidence suggests that statins can act as ROS scavengers independent of their well-documented effects on cholesterol biosynthesis [70]. Specifically, studies in dogs demonstrated that atorvastatin reduced the expression of oxidative and nitrosative stress markers (i.e., protein carbonyls, 4-hydroxy-2-noneal, 3-nitrotyrosine) and increased brain GSH levels [70, 194]. Interestingly, Cui and colleagues showed, *in vivo*, that atorvastatin administration during the acute phase of cerebral ischemia prevented increases in BBB permeability [195]. More recently, simvastatin was demonstrated to preserve barrier function following experimental intracerebral hemorrhage in an *in vivo* study involving MRI measurements of T_{1sat} , a marker of BBB integrity [196]. Taken together, these studies suggest that targeted delivery of statins may be an effective strategy for neuroprotection and/or BBB protection in the setting of stroke. We have shown, *in vivo*, that Oatp1a4 is a BBB transporter target that can be exploited to optimize CNS delivery of drugs, including statins [77, 78].

Although pathophysiological stressors can modulate BBB transporters, such changes must be controlled to provide optimal delivery of drugs. For example, we have demonstrated increased functional expression of Oatp1a4 only after 1 h hypoxia followed by up to 1 h reoxygenation [78]. If Oatp1a4 is to facilitate effective delivery of drugs (i.e., statins), its functional expression must be reliably controlled over a more desirable time course than is possible by relying solely on disease mechanisms. This objective can be accomplished by pharmacological targeting of Oatp regulatory pathways such as the TGF- β system [77, 78, 80]. TGF- β s

are cytokines that signal by binding to a heterotetrameric complex of type I and type II receptors [197]. The type I receptors, also known as activin receptor-like kinases (ALKs) propagate intracellular signals through phosphorylation of receptor-specific Smad proteins (i.e., (R)-Smads). At the BBB, only two ALK receptors (ALK1, ALK5) have been identified [28]. We have shown that pharmacological inhibition of TGF- β /ALK5 signaling can increase Oatp1a4 functional expression [77, 78]. This observation suggests that targeting of the TGF- β /ALK5 pathway may enable control of BBB Oatp1a4 expression and/or activity, thereby providing novel strategies for improved CNS drug delivery and/or BBB protection in stroke.

Optimization of drug delivery is not the only benefit that can be achieved from targeting transporters. BBB transporters mediate the flux of endogenous substrates, many of which are essential to the cellular response to pathological insult. One such substance is the endogenous antioxidant GSH. During oxidative stress, GSH is rapidly oxidized to glutathione disulfide (GSSG). Therefore, the redox state of a cell is represented by the ratio of GSH to GSSG [198]. In vitro studies using human and rodent brain microvascular endothelial cells have demonstrated that hypoxia reduces intracellular GSH levels and decreases the GSH:GSSG ratio, suggesting significant oxidative stress at the level of the BBB [199–201]. Using an in vivo model, oxidative stress was shown to cause BBB disruption characterized by altered expression/assembly of tight junction proteins occludin, claudin-5, and ZO-1 [21–23, 26, 35, 182]. These tight junction modifications correlated with increased BBB permeability to sucrose, an established vascular marker [22, 181], and dextrans [182]. Such increases in BBB permeability can result in leak of neurotoxic substances from blood into brain and/or contribute to vasogenic edema. BBB protection and/or repair in stroke are paramount to protecting the brain from neurological damage. One approach that can accomplish this therapeutic objective is to prevent cellular loss of GSH from endothelial cells by targeting endogenous BBB transporters (Fig. 6). BBB transporters that can transport GSH and GSSG include MRPs/Mrps. Both GSH and GSSG are substrates for MRP1/Mrp1 [59, 202, 203], MRP2/Mrp2 [204], and MRP4/Mrp4 [205]. It is well known that increased cellular concentrations of GSH are cytoprotective while processes that promote GSH loss from cells are damaging [206]. Therefore, it stands to reason that pharmacological targeting of Mrps during oxidative stress may have profound therapeutic benefits including vascular protection at the level of the BBB. Using the known Mrp transport inhibitor MK571, Tadepalle and colleagues showed that inhibition of Mrp1-mediated GSH transport resulted prevented GSH depletion in primary cultures of rat astrocytes [203]. Indeed, the effect of Mrp transport inhibition at the BBB and its effect on endothelial redox status and barrier integrity require further study.

Previous studies have shown that Mrp expression and/or activity can change in response to oxidative stress [59, 207]. Altered BBB expression of Mrps may prevent endothelial cells from retaining effective GSH concentrations. A thorough understanding of signaling pathways involved in Mrp regulation during oxidative stress will enable development of pharmacological approaches to target Mrp-mediated efflux (i.e., GSH transport) for the purpose of preventing BBB dysfunction in diseases with an oxidative stress component. One intriguing pathway is signaling mediated by nuclear factor E2-related factor-2 (Nrf2), a sensor of oxidative stress [188, 208]. In the presence of ROS, the cytosolic Nrf2 repressor Kelch-like ECH-associated

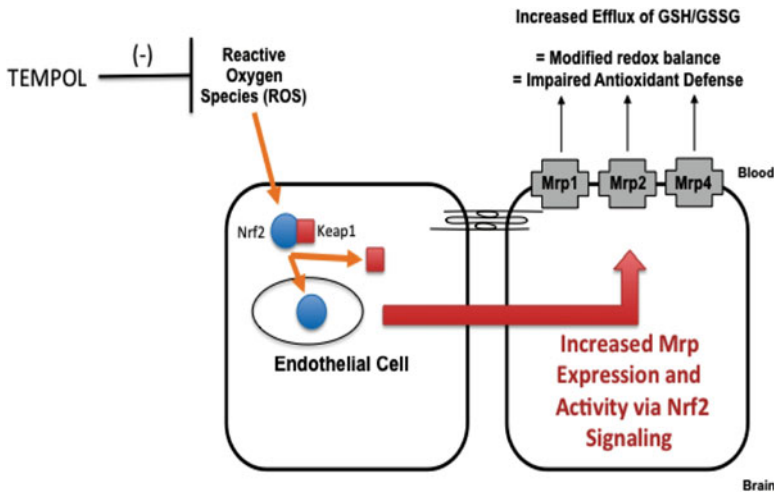


Fig. 6 Prevention of BBB Dysfunction by Targeting Mrp Isoforms in Cerebral Endothelial Cells. Results from our laboratory demonstrate increased expression of Mrp1, Mrp2, and Mrp4 at the BBB following an H/R insult. Furthermore, H/R stress is known to suppress GSH levels and increase GSSG concentrations in the brain. We propose that changes in GSH/GSSG transport occur during H/R as a result of altered functional expression of at least one Mrp isoform. Since Nrf2, a ROS sensitive transcription factor, is known to regulate Mrps, we hypothesize that this pathway is a critical regulatory mechanism for Mrps at the BBB. TEMPOL, a ROS scavenging antioxidant, is a pharmacological tool that can be utilized in order to understand how targeting activation of the Nrf2 pathway can control Mrp expression/activity

protein 1 (Keap1) undergoes structural alterations that cause dissociation from the Nrf2-Keap1 complex. This enables Nrf2 to translocate to the nucleus and induce transcription of genes that possess an antioxidant response element at their promoter [209, 210]. It has been demonstrated that activation of Nrf2 signaling induces expression of Mrp1, Mrp2, and Mrp4 [188, 207, 209, 211]. An emerging concept is that Nrf2 acts as a double-edged sword [210]: on one hand, Nrf2 is required for protecting tissues from oxidative stress; on the other, its activation can lead to deleterious effects. Therefore, an alteration in the balance of Mrp isoforms via activation of Nrf2 signaling may adversely affect redox balance and antioxidant defense at the brain microvascular endothelium. Indeed, this points towards a need for rigorous study of pharmacological approaches (i.e., use of antioxidant drugs such as TEMPOL) that can modulate Nrf2 signaling and control expression of Mrp isoforms and/or GSH transport at the BBB.

3.6 Targeting Glial Support of the BBB

In addition to the BBB endothelium, glial cells (i.e., astrocytes, microglia) are potential therapeutic targets in treatment of stroke. As noted above, glia play a crucial role in regulating BBB functional integrity in health and disease through release

of trophic factors that maintain tight junction protein complexes, release of factors that promote angiogenesis, pro-inflammatory signaling, and production of ROS. Pharmacological manipulation of glial cell biology represents a therapeutic approach that may enable control of BBB/NVU pathophysiological mechanisms during ischemic stroke and/or H/R injury.

An opportunity for cellular protection of glia in ischemic stroke involves targeting the proteinase-activated receptor (PAR) pathway. To date, four members of the PAR family (i.e., PAR-1, PAR-2, PAR-3, PAR-4) have been cloned and characterized [212]. Both PAR-1 and PAR-2 are expressed on the cell surface of astrocytes [213] and microglia [214, 215] as well as on the endothelial cell surface [216]. PAR-1 has been implicated in cytoprotective mechanisms [217, 218] while PAR-2 is involved in regulation of inflammatory responses [219]. Recent research has focused on pharmaceutical development of agonists targeted to the PAR-1 receptor such as activated protein C (APC) [216]. In a mouse model of transient cerebral ischemia, APC was shown to reduce ischemic brain damage and promote neovascularization and neurogenesis, suggesting that pharmacological targeting of the PAR-1 receptor may be an efficacious approach for the treatment of ischemic stroke [220]. Brain vascular perfusion studies demonstrated that brain accumulation of APC was reduced by 64 % in mice lacking the endothelial protein-C receptor (EPCR), suggesting that CNS delivery of APC is dependent upon saturable EPCR-mediated transport at the BBB [221]. Although native APC exhibits cytoprotection in stroke models, its use is limited by bleeding complications [222]; however, a mutant form of APC termed 3K3A-APC has been discovered that exhibits considerable cytoprotective efficacy without complications of bleeding [218]. Specifically, studies in human brain endothelial cells *in vitro* showed that 3K3A-APC protected these cells from oxygen–glucose deprivation to a significantly greater degree than APC [218]. Furthermore, 3K3A-APC improved the functional outcome and reduced the infarction size at a level that was significantly better than APC in the *in vivo* murine distal MCAO model [218], which implies that 3K3A-APC offers a safer and more efficacious alternative to APC in pharmacological targeting of the PAR-1 receptor. In the presence of r-tPA, 3K3A-APC reduced significant reduced infarct volume following focal cerebral ischemia in mice and embolic stroke in rats by up to 65 % [223]. More recently, a phase I clinical trial of 3K3A-APC demonstrated that this therapeutic was well tolerated in healthy adult volunteers [224], which provides an impetus to study the effects of 3K3A-APC in stroke patients. In the case of the PAR-2 receptor, a small molecule PAR-2 antagonist (i.e., N1-3-methylbutyryl-N4-6-aminohexanoyl-piperazine; ENMD-1068) has been shown to attenuate inflammatory responses in a dose-dependent manner [225].

Minocycline is a tetracycline with anti-inflammatory properties that directly inhibit microglial activation. Minocycline easily crosses the BBB, has a good safety profile, and a delayed therapeutic window thus rendering it an ideal candidate drug for treatment of ischemic stroke [226]. Blocking microglial activation may limit BBB disruption and reduce vasogenic edema in the context of ischemic stroke. For example, Yenari and colleagues reported that, *in vivo*, minocycline reduced infarction volume and neurological deficits as well as prevented BBB disruption and hem-

orrhage in a murine experimental stroke model [175]. In vitro, inhibition of microglial activation with minocycline limited ischemic damage in cultured endothelial cells and reduced superoxide release following oxygen–glucose deprivation [175]. More recently, combination therapy of minocycline and candesartan, a pro-angiogenic drug, was shown to improve long-term recovery in Wistar rats subjected to MCAO [227]. In vivo, minocycline was observed to reduce the frequency of hemorrhage in a murine model of cerebral amyloid angiopathy [228]. Currently, minocycline has been incorporated into clinical trials involving stroke patients. Results of these studies demonstrated that minocycline administration, both alone and in combination with r-tPA, improved functional neurological outcome following ischemic stroke [226].

TLRs are highly expressed in human CNS tissue, particularly by astrocytes and microglia [229]. Targeting these receptors has emerged as a promising goal for therapeutic control of ischemic stroke, primarily because TLRs are involved in BBB dysfunction and NVU ischemic injury [230]. While mRNA for TLRs 1–10 have been detected in murine microglia [231], all except TLR10 have been reported in human microglia [232]. Astrocytes possess a much more limited complement of TLRs since mRNA for TLRs 2, 4, 5, and 9 have been detected in murine astrocytes [233] and only TLR3 mRNA in human astrocytes [234]. While the large number of TLR receptors expressed on glial cells suggests a plethora of potential therapeutic targets for modification of glial pathology in the ischemic brain, much work needs to be done on understanding pharmacokinetics of TLR ligand binding and interactions between the TLR and the Toll/IL-1 receptor before TLR-based stroke therapeutics can reach development [230]. This field has shown promise as demonstrated by recent data in a murine model of cerebral ischemia where CNS delivery of TAT-hsp70 was shown to confer BBB protection via reduction of microglial activation, an effect that may be due to targeting of hsp70 to TLR2/4 [235].

4 Conclusion

The field of BBB biology, particularly the study of tight junction protein complexes and endogenous transport systems, has rapidly advanced over the past two decades. It is now well established that tight junction protein complexes are dynamic in nature and can organize and reorganize in response to ischemic stroke. These changes in tight junctions can lead to increased BBB permeability to small molecule drugs via the paracellular route. Additionally, many previous studies reported on the controversial ability of transporters (i.e., Oatp1a4) to act as facilitators of brain drug uptake. Now, it is beginning to be appreciated that endogenous BBB transporters can facilitate uptake of therapeutics from blood to the brain, thereby rendering these proteins, potential molecular targets for pharmacotherapy. Additionally, MRPs may represent viable molecular targets for BBB vascular protection in the setting of ischemic stroke. Molecular machinery involved in regulating these endogenous BBB transport systems (i.e., TGF- β /ALK5 signaling, Nrf2

pathway) are just now being fully characterized. These crucial discoveries have identified multiple targets that can be exploited for optimization of CNS delivery of therapeutic agents or for protection against BBB dysfunction. Perhaps targeting of currently marketed or novel drugs to influx transporters such as Oatp1a4 or to efflux transporters such as Mrp1, Mrp2, or Mrp4 will lead to significant advancements in ischemic stroke treatment. Identification and characterization of intracellular signaling pathways that can regulate the functional expression of uptake or efflux transporters provides yet another approach for pharmacological control of transporter systems in an effort to precisely deliver therapeutics to the CNS. Additionally, identification and characterization of novel targets on glial cells (i.e., astrocytes, microglia) provide yet another opportunity for the design and development of therapeutics aimed at protecting the BBB/NVU during ischemic injury and, by extension, controlling CNS drug delivery. Future work will continue to provide more insight on the interplay of tight junction protein complexes, transporters, and intracellular signaling pathways at the BBB/NVU and how these systems can be effectively targeted for improved stroke therapy.

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Barrier Mechanisms in Neonatal Stroke

Zinaida S. Vexler

1 Introduction

The concept that the fetal and newborn blood–brain barrier (BBB) is “immature,” implying that the BBB is not formed, leaky or even absent has dominated the field for a long time. However, such a simplistic idea has been challenged in multiple species, demonstrating in naïve brain that many functional barrier elements are already in place in fetal brain and, together with placenta, are effectively preserving brain metabolism and limiting brain edema that is associated with entrance of peripheral components. It has also become apparent that BBB integrity does not linearly change during postnatal brain maturation and that infection, inflammation, brain trauma, or stroke during postnatal development exert age-specific “susceptibility signatures” of BBB disturbances, processes that we will discuss in this chapter.

2 Development of the BBB and Other Brain Barriers

There are multiple brain barriers, including the BBB, blood–CSF barrier across choroid plexus (BCSFB) and the pia-arachnoid barrier [1]. There is also a barrier only present in the embryo, the CSF-ventricular zone interface lining the ventricular system. Cumulatively, these barriers limit exchange of solutes and various molecules between compartments as well as establish efflux/influx mechanisms that control ionic, nutrient, glucose, and other gradients. In the adult, the BBB exists at all levels

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of the vasculature within the CNS, including the penetrating arteries and arterioles, the capillary bed, the postcapillary venules and the draining venules and veins [2, 3].

The development of the BBB is believed to be a multi-step process. Both the development and maintenance of the BBB are governed by intracellular endothelial mechanisms in conjunction with multiple extracellular mechanisms that link endothelial cells to the extracellular matrix (ECM), thereby supporting their proper positioning and functionality. Astrocytes, pericytes, and ECM components provide both structural and functional support to the BBB. BBB development starts mid-gestation and temporally occurs after establishment of a local microglial pool in the brain and before pericytes and astrocytes take their respective positions surrounding the vessels [4–7]. Thus, endothelial sprouts are guided by microglia/macrophages in fetal brain, largely via VEGF-dependent mechanisms [8]. Lack of or reduced number of PU.1 or SDF-1 microglia/macrophage leads to distorted vasculature and embryonic lethality [8]. Mice deficient in VEGF or VEGF receptor 2 (Flk-1^{-/-}) fail vessel formation, which leads to embryonic lethality [9]. Lack of a single VEGF allele also leads to embryonic lethality despite partial blood vessel development [10]. Several VEGF-independent mechanisms that regulate embryonic angiogenesis and BBB formation were identified. Integrin/TGF β signaling, for example, suppresses vascular branching/sprouting and prevents germinal matrix hemorrhage [11]. Excessive vascular sprouting, in turn, underlies cerebral hemorrhage in mice lacking α V β 8-TGF β signaling in the brain [11].

Wnt- β -catenin signaling has been shown to be important in brain angiogenesis and BBB formation, but not in barrier formation in peripheral vessels [12]. The canonical Wnt pathway was demonstrated to play a key role in BBB formation via induction of the BBB genes, such as the nutrient transporter Glut-1 [13], while β -catenin was shown to regulate angiogenesis in the developing brain, in part by inducing expression of the death receptors Dr6 and Troy. Endothelial β -catenin was also shown to play a key role in embryonic and postnatal BBB maturation by regulating the formation of tight junctions (TJs) [14].

Pericytes contribute to BBB development and function in a number of ways, primarily by regulating capillary diameter and CBF and controlling BBB integrity [15]. They also contribute to vessel stability, possibly by affecting both TJs and transcytosis routes [5]. However, the existence and the role of multiple phenotypes are still poorly understood. Pericyte coverage gradually increases starting late gestation and continues through postnatal ages [4, 5]. Astrocytes are another cell population central to the maturing BBB in several ways [15]. Perivascular astrocytic end-feet at the abluminal side of brain vessels are polarized structures that are enriched with water channel aquaporin-4 (AQP-4) and several ion pumps. Astrocytes modulate TJs and are also an essential part of antioxidative mechanisms in the brain, recycling GSSG, which, as we discuss later, is of particular importance after stroke and other diseases. They also provide nutrition for neurons and neurotransmitter clearance and recycling.

Cells of the BBB/neurovascular unit do not exist in isolation. The ECM plays a key role both in the development and maintenance of the BBB under normal and disease conditions. Mutant ECM proteins, such as collagen-IV α 1 isoform and, to a

lesser extent, $\alpha 2$ isoform adversely affect endothelial communications with the matrix, leading to mortality or hemorrhagic transformation [16]. During embryonic angiogenesis, TGF β signaling suppresses vascular branching/sprouting and prevents hemorrhages [11]. Loss of TGF β R2/SMAD3 enhances vascular sprouting, branching, and hemorrhages during embryonic brain development but these effects are independent of BBB dysfunction [11]. Multiple lines of evidence show an important role for endothelial TGF β /TGF β R2 signaling for vascular integrity during embryonic brain development [17]. Endothelial specific depletion of Alk5 leads to a similar cerebral hemorrhagic phenotype and embryonic lethality [17, 18], consistent with a requirement for both TGF β R1 and TGF β R2 proteins as a heteromeric receptor complex in endothelial cells. Mutations in other members of the TGF β family, endoglin, (ENG) and activin receptor-like kinase 1 (ALK1), were demonstrated as major contributors to intracerebral hemorrhages of brain arteriovenous malformations in children [19].

Although individual neurovascular unit components form at different paces during gestation, likely contributing to age-dependent susceptibility to infection, inflammatory stress and stroke, the fetal brain has surprisingly effective influx/efflux mechanisms. Efflux mechanisms are active across brain barriers in the developing brain. Multiple transcription factors involved in metabolic- or cell stress-induced detoxifying enzymes and transporters are expressed throughout embryonic brain development. For example, several proteins of the ATP-binding cassette (ABC) family, which reduce the entry of compounds from blood into the brain by active efflux, are already expressed in cerebral blood vessels and in choroid plexus in the fetal and neonatal rat brain [20]. Among ABC-transporters, Pgp/ABCB1 expression is lower in the fetus, whereas MRP1/ABCC1, MRP4/ABCC4, and BCRP/ABCG2 are expressed at comparable levels in fetal and adult brains. Data on amino acid transporters in endothelial cells in the developing brain are sparse but available data indicate that the function of several amino acid and inward transporters is higher than in the adult. For example, transporter mannose 6-phosphate receptor is expressed in developing rodent and rabbit brain but is progressively lost with age [21].

The BCSFB plays a prominent role in fetal and newborn brain. Ion pumps are active in the BCSFB. Essential nutrients and other molecules important for growth and differentiation of the brain enter the brain via uptake from the CSF. A recent transcriptome approach allowed insights into transport mechanisms at the developing mouse blood–CSF interface [22].

3 Blood–Brain Barrier and Adult Stroke

There is ample evidence of BBB disruption after acute stroke. Several recent reviews comprehensively discussed various mechanistic aspects of neurovascular state in adult stroke, including contribution of the ECM, its communication with endothelial cells, interaction between individual cell types, including endothelial cells,

astrocytes and pericytes, as well as various individual intracellular and intercellular mechanisms of adult stroke [23–25]; these aspects will not be discussed here. It is, however, important to mention that several long-term concepts in the stroke field are currently being revisited and reconsidered. Three most notable examples of revisited questions are the relative role of the paracellular compared to the transcellular route as paths for leukocyte trafficking [26], the ability of neutrophils to reach injured parenchyma and signal from within, rather than signal from the perivascular space [27], and beneficial, rather than purely toxic role of inflammation and microglial involvement in particular [25].

4 Maturation-Dependent Susceptibility of the BBB to Inflammation

Neuroinflammation is a characteristic feature of stroke progression in the adult and is a major contributor to brain injury [25]. Parenchymal, perivascular, and peripheral circulating cells independently and in concert contribute to stroke-induced production of inflammatory mediators and neuroinflammation [25] and activation of endothelial cells [28, 29]. Perivascular macrophages, microglial cells and mast cells, which are strategically positioned around brain vessels, further contribute to BBB disruption by induction and release of signaling molecules and proteases that promote vascular permeability.

A comparative study of the effects of recombinant interleukin-1 beta (IL-1 β) injected into the brain parenchyma of adult, juvenile and newborn rats showed that while in the adult intense meningitis and disruption of the BCSFB, but not leukocyte recruitment, occur within 4 h, in the juvenile rats, a 500-fold lower IL-1 β dose causes a large BBB leakage and neutrophil recruitment into the tissue around the injection site within the same time frame [30]. In the newborn rat (2-h-old rat), a similar low IL-1 β dose gives rise to an increase in permeability in the meninges, but no increase in BBB permeability. The IL-1 β -induced increases in vessel permeability in the meninges, parenchyma, and choroid plexus are neutrophil-dependent [30]. These findings demonstrate clear differences in the age-specific response of the BBB and barrier-type specific responsiveness to IL-1 β . Another study utilizing prolonged systemic inflammation in early development in the rat showed induction of long-term changes in BBB permeability that ultimately led to white matter lesions [31]. LPS given at postnatal day 0 (P0)-P8 induced BBB permeability to small (sucrose and inulin) and large (protein) molecules during and immediately after the inflammatory response. Permeability to protein was increased only transiently whereas increased permeability to ¹⁴C-sucrose and ¹⁴C-inulin was more persistent, demonstrating size selective increases in BBB in the early postnatal period in response to prolonged systemic inflammation [31].

5 Models of Neonatal Focal Stroke and Hypoxia–Ischemia

While models of stroke in adult rodents have existed for several decades, models of perinatal focal arterial stroke have been established relatively recently. These models include permanent MCA ligation [32] and suture tMCAO in P7–P10 rats [33, 34] and P9 mice [35]. Varying the duration of tMCAO has allowed for induction of injuries of different severity [35] and MRI-based confirmation of recirculation following suture retraction in the tMCAO model [36] has allowed for studies of reperfusion, which frequently occurs in arterial stroke in term infants. The vast majority of experimental data related to ischemia-induced brain injury in neonates this far have been obtained using a hypoxia–ischemia (HI) model in P7–P9 rats and mice, a group of models that more closely mimic hypoxic-ischemic encephalopathy in term human babies than focal arterial stroke [37–40]. We and others recently reviewed the most important pathophysiological findings in the neonatal HI models [39, 41–43]. Thus, we will limit our discussion to available data on stroke and HI-induced changes in BBB permeability in the neonate [44, 45] and effects of neurovascular changes for injury and repair [46–48].

6 BBB Integrity After Neonatal Focal Stroke and Hypoxia–Ischemia

The data on BBB function after ischemia-related brain injury in the neonate are rather scant. Considering that endothelial TJs are already present during early embryonic development [49], specific BBB transporters are present in the brain endothelium during mid-gestation, with no fenestrations are observed at birth [50], that astrocytes and pericytes cover brain vessels in postnatal brain [5], at least to some extent, we examined whether neonatal stroke disrupts functional integrity of the BBB. To test if compared to adult stroke, the assumed more extensive magnitude of BBB disruption after stroke occurs in neonates, we determined BBB leakage in adult and P7 rats subjected to a 3-h tMCAO. We administered Evans Blue or fluorescent intravascular trace 3, 70 kDa dextran or TRITC-albumin at 24 h after reperfusion and observed significant extravasation of Evans Blue, 70kD and TRITC-albumin in injured regions of adult rats. Extravasation of these molecules was significantly lower following acute tMCAO in P7 rats, showing a strikingly better-preserved BBB integrity in injured neonatal brains [44]. We also used Gd-DTPA-enhanced T1W to further test BBB permeability in neonates and showed that contrast enhancement in injured regions was negligible at 24 h, within 10% [44]. Fig. 1a demonstrates increased leakage of 70 kDa dextran in injured adult brain and only minor leakage in injured neonatal rats. In contrast to tMCAO, permanent MCAO in P7 rats resulted in rapid BBB disruption and leukocyte extravasation [51], suggesting that

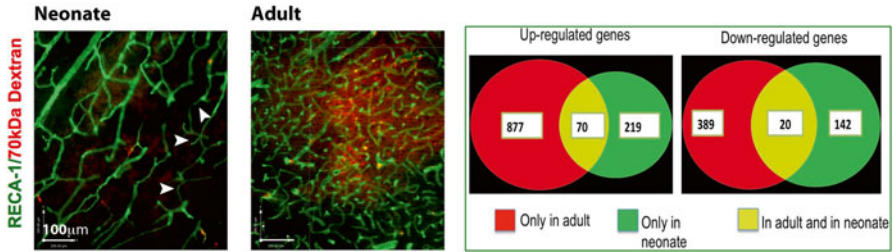


Fig. 1 Acute focal arterial stroke differentially affects BBB integrity in the adult and neonate. **(a)** Representative examples of the spatial distribution of intravenously administered TRITC-conjugated 70 kDa dextran in injured neonatal (*left*) and adult (*right*) brain regions 24 h after a 3-h tMCAO. **(b)** Differential effect of acute focal stroke on gene expression in endothelial cells of adult and neonatal rats. The endothelial transcriptome data were obtained in endothelial cells purified by sequential negative and positive selection from injured and contralateral tissue of adults and neonates 24 h after reperfusion. Shown are genes with >2-fold change in expression compared to that in contralateral hemisphere

persistent lack of cerebral microcirculation contributes to BBB collapse. Two studies that utilized tMCAO model in P10 rats observed increased BBB permeability during a sub-chronic injury phase, 72 h following tMCAO [52, 53]. The affected region was larger in spontaneously hypertensive pups [52] than in normotensive pups of the same age [53]. A recent study studied changes to the BBB in conjunction with CBF in a neonatal mouse HI model [45]. HI increased BBB permeability to small and large molecules, peaking at 6 h after the insult followed by normalization by 24 h. The opening of the BBB was associated with changes to BBB protein expression. Brain pathology was closely related to reductions in CBF during the hypoxia as well as the areas with compromised BBB [45]. Taken together, these data demonstrate a limited extent of BBB opening after ischemia-related injury in neonatal brain.

While the phenomenon of a better-preserved BBB integrity in neonatal stroke has been established, the underlying mechanisms are poorly understood. Comparative endothelial transcriptome data obtained in adult and neonatal rats subjected to tMCAO provided some mechanistic insight of age differences in BBB susceptibility to stroke. It appeared that, strikingly, the patterns of up- and down-regulated endothelial genes are largely non-overlapping between the two ages (Fig. 1b) [44]. Transcript levels of several adhesion molecules and ECM components were differentially affected by injury in immature and adult brain, including E-selectin and P-selectin. Gene expression of Mmp-9 was significantly upregulated in injured adults and, while high transcript levels of collagen type IV $\alpha 1$ (Col4a1) and Col4a2 remained unaltered in neonates, a significant increase of these two genes was evident in injured adult rats. Interestingly, transcripts of angiogenic regulators Vegfr-2 and Angpt2 were increased after stroke in adults but not in neonates [44]. Comparisons of protein expression of occludin, claudin-5, and ZO-1 between

adult and neonatal rats after tMCAO showed better-preserved expression in neonates than in adults [44]. Endothelial-ECM interaction via $\beta 1$ integrins regulates the expression of claudin-5 and BBB tightness whereas other ECM proteins, like galectin-3, mediate integrin-induced stabilization of focal adhesions and activate cytokine receptors to enhance actions of growth factors [54]. Laminin degradation occurs after focal stroke in adults and causes detachment of astrocytic end-feet, disrupts BBB, and induces ICH [55], while in neonates, expression of this ECM protein is not reduced acutely [44]. The role of other ECM proteins in injured neonates is less studied but the opposite effects of galectin-3 in adult stroke and HI have been demonstrated [56, 57]. Together, these data suggest that intrinsic developmental differences in basement membrane and ECM formation may contribute to a better-preserved BBB integrity after acute neonatal arterial stroke.

Astrocytes and pericytes may contribute differently to neonatal and adult stroke. In adults, astrocyte swelling, mediated by water channels (aquaporin-4, AQP-4) present in astrocyte end-feet, is one of the mechanisms involved in the generation of cytotoxic edema during the early phase after stroke [58]. Retraction of astrocyte end-feet from the parenchymal basal lamina of vessels and cellular redistribution and/or degradation of AQP-4 at later stroke stages are involved in BBB leakage and formation of vasogenic edema [59]. While coverage of vessels with astrocytic end-feet begins before birth, it continues to increase during the first postnatal week [5]. However, these particular mechanisms have not yet been specifically characterized after neonatal stroke.

7 Leukocyte and BBB Integrity After Neonatal Stroke

Interactions between activated endothelial cells and peripheral leukocytes contribute to BBB disruption after stroke. Compared to the adult, neutrophil infiltration in neonates is negligible after tMCAO [44]. Following HI, neutrophil infiltration was shown limited [60] or brief [61]. Neutropenia was shown beneficial when induced before, not after, HI [60]. The exact mechanisms that restrict neutrophil infiltration in the injured neonatal brain are not well understood. The different or uncoordinated patterns of expression of adhesion molecules and MMPs between neonatal and adult rodents may be of critical importance. Particulars of chemokine gradients between the brain and the blood may affect neutrophil extravasation in the neonate [44]. For example, cytokine-induced neutrophil chemoattractant (CINC-1) is a major chemoattractant for neutrophils. Neutralization of peripheral CINC-1 following tMCAO in the adult halted neutrophil transmigration, reduced brain edema and protected [62] but CINC-1 neutralization in neonatal rats subjected to tMCAO promoted neutrophil infiltration. Alterations in the blood–brain gradient of CINC-1 not only led to increased presence of neutrophils in the brain parenchyma but also increased BBB permeability and injury volume [44], suggesting that neutrophils mediate BBB damage in association with transmigration.

Compared to adult stroke, infiltration of circulating monocytes across the BBB was also low during the acute phase after stroke in the neonate [63]. The exact mechanisms of low monocyte infiltration in the ischemic neonatal brain are not completely understood, and it remains unclear whether the higher resistance of the neonatal BBB to stroke is a cause or a consequence of reduced transmigration. Monocytes have been recently shown to have a dual role, elicit both inflammatory effects and maintain BBB integrity following cerebral ischemia [64]. T and B cell infiltration may be less profound [61] or transient [51] in injured neonates than in adults [65, 66] but there have been no studies that examined effects for BBB integrity.

8 BBB Integrity, Angiogenesis and Brain Repair After Neonatal Stroke

Physiological angiogenesis continues during the first two postnatal weeks in the rat brain [44]. Endothelial cell proliferation and endothelial tip cells are abundantly present [67, 68]. Interestingly, following stroke in P7 rats, angiogenesis is essentially arrested in injured brain regions up to 14 days after injury, and only subtle angiogenic response is detected in the ischemic boundaries in the cortex [69]. Figure 2 summarizes these findings. Considering that in adults endothelial cell proliferation and vascular outgrowth have been reported as soon as 24 h after stroke [70–72],

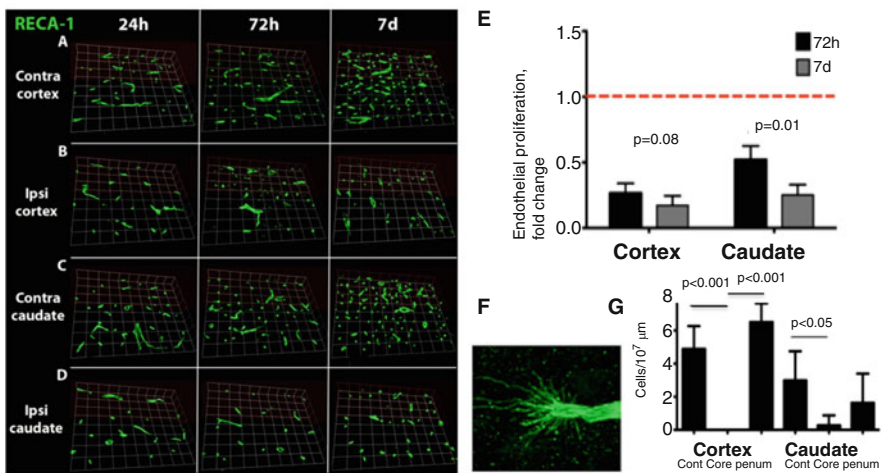


Fig. 2 Reduced vascular density and endothelial proliferation in sub-chronic injury phase after stroke in neonatal rats. **(a–d)** Examples of RECA-1+ vessel distribution in the ischemic core in the cortex **(a, b)** and caudate **(c, d)** at 24, 72 h, and 7 days after reperfusion. **(e)** Quantification of proliferating endothelial cells (BrdU+/RECA-1+) in the ischemic cortex and caudate. **(f)** Examples of PECAM-1+ endothelial tip cells (defined by extended filopodia). **(g)** Quantification of the number of PECAM-1+ endothelial tip cells in the contralateral and ischemic cortex and caudate

the response of neonatal brain to stroke differs to that in the adult in this aspect as well. Gain- and loss-of-function stroke studies in the adult have demonstrated that neuroblast migration occurs in association with remodeling of blood vessels [73], via a link between angiogenesis and neurogenesis within the “neurovascular niche,” and that blockage of angiogenesis abolishes neurogenesis after adult stroke [73]. Brain vessels with active endothelial proliferation in the ischemic boundaries of the injured regions 14 days after neonatal stroke showed abnormal expression of the endothelial barrier antigen (EBA) [69], a protein necessary for proper BBB function in adolescent and adult rats [74–76] and which expression is maturation-dependent [77, 78]. It is tempting to speculate that the relatively preserved BBB after neonatal stroke may negatively impact angiogenesis and account for a delay in angiogenesis and ultimate endogenous neurogenesis, but the relationships between the processes are still poorly understood.

Consistent with the findings in injured P7 rats, the extent of angiogenesis was also limited 1 and 2 weeks after tMCAO in P10 [46, 48]. Angiogenesis was low despite rapid induction of first neuronal and, then, of astrocytic VEGF [34]. Angiogenesis and neurogenesis were enhanced by delayed administration of rhVEGF and disrupted by VEGFR2 inhibition, demonstrating the role of VEGF signaling for the maintenance of angiogenesis and repair in the injured ischemic core [46, 48].

9 Conclusions and Future Directions

Successful development of therapeutics for neonatal stroke depends on accurate understanding of the neurovascular interface. Recent studies have improved our understanding of the events at the BBB after neonatal ischemic injury by revealing that the developmental step of the BBB at the time of ischemic insult contributes to the differing patterns of brain damage between neonates and adults and that careful consideration should be given about whether the BBB is in fact disrupted, allowing therapies to reach an injured neonatal brain. However, it is important to recognize that there are too many unknowns. We know little about relationships between BBB integrity and white matter maturation after stroke. No studies systematically examined whether these processes are sex-dependent. There have been no studies looking in depth at the effects of hypothermia on neurovascular integrity in injured neonatal brain. Future studies should also shed light on relationships between neurovascular integrity and interaction with neuroprogenitors, endogenous or engrafted, and migration and differentiation of neural progenitors during stroke-induced neurogenesis. Although BCSFB is critically important for functioning of a neonatal brain, there have been no studies on addressing functionality of this barrier in injured neonatal brain.

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Angiogenesis: A Realistic Therapy for Ischemic Stroke

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1 Introduction

Stroke is the fourth leading causes of death and the leading cause of adult disability in the Western society. Currently, thrombolytic therapy within a 3–4.5 h of narrow time window is the only acute therapeutic intervention for ischemic stroke, and development of effective therapies is urgently required [1–3]. During the past two and half decades, neuroprotection has been a major focus of ischemic stroke therapy research, yet it has also been most perplexing to investigators. All drugs that were neuroprotective in animal studies failed to show significant effectiveness in stroke patients, implying that solely focusing on neuroprotection is not sufficient and other methods for treating ischemic stroke should be explored. Thus, greater attention has been paid to the local environment of the surviving neuron, such as the cerebral microvasculature and other non-neuronal brain elements [4–8].

Angiogenesis is a normal physiological process in tissue growth and development that may also occur as a natural defense response against neurological diseases, such as stroke. Extensive animal studies have shown that post-ischemic angiogenesis plays an important role in the recovery of blood flow in affected brain tissue [9–14]. It is well known that angiogenic vessels in the ischemic boundary zone (IBZ) may contribute to recovery of tissue-at-risk by restoring metabolism in surviving neurons as well as provide the essential neurotrophic support to newly generated neurons. Robust increase of microvessel density has been observed in the penumbral areas, and the number of new angiogenic vessels is correlated with longer survival in stroke patients [15, 16], suggesting active angiogenesis may be beneficial for neurological functional recovery. These results strongly support inducible

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angiogenesis (drug-based) as a potential therapy for acute ischemic stroke. In addition to endogenous angiogenesis, therapeutic angiogenesis is a novel treatment for ischemic stroke. Through increasing blood flow in or around the ischemic area, in particular the penumbra which likely is salvageable only through quick efforts to restore tissue perfusion after stroke onset, the aim of therapeutic angiogenesis is to promote or improve tissue preservation and regeneration [9, 10, 17–19].

During stroke, ischemic insults rapidly trigger transcription of various genes and proteins that may be associated with the process of angiogenesis [9–14]. Among them, vascular endothelial growth factor (VEGF) acts as a central mediator in post-ischemic angiogenesis. Infusion of VEGF into brain tissue or overexpression of VEGF in transgenic mice has been documented to promote angiogenesis, decrease infarct volume, and reduce neurological deficits after focal cerebral ischemia [20–22]. In brain tissue, VEGF is produced and secreted by many neurovascular cells (neurons, astrocytes, and vascular endothelial cells) [9, 23–25]. VEGF from various cellular sources binds to its receptors on nearby vascular endothelial cells to directly initiate an angiogenic response. The binding of VEGF to its receptors (VEGFR-1 or -2) on the surface of endothelial cells activates intracellular tyrosine kinases and triggers multiple downstream signals (PI3K/Akt and MEK/ERK protein kinase pathways) that promote angiogenesis [10, 26]. Increased levels of VEGF have been identified in human brains and serum after ischemic stroke [27, 28].

Indeed, it has been well established that stimulation of angiogenesis can be therapeutic in ischemic heart or cerebrovascular disease, peripheral arterial disease, and wound healing. In the healthy condition, angiogenesis is strictly controlled by a dynamic balance between pro-angiogenic and anti-angiogenic factors. When this balance is disturbed, it usually causes pathological angiogenesis which results in increased blood-vessel formation in diseases that depend on angiogenesis. Accumulating endogenous pro-angiogenic molecules have been identified, including matrix metalloproteinases (MMPs), cytokines, integrins, and growth factors such as VEGF, fibroblast growth factors (FGF), transforming growth factors (TGF), epidermal growth factor (EGF), and angiogenesis-regulating microRNAs (Angiomirs) [9, 10, 13, 14, 19].

In this review article, we summarize the research progress describing the role and molecular regulating mechanisms of post-stroke angiogenesis. We also discuss the potential clinical applications of therapeutic angiogenesis as new treatment approaches for ischemic stroke.

2 Overview of Angiogenesis

Angiogenesis is defined as new vessels that are formed in the pathological and physiological process from preexisting vessels in a multistep process in order to deliver nutrients and oxygen to various organs and tissues [9], which is a key restorative mechanism in response to ischemia [29, 30]. Under ischemia condition, hypoxia is a key stimulus for angiogenesis and involved in the angiogenesis through the

activation of hypoxia-inducible factor-1 α [31, 32]. Besides, pro-angiogenic molecules such as vascular endothelial growth factor-A (VEGF-A) and VEGF receptor 2 (VEGFR-2), angiopoietins (Ang-1 and -2) and cognate receptor Tie-2, neuropilin-1, and basic fibroblast growth factor (bFGF) are also stimulated [26, 33]. These growth factors activate otherwise quiescent endothelial cells to start the angiogenic cascade. Among these factors, some of them act in a synergic manner but some in a direct-opposed manner. In most cases, angiogenesis exerts preventive effect in organs. For example, angiogenesis could significantly reduce ischemia injury after cerebral and myocardial infarction [34]. However, in other case, angiogenesis could accelerate tissue injury. For example, angiogenesis could exacerbate the injury of tumor tissues [35]. However, therapeutics which reduced angiogenesis is proved to improve tumor prognosis. In the healthy condition, angiogenesis is strictly controlled by a dynamic balance between pro-angiogenic and anti-angiogenic factors. Increased or decreased blood-vessel formation in diseases could be caused when this balance is disrupted.

3 Angiogenesis After Ischemic Stroke

Ischemic stroke occurs immediately after the supply of brain is stopped. Part of the natural recovery process after stroke is angiogenesis [36], which is a biological process involving the growth of new blood vessels from preexisting vessels [13, 14]. The formation of new blood vessels around the infarct may be important for restoration of adequate perfusion for brain tissue. Subsequently, re-establishment of the functional cerebral microvasculature network will promote stroke recovery. In infarction region, neurons in the core of infarction were usually dead. But many apoptotic neurons existed in the peri-infarction, which is called penumbra. Vascular angiogenic remodeling mainly occurred in the penumbra which was beneficial to reduce the apoptosis of neurons and subsequently improved the prognosis of stroke [37]. Thus, it is important for stroke recovery to restore penumbra after stroke.

Although angiogenesis is completely suppressed under normal physiological conditions in adult brains, studies from human and experimental stroke indicate that neovascularization is present in the adult brains after cerebral ischemia [13, 14, 38–40]. Accumulating studies have showed that promotion of angiogenesis was helpful in reducing infarction in experimental animals [41–43]. It has been reported that cerebral vascular endothelial cells start to proliferate in the peri-infarcted region as early as 12–24 h after the onset of stroke [38–40]. This ischemia/hypoxia-induced vascular remodeling leads to increased microvessel density surrounding the infarcted brain area 3 days following ischemic injury. Indeed, many factors such as hemeoxygenase-1 and VEGF have been proved to improve neurological outcome in MCAO animals by enhancing angiogenesis [44].

Consistent with basic research in experimental stroke animal models, it is demonstrated that angiogenesis also occurred in stroke patients. It has been previously reported that young stroke patients had a better prognosis than old stroke patients

[45]. This means angiogenesis is easy to occur in young patients and old patients have worse ability in vessels formation. Other studies showed that stroke patients who had a history of transient ischemic attack (TIA) had a better prognosis than those without TIA history [46]. It is speculated that signals for angiogenesis should be activated when TIA occurred. Mechanistically, stroke patients with greater cerebral blood vessel density appear to make better progress and survive longer than patients with lower vascular density, suggesting that active post-stroke angiogenesis is helpful for long-term neurological functional recovery. Of importance, angiogenesis usually took place after some time when stroke is onset in patients. Then, many neurons were dead especially in those with occlusion in large vessels.

4 Regulation of Angiogenesis After Ischemic Stroke

Since angiogenesis is important for the recovery of stroke, it is important to know how to regulate angiogenesis. Cerebral angiogenesis is strictly controlled by many key angiogenic factors that play an important role in the development of angiogenesis in response to various pathological conditions such as stroke [9–14]. These stroke-associated angiogenic factors include VEGF and its receptors [20, 21, 25, 40, 47], bFGF and its receptors [48, 49], platelet-derived growth factor (PDGF) and its receptors [50, 51], transforming growth factor beta (TGF β) [52, 53], MMPs [54], thrombospondin-1 (TSP-1) [55], angiopoietins and Tie receptors [56–58], endothelial nitric oxide synthase (eNOS) [59–61], hypoxia-inducible factor 1 [32], and many others [62].

In addition to the classic angiogenic regulators described above, recent studies have revealed important roles for microRNAs (miRs) in regulating angiogenesis [13, 14, 63–65]. MiRs are small endogenous RNA molecules (~21–25 nt) that repress gene translation by hybridizing to 3'-UTRs of one or more mRNAs in a sequence-specific manner. The discovery of miRs has shed light on how non-coding RNAs play critical roles in angiogenesis [13, 14, 63–65]. The initial support for the regulation of angiogenesis by miRs arose from genetic manipulations of the Dicer gene [66, 67]. Dicer knockout mice exhibit embryonic lethality because of abnormal vascular wall structure and arrangement. Mice with vascular-selective Dicer knockout have been reported to show impaired angiogenic ability, such as reduced endothelial tube formation and slowed EC migration, which may result from alterations of key angiogenesis-related genes. Moreover, an increasing number of individual miRs have been shown to regulate angiogenesis signaling pathways, thereby modulating endothelial migration, proliferation, and vascular-forming patterns [13, 14, 63–65]. Angiogenesis-related miRs can be classified into two groups with often opposing effects: pro-angiogenic and anti-angiogenic miRs. Among them, the miR-17-92 cluster, let-7, miR-27b, miR-126, miR-130a, miR-210, miR-296, miR-378, miR-21, and miR-31 exhibit pro-angiogenic effects, whereas miR-15/16, miR-424, miR-221/222, miR-92a, miR-320, miR-200b, miR-217, miR-503, miR-34, and miR-214 are considered anti-angiogenic [13, 14, 63–65].

Accumulating evidence has shown that miRs play a critical role in the pathogenesis of ischemic stroke [13, 14, 68–74]. Altered cerebral miR profiles have been reported in rodent focal and global cerebral ischemia models [75–78]. Ischemic preconditioning was also shown to change miR expression, including miR-132 [79], the miR-200 family, miR-182 family [80], and others [81], which may promote ischemic tolerance via neuroprotective signaling pathways. Altered miR levels were also found in blood samples of rodent stroke models [76, 78] and stroke patients [82–84] and can serve as potential biomarkers. We and others are among the first to identify the function of individual miRs in stroke pathology. Notably, we have defined that miR-497, a miR-15a homolog, is dramatically induced and promotes ischemic neuronal death *in vitro* and *in vivo* by inhibiting *bcl-2* and *bcl-w* [85]. Moreover, we provided the first evidence that inhibition of endothelial miR-15a contributes to the PPAR δ -mediated cerebral vasoprotective role in stroke [86]. Other miRs including miR-145 [75], miR-320a [87], miR-21 [88], miR-223 [89], miR-23a [90], miR-181 [91, 92], *Let7f* [93], miR-331 [94], miR-146a [95], miR-199a-5p [96], miR-124a [97], miR-124 [98], miR-133b [99], miR-29c [100], and miR-17-92 cluster [101] also mediate ischemia-induced neuronal death, neurogenesis, and neurological recovery.

Although many angiogenesis-regulating miRs play a role in development, cancer, and cardiovascular diseases, none of them have been systematically studied in the ischemic brain [13, 14]. In animal experiments, our group found that vascular endothelial cell (EC)-enriched miR-15a/16-1 cluster, a kind of stroke-associated miR, could suppress tube formation, cell migration, and cell differentiation in ECs [43]. Moreover, we found that EC-selective miR-15a transgenic overexpression in mice led to reduced blood vessel formation and local blood flow perfusion in the ischemic hindlimbs at 1–3 weeks after hindlimb ischemia [43]. Of note, we found that miR-15a exerts its anti-angiogenic effect by directly inhibiting FGF2 and VEGF activities [43]. This means miR-15a is an anti-angiogenic miR. A very recent study showed that miR-107 was strongly expressed in IBZ after permanent middle cerebral artery occlusion in rats and inhibition of miR-107 could reduce capillary density in the IBZ after stroke [102]. Mechanistically, they found that miR-107 could directly downregulate *Dicer-1*, a gene that encodes an enzyme essential for processing miRNA precursors. This resulted in translational inhibition of VEGF mRNA, thereby increasing the expression of endothelial cell-derived VEGF (VEGF₁₆₅/VEGF₁₆₄), leading to angiogenesis after stroke. Using HUVECs and MCAO rat, Li and his colleagues showed that miR-376b-5p repressed angiogenesis *in vivo* and *in vitro*, and miR-376b-5p inhibited angiogenesis in HUVECs by targeting the HIF-1 α -mediated VEGFA/Notch1 signaling pathway [103]. Other groups found that miR-210, a hypoxia-induced miR, is significantly upregulated in adult rat ischemic brain cortexes in which the expression of Notch1 signaling was also increased. Gain-of-miR-210 function in cultured HUVE-12 cells caused activation of the Notch1 signaling cascade and induced endothelial cells to migrate and form capillary-like structures. These data may imply that miR-210 is a kind of pro-angiogenic miR [104]. However, this study does not establish a causal link between miR-210 and post-stroke cerebral angiogenesis. Consistent with this finding,

miR-210 has also been shown to regulate angiogenesis in normal brain tissue [105]. Besides brain miRNAs, many circulating miRNA has close relationship with ischemic stroke. For example, previous studies showed that plasma miRNA such as miR-124 and miR-290 were increased from 6 to 24 h in stroke animals [76, 106]. These data showed the potential biomarker of miRNA for onset of ischemic stroke and the potential role of brain-specific miRNAs to serve as biomarkers of tissue injury in MCAO animals.

In human subjects, it is demonstrated that many circulating miRs are associated with vascular endothelial function and angiogenesis and vascular remodeling under different ischemic conditions in stroke patients. In young stroke patients, circulating plasma miRs are associated with vascular endothelial function and angiogenesis (hsa-let-7f, miR-130a, -150, -17, -19a, -19b, -20a, -222, and -378) and vascular remodeling (miR-21, -126, and -150) have been found to be differentially regulated under ischemic conditions. Similarly, miRs that are expressed in hypoxic conditions (miR-23, -24, -26, -103, -107, and -181) and cardiac ischemia/reperfusion (miR-15, -16, -21, -23a, -29, -30a, -150, and -195) have also been detected in blood samples from stroke patients [82]. These miRNA plays an important role in angiogenesis, neuroinflammation, immune activation, leukocyte extravasation, and thrombosis [107]. The expression of certain miRNA may be different in different types of stroke. For example, it has been reported that circulating let-7b decreased in patients with large-vessel atherosclerosis than healthy controls, but it increased in patients with other kinds of ischemic stroke [107]. Moreover, circulating miRNA even could be used to predict the outcome of patients. Decrease of miR-210 was associated with poor outcome [108]. Conversely, reduction of anti-miR such as miR-30 predicts good outcome in stroke patients.

In addition to miRNAs, it has been demonstrated that cerebral ischemia rapidly changes the expression profiles of other classes of ncRNAs, such as lncRNAs and piRNAs, and imply that manipulation of these ncRNAs may regulate post-ischemic cerebrovascular remodeling. Till now, little is known regarding the expression and functional role of cerebral vasculature and/or endothelium lncRNAs and piRNAs after ischemic stroke [13, 14].

5 Functional Roles of Angiogenesis in Stroke Long-Term Remodeling and Functional Recovery

Angiogenesis, neurogenesis, and synaptogenesis could be occurred simultaneously, and function synergistically to improve long-term brain remodeling and promote functional recovery after ischemic stroke. It has been reported that NF- κ B could enhance both angiogenesis and synaptogenesis in experimental stroke models [109]. Moreover, Avraham et al. found that following stroke, administration of exogenous leptin significantly induces neural stem cells in the cerebral cortex near the lesion site [110]. An increase in the number of newborn neurons and glia were observed in

leptin-treated animals as well. Leptin treatment also significantly increased the density of brain blood vessels in the penumbral cortex. In addition to neurogenesis and synaptogenesis, angiogenesis could also promote oligodendrogenesis, evidenced by a recent report by Zhang and colleagues that recombinant human erythropoietin could increase differentiation of SVZ neural progenitor cells into oligodendrocytes and effectively enhance neurological outcomes in stroke rats [111].

Accumulative studies have shown that angiogenesis appears to improve long-term functional recovery by promoting brain remodeling, repair, and plasticity. Cerebral injection of AAV-Netrin-1 gene has been shown to effectively improve neurological outcomes in stroke rats during 1–4 weeks [112], with a simultaneous increase in newly generated blood vessels after treatment. Striking findings from another group documented that BSc2118, a proteasome inhibitor, could enhance angiogenesis and improve functional outcomes even at 3 months after ischemic stroke [113]. Consistent with the described-above findings, a recent study also showed that fat-1 transgenic mice that cause overexpression of omega-3 polyunsaturated fatty acids (*n*-3 PUFAs) exhibited long-term histological and behavioral protection against ischemic stroke [114]. Mechanistically, *n*-3 PUFAs induced upregulation and release of a pro-angiogenic factor, angiopoietin 2 (Ang 2), in astrocytes after ischemic stimuli both *in vitro* and *in vivo*. Ang 2 facilitated endothelial proliferation and barrier formation *in vitro* by potentiating the effects of VEGF on phospholipase C γ 1 and Src signaling. Blockade of Src activity with a Src inhibitor AZD0530 impaired *n*-3 PUFA-induced angiogenesis and exacerbated long-term neurological outcomes in fat-1 Tg mice following ischemic stroke. These findings provide strong evidence that post-stroke cerebral angiogenesis contributes to long-term functional recovery in experimental stroke models.

6 Therapeutic Angiogenesis in Ischemic Stroke

Therapeutic angiogenesis is to employ clinical methods to enhance or improve blood vessels growth or development of collateral blood vessels within the ischemic tissue. Currently, there are three major ways to promote angiogenesis, namely, pharmacological/genetic approaches, physical activity, and cell therapy (Fig. 1).

Since cerebral neovascularization is involved in the pathogenesis of stroke, it could be helpful in the recovery of cerebral injury by modulating angiogenesis. Indeed, cumulative studies showed that angiogenesis is a therapeutic target of ischemic stroke [115]. Many pharmacological/genetic approaches could be used to serve as modulating angiogenic factors for angiogenesis in ischemic stroke. These factors reduced ischemic injury and improved neurologic outcomes via promoting angiogenesis under ischemic conditions. There are two main kinds of delivery way, namely direct delivery and virus-gene-associated delivery. Accumulating studies showed that many angiogenic factors such as VEGF [116], angiopoietin 1 [117], fibroblast growth factor (FGF-2), tumor necrosis factor α [118], hepatocyte growth factor (HGF) [119], bFGF [120], heparin-binding epidermal growth factor-like

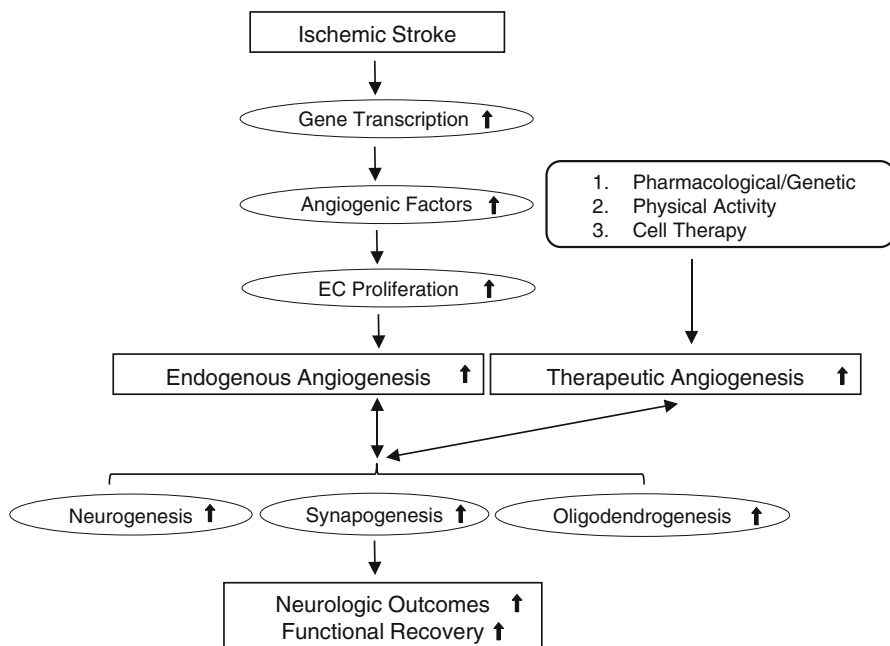


Fig. 1 Schematic representation of the post-stroke angiogenesis and therapeutic angiogenesis after ischemic stroke. Ischemic stimuli in brains trigger gene transcriptions and translations of pro-angiogenic factors, which subsequently induce endothelial cell proliferation and promote endogenous angiogenesis. Newly generated brain microvessels function synergistically with cerebral neurogenesis, synaptogenesis, and oligodendrogenesis to perform long-term neural/vascular remodeling and improve functional recovery after ischemic stroke. Therapeutic angiogenesis via pharmacological/genetic approaches, physical activity, and cell therapy further enhance post-stroke angiogenesis, thereby resulting in therapy-promoted functional recovery after ischemic brain injury

growth factor (HB-EGF), granulocyte-colony stimulating factor (G-CSF) could also decrease ischemic brain damage in animal models [9, 10]. Besides the direct effect of angiogenic factors, some proteins could enhance the expression of angiogenic factors and consequently improve ischemic stroke. For example, Zan and his colleague found that phosphorylation of Src at Y418 was increased as early as 3 h and peaked at 6 h following ischemic injury, after decreasing, it peaked again at 3–7 days [121]. Increases in Src mRNA and phosphorylation correlated positively with levels of VEGF and Ang-2, and negatively with levels of angiopoietin-1 (Ang-1) and zonula occludens-1. More importantly, increase in angiogenic factors was helpful in reducing vascular permeability, cerebral infarct size, and neurologic dysfunction. Moreover, many pharmacological approaches, including phosphodiesterase type-5 (PDE-5) inhibitors, statins, adrenomedullin, erythropoietin, kallikreins, nitric oxide donors, angiotensin II type 1 receptor blockade, have also shown to increase cerebral angiogenesis around the infarcted area and improved functional recovery in rodent stroke models [10].

Additionally, it has been reported that exercise could induce angiogenesis and improve motor deficit in animal models of stroke. For instance, Ma et al. reported that exercise enhanced matrix metalloproteinase 2 (MMP2) and VEGF-related genes and proteins expression while improving regional cerebral blood flow and neurobehavioral score [122]. Another group found that pre-ischemic exercise reduces brain injury in rats subjected to 2 h MCA occlusion followed by 48 h reperfusion via upregulating angiogenic factors such as VEGF [123]. Also Gertz et al. [124] demonstrated that continuous voluntary running in stroke mice led to long-term upregulation of eNOS in the vasculature and of EPCs, which was associated with higher numbers of circulating EPCs and an enhanced cerebral neovascularization, and resulted in an improved functional outcome and increased density of perfused microvessels.

Cell therapy is a novel method in which many angiogenic factors-producing cells are used to promote angiogenesis in ischemic tissue [18, 125]. A variety of cell types have been used for the purpose of therapeutic angiogenesis in experimental stroke models such as monocytes, endothelial progenitor cells, marrow stromal cells, mesenchymal stem cells, umbilical cord blood, and neural stem cells [10, 18]. Mechanistically, these cells are able to stimulate angiogenesis at the ischemic boundary region through direct expression and release of pro-angiogenic factors induced by local paracrine stimuli from ischemia. These cells can also act as a shuttle to transfer an angiogenic gene toward ischemic brain tissue [10, 18].

In summary, recent therapeutic angiogenesis through pharmacological/genetic delivery of various angiogenic factors, physical activity, and cell therapy has shown us several promising results in promoting cerebral neovascularization, blood flow recovery, and neurological function recovery in ischemic stroke [10, 18]. Although selective regulation of post-stroke cerebral angiogenesis appears to be a promising restorative therapy for ischemic stroke, there are several potential challenges or limitations for application of therapeutic angiogenesis in stroke. These obstacles include incomplete understanding of the function and underlying mechanisms of cerebral neovascularization in the stroke pathogenesis, brain edema, ineffective delivery system to achieve cell-specific delivery in vivo, limited target specificity, inflammatory responses, vascular malformation, hypotension, and off-target tumorigenesis [17, 18]. Therefore, a better understanding of the cellular and molecular signaling cascades of post-stroke cerebral neovascularization is needed for successful application of angiogenesis-based neurorestorative therapies after ischemic stroke.

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Part II
Glial Cells in Stroke

Astrocytes as a Target for Ischemic Stroke

Shinghua Ding

1 A Brief Overview of Astrocytes

Astrocytes are star-like cells and the most diverse glial cell type in the central nervous system (CNS). The classical definition of astrocytes is based on morphology and specific protein markers. Although there are many types of astrocyte in a normal brain, our discussion in this chapter will focus on two major types of astrocytes in the adult brain: Fibrous astrocytes present in white matter tracts such as the corpus callosum and protoplasmic astrocytes present in gray matter such as the cortex. Kimelberg proposed the following eight criteria to define mature astrocytes [1]:

1. Electrically non-excitabile.
2. A very negative membrane potential determined by the transmembrane K^+ gradient.
3. Expression of functional transporters for glutamate and GABA uptake.
4. A large number of intermediate filament bundles, which are the sites of the astrocyte-specific protein glial fibrillary acidic protein (GFAP).
5. Glycogen granules.
6. Processes from each cell surrounding blood vessels.
7. Many more processes from each cell surrounding synapses.
8. Linkage to other astrocytes by gap junctions consisting of connexins 43 and 30.

Astrocytes are traditionally considered as the housekeeping cells needed to maintain homeostasis in CNS. Growing evidence suggests that astrocytes also play an active role in modulating neuronal function. Protoplasmic astrocytes exhibit the following functions under normal conditions [1, 2]:

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1. Extracellular K^+ buffering.
2. Control of extracellular H^+ and brain pH.
3. Uptake of glutamate and GABA with their transporters.
4. Mobilizing intracellular Ca^{2+} stores by activation of G-protein coupled receptors (GPCRs) such as mGluR5 and P2Y.
5. Control of cerebral blood flow.
6. Control of water transport by aquaporins.
7. Astrocyte-neuron lactate shuttle.
8. Modulation and control of synaptic activity.

Many astrocytes conform to the aforementioned criteria and functional roles, but astrocytes are heterogeneous in morphology, molecular expression, and physiological function [3, 4]. Morphologically, protoplasmic astrocytes are highly branched. A protoplasmic astrocyte in gray matter has several primary processes each with elaborated sub-branched fine process arborizations to form a bush-like astrocytic territory. Protoplasmic astrocytes exhibit distinct territorial domain with little overlap between neighboring astrocytes [5, 6]. Fibrous astrocytes have thicker and less branched processes. They exhibit a high degree of overlap. Intermediate filament protein GFAP is primarily expressed in the thick main processes in astrocytes and has been considered as a “pan-astrocyte” marker; however, its expression levels are quite different between fibrous and protoplasmic astrocytes. GFAP is expressed in almost every individual fibrous astrocyte, but protoplasmic astrocytes in the cortex express much lower levels of GFAP than do protoplasmic astrocytes in the hippocampus [7]. Recent transcriptome study revealed that the *Aldh1L1* gene is most widely and homogeneously expressed in astrocytes, and immunostaining revealed that the *Aldh1L1* protein is highly expressed in the cell body and extensive processes of an astrocyte [8]. Therefore, *Aldh1L1* is now considered as a new “pan-astrocyte” marker.

Astrocytes exhibit different electrophysiological properties. Astrocytes can be classified as outward rectifying astrocytes and variably rectifying astrocytes based on current–voltage relationship [9]. Two-photon (2-P) microscopy revealed that astrocytes also exhibit different properties of Ca^{2+} signaling *in vivo*; for example, astrocytes in the cortical layer 1 (L1) nearly double the Ca^{2+} activity compared to the astrocytes in L2/3 in anaesthetized rats; moreover, Ca^{2+} signals in the processes in the same astrocyte are asynchronous in L1 while those in L2/3 are more synchronous [10]. The morphological, molecular, and functional heterogeneity of astrocytes indicates that astrocytes play diverse and complex physiological and pathological roles in the CNS.

Astrocytes are electrically non-excitabile, but their excitability is manifested by Ca^{2+} signaling through the activation of various GPCRs. Astrocytes express GPCRs for glutamate, γ -aminobutyric acid (GABA), ATP, serotonin, norepinephrine, and dopamine. Astrocytes can mediate Ca^{2+} signaling and intercellular waves *in vivo* by the activation of metabotropic glutamate receptors (mGluRs) [11–13], P2Y receptors [13–17], GABA_B receptors (GABA_BRs) [14, 18], noradrenergic receptors [19], and dopamine receptors [20, 21]. GPCR stimulation activates phospholipase-C (PLC) with subsequent IP₃R-mediated Ca^{2+} release from internal stores in endoplas-

mic reticulum (ER) [22–24]. Among the three types of IP₃R (IP₃R1-3), IP₃R2 seems to be the predominant type in astrocytes in the rodent brain [25–27]. IP₃R2 knock-out (IP₃R2 KO) mice do not exhibit GPCR agonists-evoked Ca²⁺ release in astrocytes in brain slice and in vivo, demonstrating that IP₃R2 is a key mediator of intracellular Ca²⁺ release in astrocytes [24, 28, 29]. Ca²⁺ signaling is now considered as a primary form of cellular excitability in astrocytes that can be measured by fluorescent imaging.

2 Astrocytes in Ischemic Stroke

Focal ischemic stroke (FIS) is initiated by the mechanical occlusion of cerebral vessels by thrombus or embolus thereby reducing or blocking blood flow; it is also called focal cerebral ischemia. FIS accounts for approximately 80 % of all human strokes and causes brain infarction and human disability. It is the leading cause of human death and disability and has a major impact on public health. The degree of blood flow reduction largely determines the brain infarction. Due to the lack of glucose and oxygen in the ischemic region, synthesis of ATP through glycolysis and oxidative phosphorylation is impaired. Thus, ischemia results in energy depletion and subsequent loss of ionic gradients and membrane potential depolarization in neurons and astrocytes. This further induces the release of glutamate and other neurotransmitters from presynaptic terminals to the extracellular space. This energy depletion also contributes to the malfunction of astrocytic and neuronal glutamate transporters, which are responsible for the clearance of the glutamate released into the synaptic cleft.

Cells in the ischemic core region die rapidly from the onset of ischemia due to severely impaired energy production and the ensuing breakdown of ionic homeostasis, but in the penumbra, the region surrounding the core of a focal ischemic locus, the brain tissue is hypoperfused with collateral blood flow with partial preservation of energy metabolism. This region can completely or partially recover after reperfusion, but can progress to infarction without treatment due to the ongoing excitotoxicity [30], thus the primary goal for stroke therapy is to salvage the penumbra in the acute phase of ischemic stroke. Clinically, current strategies to treat ischemic stroke are aimed at restoring blood supply by administration of thrombolytic drugs, which is only effective within a narrow window of about 3–4 h following the onset of ischemia [31, 32]. Thus, identifying novel therapeutic targets and elucidating cellular and molecular mechanisms by which ischemia induces neuronal death and brain damage are of great importance to provide effective therapeutic avenues.

Astrocytes respond to almost all neural diseases including FIS; however, compared with neurons, the role of astrocytes in ischemia is less understood. After the onset of ischemia, astrocytes undergo numerous pathological processes [33–35]. Astrocytes rapidly swell after ischemia, which is contributed to by a number of factors [36–38]. Swelling spreads from the ischemic core to the penumbral tissue. Many of swollen astrocytes in the ischemic core eventually lyse, while astrocytes in the penumbral region exhibit reversible swelling [37]. Astrocytes are generally

more resistant to ischemic insults than neurons [39] primarily due to utilization of glycogen stores as an alternative energy source [30, 40–42]. Astrocytes can supply energy to neurons through lactate shuttle [43]. Astrocytes can also convert glycogen to glucose-1P and produce ATP by glycolysis [41]. Astrocytes become reactive over time after ischemia, a process called reactive astrogliosis [33, 44–47]. Reactive astrocytes eventually form a glial scar around the ischemic core. Some reviews have provided various aspects of astrocytes in stroke and therapeutic potential [35, 48–51]. This chapter focuses on discussing the Ca^{2+} signaling in astrocytes in the acute phase of ischemia, dynamic changes of morphology and proliferation in the sub-acute phase, signaling pathways of reactive astrogliosis, stem cell properties of reactive astrocytes, and the potential of astrocytes as a target in stroke therapy.

2.1 Astrocyte and Glutamate Uptake in Ischemic Stroke

Glutamate excitotoxicity is considered the primary mediator of acute neuronal death [52]. The excessive accumulation of extracellular glutamate overstimulates ionotropic and metabotropic glutamate receptors, especially ionotropic *N*-methyl-D-aspartate (NMDA) receptors (NMDARs), and results in pathological overloading of intracellular Ca^{2+} in neurons [53, 54], which is linked to the triggering of activations of downstream phospholipases and proteases that cause degradation of membranes and proteins and eventually neuronal death [39, 55]. Thus, glutamate excitotoxicity not only causes acute neuronal death, but it also initiates molecular events that lead to delayed neuronal death and brain damage [55]. It has been known that glutamate transporters in astrocytes are impeded due to energy depletion. Glutamate can even be released from astrocytes by the reversed operation of transporter in ischemia [56, 57], leading to elevation of extracellular glutamate and aggravating excitotoxic neuronal death. From study using cortical cultures, neuronal vulnerability to glutamate is 100-fold greater in astrocyte-poor cultures than in astrocyte-rich cultures. Knockdown of glial glutamate transporter (GLT-1) significantly increases neuronal death in a global ischemia mouse model [58]. These studies demonstrate that astrocytes play an important role in reducing glutamate excitotoxicity in ischemia through uptaking glutamate by their glutamate transporters. Thus, enhancing the glutamate transporter function is a promising therapeutic strategy to reduce acute neuronal death and brain damage after ischemic stroke.

2.2 Ca^{2+} Signaling in Astrocytes After Ischemic Stroke

Neuronal glutamate and Ca^{2+} excitotoxicity is widely acknowledged [52, 59]; however, whether and how astrocytic Ca^{2+} is altered after ischemia has been less understood. Since astrocytic Ca^{2+} signaling is the signature feature of astrocytic excitability and due to the tripartite nature of synapse, Ca^{2+} -dependent gliotransmitter release is

expected to affect neuronal excitotoxicity during a stroke. Several studies using *in vitro* brain slice and *in vivo* animal models reported astrocytic Ca^{2+} dysregulation after ischemia.

Using acute brain slices which can mimic tissue environment, Duffy and MacVicar [60] found that a short episode (5 min) of simultaneous hypoxia and hypoglycemia can induce intracellular Ca^{2+} increase in astrocytes in acute brain slices within an average of 7.5 min. Their report also recorded 2.5 min as the time needed to reach a peak. Ca^{2+} levels in astrocytes remained elevated for a variable period of time ranging from several minutes to 1 h after reoxygenation. Astrocytic Ca^{2+} elevation could be still detected in the absence of extracellular Ca^{2+} albeit with a relative consistent duration. They further showed using electrophysiological recordings that hypoxia and hypoglycemia can also depolarize astrocytes. Their study suggests that astrocytes can mediate Ca^{2+} increase from the internal store release and influx of a voltage-dependent Ca^{2+} channel.

Dong et al. showed that OGD can induce slow inward currents (SICs) mediated by extrasynaptic NMDA receptors in rat CA1 pyramidal neurons in brain slices [61]. SICs can be inhibited by dialysis of the Ca^{2+} chelator BAPTA into astrocytic network, indicating that the activation of extrasynaptic NMDA receptors depended on astrocytic Ca^{2+} activity. Using 2-P microscopy, it was found that frequent astrocytic Ca^{2+} elevations were observed during OGD, with over 60 % of astrocytes displayed detectable Ca^{2+} elevations within the 10 min of OGD. In addition, most astrocytes displayed more than two transients. They further demonstrated that astrocytic Ca^{2+} elevations and the frequency of SICs during OGD were largely reduced in $\text{IP}_3\text{R2}$ KO mice as compared with wild type (WT) mice.

Ding et al. for the first time determined whether astrocytes exhibit altered Ca^{2+} signaling *in vivo* after ischemia [14]. Using 2-P microscopy, they imaged astrocytic Ca^{2+} signals after photothrombosis (PT)-induced focal ischemia in urethane-anesthetized adult mice. They found that astrocytes exhibit enhanced Ca^{2+} signaling characterized as intercellular Ca^{2+} waves, which starts ~20 min after PT, and Ca^{2+} signals reach the plateau 60 min after PT. Both amplitude and frequency of PT-induced astrocytic Ca^{2+} signals were dramatically increased compared to relative quiescent Ca^{2+} signaling prior to PT. Most Ca^{2+} signals were initiated and returned to the basal level at the same time among astrocytes in the imaging field, i.e., they were highly synchronized transient signals. To further determine the nature of astrocytic Ca^{2+} signals in ischemia, antagonists for GPCRs including mGluR5, GABA_BR , and P2Y receptors were administered after emergence of Ca^{2+} signals. Both 2-methyl 6-(phenylethynyl)pyridine hydrochloride (MPEP), an antagonist of mGluR5, and CGP54626, the antagonist of the GABA_B receptor, significantly reduced Ca^{2+} signals (~55 %). But suramin, a non-specific inhibitor for P2Y receptors, did not reduce the PT-induced Ca^{2+} signals. A general P2 receptor antagonist, pyridoxalphosphate-6-azophenyl-2',4'-disulfonic acid (PPADS), did not attenuate the astrocytic Ca^{2+} signal either. Similarly, inhibition of adenosine receptor A1 did not inhibit PT-induced astrocytic Ca^{2+} signals. Thus, the pharmacological study suggests that glutamate and GABA are likely to be released to the extracellular space following PT to stimulate intercellular Ca^{2+} waves in the astrocytic network through

mGluR5 and GABA_BR. It is surprising that neither P2Y nor A1 receptors contribute to the enhanced Ca²⁺ signaling after PT, presumably due to the rapid degradation of ATP and adenosine by enzymes such as ectonucleotidases after ischemia [62]. Furthermore, application of BAPTA-AM significantly reduced brain infarct 24 h after PT. The results represent the first *in vivo* study of astrocytic Ca²⁺ responses after ischemia and suggest that a blockade of Ca²⁺ increase in astrocytes can reduce the release of glutamate from astrocytes and cause neuronal death and brain damage. The emergence of Ca²⁺ increase in the acute phase might represent the initial response of astrocytes to ischemic insults. In future study, it is important to determine the time course of neuronal and astrocytic Ca²⁺ signaling and overloading after ischemia as neuronal Ca²⁺ overloading is one of the major events that cause excitotoxicity.

Astrocytes also exhibit altered Ca²⁺ signaling in the chronic phase of ischemia. Winship et al. studied the neuronal and astrocytic Ca²⁺ signaling and functional rewiring in somatosensory neurons in the ipsilateral hemisphere after PT [63]. They examined astrocytic Ca²⁺ responses in anesthetized mice to study whether the prevalence of these responses changed in the peri-infarct cortex half, 1 and 2 months after PT. A significant increase was found in the prevalence of responses to preferred limb stimulation. The selectivity of astrocyte responses in regions with overlap between contralateral hindlimb- and contralateral forelimb-evoked intrinsic optical signal and regions without intrinsic optical signal overlap were also significantly different. These results suggest that mechanisms for rapid neuron–astrocyte communication are preserved, or even enhanced, after stroke in the penumbra.

Astrocytes also play a role in the functional recovery within the contralateral hemisphere. Takatsura et al. performed *in vivo* Ca²⁺ imaging to examine the neuronal and astrocytic Ca²⁺ responses in the region contralateral to the stroke site at different times following PT [64]. Their results showed that the number of astrocytes with a Ca²⁺ response to limb stimulation significantly increased in the contralateral somatosensory cortex responding to ipsilateral limb stimulation during the first and second week after infarction as compared with the sham group. A significantly larger number of astrocytes responded only to the single-limb stimulation in the sham group as compared with stroke groups, but a smaller number of astrocytes responded to multiple-limb stimulation in the sham group as compared to the stroke groups. Interestingly, unlike neurons, astrocytes showed no preference in response to contralateral and ipsilateral limb stimulation. The amplitude of Ca²⁺ response was also increased in stroke groups as compared with sham groups. A large increase in glutamate concentration were observed 2 weeks after the stroke compared with the sham and 1-week group based on a microdialysis study; however, glutamine concentration was much higher in the contralateral side in the 1-week group than in the sham and 2-week group. Astrocytic plasma membrane glutamate transporter 1 (GLT-1) may contribute to the uptake of glutamate in the 1-week groups. These findings demonstrate that activated astrocytes increased the uptake of glutamate by glutamate transporters during the first week, and indicate that astrocytes play an important role in functional recovery and cortical remodeling in the area contralateral to ischemic lesion in the post-ischemic period.

IP₃R2 knockout mice exhibit smaller infarct volume than WT littermates in acute ischemia [61, 65] as well as in chronic phases of ischemia [65], indicating that IR₃R2-mediated Ca²⁺ signaling contributes to neuronal excitotoxicity and acute brain damage after ischemia. IP₃R2 KO mice also exhibited less neuronal apoptosis, reactive astrogliosis, and tissue loss than WT mice [65]. The study from Li et al. further shows that IP₃R2 KO mice exhibited reduced functional deficits after PT using behavioral tests, including cylinder, hanging wire, pole and adhesive tests [65]. These studies demonstrate that disruption of astrocytic Ca²⁺ signaling has beneficial effects on neuronal and brain protection and functional deficits after stroke by reducing Ca²⁺ dependent glutamate release and revealed a novel non-cell-autonomous neuronal and brain protective function of astrocytes in ischemic stroke, whereby suggest that the astrocytic IP₃R2-mediated Ca²⁺ signaling pathway might be a promising target for stroke therapy.

Astrocytic Ca²⁺ signaling may play different roles in neuronal death and brain damage in different brain injury models. In a tiny ischemic lesion model generated by a single vessel laser irradiation, an increase in astrocytic Ca²⁺ can stimulate energy metabolism and ATP production in astrocytic mitochondria and, thus, reduce brain damage in WT mice as compared with IP₃R2 KO mice [66]. In the stab wound injury (SWI) model which also produces a tiny lesion, IP₃R2 KO mice had increased neuronal loss as compared with WT mice [67]. In MCAo and PT models, which cause large infarction, WT mice exhibit larger infarction than IP₃R2 KO mice [61, 65]. Thus, astrocytic Ca²⁺ signals have different effects on neuronal death and brain damage in mild vs. severe brain injury models.

2.3 Morphology of Reactive Astrocytes After FIS

Since the clinical aim of stroke therapy is to salvage the penumbral cells, understanding the spatial and temporal changes of reactive astrocytes at molecular and cellular levels will provide therapeutic insights for brain repair after stroke. The hallmark of reactive astrogliosis after FIS is the morphological changes and the increased expression levels of GFAP [33, 44, 46, 50, 68, 69]. These changes include the eventual formation of a glial scar, which establishes both a physical and biochemical barrier that separates the ischemic core from the vital tissues. Although there is little expression of GFAP in the astrocytes within the cortex under normal condition [45, 68, 70], cortical astrocytes undergo dramatic increases in GFAP levels over time after FIS. Thus, GFAP expression has been used to study the morphological changes of reactive astrocytes following ischemia. Using endotelin-1 collagenase type IV-S-induced ischemia model, Mesttriner et al. [71] conducted a detailed study on morphology of reactive astrocytes in the penumbra in the chronic stage, i.e., 30 days after ischemic stroke in rats. Their study showed that not only GFAP+ astrocyte density significantly increased after stroke but the ramification and length of primary processes also increased compared with the astrocytes in a sham control group. Wagner et al. studied the morphology of reactive astrocytes in

rats at early stage, i.e., day 4 after middle cerebral artery occlusion (MCAo) and showed that the process volume, diameter, length, and branching levels in reactive astrocytes in the penumbra were increased compared with the astrocytes in contralateral hemisphere and in the distant regions from the ischemic core [72]. Moreover, the process volume and diameter in the penumbra were larger than those in the distant regions. The mean process length of astrocytes in the contralateral hemisphere was longer than reactive astrocytes in the penumbra and the remote region, while the mean process length of reactive astrocytes in the remote region was slightly but significantly longer than those in the penumbra. On the other hand, the mean process branching was similar in the penumbra and in the remote region. The data shows that reactive astrocytes become hypertrophic rather than increase the length of their process at a relatively acute stage of ischemia.

Reactive astrogliosis after ischemia is highly dependent on time and location [45, 46, 73]. Li et al. examined the dynamic changes of reactive astrocytes in the cortex using a photothrombosis-induced mouse ischemia model with a high temporal resolution [68]. GFAP expression was examined in mouse brains after 2, 4, 6, 8, 10, 12, and 14 days following PT using immunostaining and confocal microscopy. A significant increase of GFAP expression was observed 2 days after PT, suggesting astrocytes had been activated. Reactive astrocytes exhibited a stellate morphology and more hypertrophy and expressed higher GFAP up to 4 days after PT. After day 6 post PT, reactive astrocytes became densely packed and exhibited a stream-like structure with their elongated processes pointing towards the ischemic core—a feature of reactive astrocyte polarization and astroglial scar formation, but reactive astrocytes became less hypertrophic after day 6 post PT. Ten days after PT, reactive astrocytes in the penumbra remained similar but with longer processes, indicating the maturation of a glial scar. A significant increase in GFAP expression was also observed in the regions further away from the scar border but with similar morphology to the astrocytes under normal conditions. Thus, the morphology of GFAP+ astrocytes in the penumbra underwent dynamic modifications over time following PT.

Astrocytes exhibit non-overlapping domains in healthy mouse brains [6]. In an epileptic brain, reactive astrocytes interdigitate, i.e., exhibit overlapping domains between the neighboring astrocytes [74]. In an electrically induced mouse lesion model, reactive astrocytes in the cortex exhibit minimal interdigitation, which is similar to those in the cortex of control animals [5]. Thus, it is likely that the severity of brain injury determines the degree of overlapping between reactive astrocytes. It will be interesting to investigate whether reactive astrocytes interdigitate after FIS and whether the degree of interdigitation is temporal and spatial dependent.

2.4 Proliferation of Reactive Astrocytes After FIS

Reactive astrocyte proliferation after FIS has been well documented using PT-induced focal ischemia and MCAo models [45, 46, 73, 75–78]. Glial scar formation is associated not only with morphological changes of reactive astrocytes but

also with substantial tissue shrinking [45]. Spontaneous recovery from brain injury in the chronic phase of ischemia may involve reactive astrogliosis [79]. While it is well known that the common feature of reactive astrocytes is increased GFAP expression levels, whether reactive astrogliosis is merely through the upregulation of GFAP or through cell proliferation is less understood. Bromodeoxyuridine (BrdU) labeling and immunostaining are the general approaches used to assess the spatial and temporal changes of reactive astrocyte proliferation [44–46, 73, 75, 80]. Li et al. [68] studied the dynamic change of proliferation of astrocytes and microglia after FIS. They designed a “time-block” protocol to label proliferating cells at different times after PT. Mice were administered with BrdU at days 1, 3, 4, 5, 9, 11, and 13 post-PT for 2 consecutive days and transcardially perfused after 1 day from the last injection. The proliferation of reactive astrocytes was studied using GFAP and BrdU double staining. BrdU+ cells and GFAP+ astrocytes dramatically increased from day 2 post PT and reached their peak at day 4; however, overall, the GFAP+BrdU+ proliferating astrocytes only accounted for a small percentage of total BrdU+ cells, which reached a peak value of about 6% from days 3 to 4 post PT and then declined sharply. Based on the labeling protocol, the GFAP+BrdU+/BrdU+ ratio at each time point represents the relative rate of the generation of reactive astrocytes. On the other hand, the GFAP+BrdU+/GFAP+ ratio also reached the highest level within days 3–4 post PT. These data demonstrated that FIS increases the population of proliferating reactive astrocytes in a highly temporal dependent manner. The results indicate that glial scar is formed largely from the existing astrocytes through the upregulation of GFAP, rather than from newly generated astrocytes through proliferation. Although proliferation rate reduces dramatically after 8 days following PT, the morphology of reactive astrocytes maintains straight processes pointing to ischemic core for a prolonged time based on the GFAP expression [68, 73, 75]. This phenomenon suggests that the expression of certain genes is irreversible in the penumbra during recovery from focal ischemic injury, and glial scar formation causes substantial tissue remodeling and permanent and persistent structural changes. Although the approach might underestimate the total number of proliferating cells since the injection protocol may not label all the proliferating cells [81], the GFAP+BrdU+/BrdU+ ratio should represent the ratio of newly generated reactive astrocytes out of the total proliferating cells.

2.5 Signaling Pathways of Reactive Astrogliosis After Ischemia

Advancements in genetics and molecular biology have helped to extensively characterize reactive astrocytes after brain injury. A plethora of molecular markers and signaling pathways associated with reactive astrogliosis have been identified [80, 82–85]. The following is a brief summary of the major signaling pathways that have been identified to regulate reactive astrogliosis in ischemic stroke.

Intermediate filaments. The intermediate filament GFAP is expressed both in protoplasmic and fibrous astrocytes. Following ischemic stroke, the peri-lesional

astrocytes become reactivated and dramatically upregulate their GFAP in a temporally and spatially dependent manner. Genetic deletion of GFAP is neuroprotective against metabolic insult and excitotoxicity which is attributed to increased production of the glial-derived neurotrophic factor (GDNF) by GFAP knockout astrocytes [86]. GFAP^{-/-} mice have exhibited a more significant decrease in cortical cerebral blood flow, and a relatively larger infarct volume than WT mice after MCAo, indicating that GFAP-null mice have a high susceptibility to cerebral ischemia, which indicates that astrocytes and GFAP play an important role in the progression of brain damage after ischemia [87]. However, studies using GFAP^{-/-}/Vimentin^{-/-} double knockout mice show that in acute ischemia, double deletion of GFAP and vimentin results in decreased glutamate uptake, increased susceptibility to oxidative stress, and significantly increased infarct volumes (two to three times compared with that of WT mice). Hence, the deletion of GFAP and vimentin in the acute stage disrupts glial scar formation and promotes neuronal regeneration after ischemia [88]. Whereas in chronic stages of stroke, attenuation of astroglial reactivity by GFAP and vimentin deletion impaired axonal remodeling and functional recovery [89]. Together, these studies suggest that reactive astrocytes have a protective role in brain ischemia. GFAP and vimentin are essential to retaining the beneficial function of reactive astrocytes in ischemic as well as recovering nervous tissues. Thus, manipulation of astrocytic reactivity may represent a therapeutic target for neurorestorative strategies in ischemic stroke.

p38 MAPK pathway. Mitogen-activated protein kinases (MAPK) are a family of enzymes transducing a wide range of extracellular signals such as inflammation, growth factors, and toxic stimuli as well as integrating corresponding cellular responses [90]. Among p38, MAPKs is of particular importance since it transduces cellular inflammation [91]. p38 MAPK can become activated in neurons, microglia, and astrocytes after ischemic stroke [92–94]. A recent study using an astrocyte-specific p38 MAPK knockout mouse model demonstrated that delayed activation of p38 MAPK following ischemic injury corresponds with upregulation of GFAP in the penumbra region [95], suggesting an important role of p38 MAPK in signaling reactive astrogliosis following ischemic stroke. Furthermore, conditional deletion of using GFAP-Cre/LoxP p38 MAPK mice resulted in a significant reduction of reactive astrogliosis in the penumbra following ischemia compared to WT animals, but had no significant effect on motor functional recovery after ischemia. p38 MAPK signaling may be a critical transducing pathway to modulate reactive astrogliosis in ischemic stroke.

Notch signaling pathway. Notch signaling is a highly conserved pathway critical for the maintenance and self-renewal of progenitor cells and to inhibit precocious neurogenesis [96]. In ischemic stroke, Notch 1 signaling was activated in astrocytes in the peri-infarct region by 24 h [97], and conditional deletion of Notch 1 from GFAP positive cells resulted in a decreased number of proliferating astrocytes and an increased number of invading CD45+ immune cells following ischemic injury [98]. Further study showed that Notch1–STAT3–ETBR axis connects a signaling network that promotes reactive astrocyte proliferation after a stroke [99]. Genetic fate mapping analysis revealed that a subpopulation of reactive astrocyte activated

Notch1 signaling in response to injury and became proliferative [97, 100]. A recent study also showed that Notch1 signaling is important in reactive astrocyte differentiation to neurons in an ischemic brain [101]. These studies suggest that Notch 1 plays different roles including neurogenesis, reactive astrocyte proliferation, and transdifferentiation after brain injury. Some outstanding questions which will be mandatory in future studies are: Why does only a subset of reactive astrocytes express Notch 1 signaling and become proliferative? What is the spatiotemporal pattern of Notch 1 signaling after ischemia? Does inhibiting Notch 1-induced astrocyte proliferation affect functional recovery? Answering these questions will better define the role of Notch 1 in reactive astrogliosis and provide novel therapeutic insights for stroke therapy.

STAT3 signaling. Signal transducer and activator of transcription 3 (STAT3) is a member of the Jak-STAT family and activated by hormones, growth factors, or cytokines [102]. The role of STAT3 in reactive astrogliosis has been extensively studied in spinal cord injury models [81, 103–105]. In ischemic stroke, reperfusion-induced increases in ROS and inflammatory cytokines are known to be powerful stimulants of STAT3 signaling [106, 107], but it remains a matter of extensive debate regarding cell type specific activation of STAT3 in the post ischemic brain. Some reports suggest neurons are the predominant source of activated STAT3 [108, 109], while others have shown that STAT3 is also activated in astrocytes along with neurons [110, 111]. More recently, it suggests that STAT3 can regulate reactive astrogliosis induced by neurotoxic insults where activation of STAT3 in astrocytes and subsequent increases in GFAP expression can be induced by direct neuronal death even in the absence of astrocyte damage [112, 113]. It appears that its role in driving reactive astrogliosis in ischemic stroke remains to be explored.

Transforming growth factor-Beta (TGF- β) signaling. TGF- β is an injury-related peptide and has been shown to increase immunoreactivity in both the infarct and penumbra in stroke patients [114] as well as animal models of FIS [115–117]. TGF- β has been mostly regarded as neuroprotective particularly against the *N*-methyl-D-aspartate receptor (NMDA)-induced excitotoxicity [116, 118]. It is suggested that the protective action of TGF- β is mostly mediated by astrocytes [119, 120]. Most recent data suggests that disrupting astrocytic FGF receptors results in reduced scar size, increased astrocyte activation, and inflammation, but did not affect infarct volume [121, 122]. These studies suggest that TGF- β signaling in astrocytes can limit neuroinflammation, reactive astrogliosis in the peri-infarct cortex and preserve brain function during the subacute period after stroke.

Despite the extensive characterization of reactive gliosis, high heterogeneity and the context-dependent nature of reactive astrogliosis in the injured brain (e.g., acute phase vs. chronic phase, white matter vs. gray matter, penumbral region vs. distant region) complicate our understanding of signaling pathways for regulating this process [3, 85, 123]. Recent studies on the genome-wide transcriptome profiles of pure glial cells using microarray and RNA-Seq technologies provide an unprecedented opportunity to study glia phenotypes and functions in health and disease [8, 80, 124, 125]. Data from these studies open a window for a comprehensive view of complex mechanisms by which astrocytes contribute to brain protection and repair in ischemia.

Especially when using RNA-Seq, a large number of signaling pathways that trigger reactive astrogliosis and molecular changes of astrocytes can be identified. Changes in transcriptome profiles at different times after ischemic stroke will help to identify novel molecular pathways by which astrocytes respond to ischemic insults at different stages [80]. These data will shed new light on the roles of astrocytes in brain protection and repair and provide information for astrocyte-targeting therapeutic strategies for ischemic therapy.

2.6 Stem-Cell-like Properties of Reactive Astrocytes and Endogenous Neuronal Differentiation of Reactive Astrocytes After Ischemic Stroke

Growing evidence has shown that ischemic stroke dramatically increases neurogenesis in the subventricular zone (SVZ) and subgranular layers in dentate gyrus [126]. Endogenous progenitor cells proliferate and differentiate into neurons in response to ischemic insult; the newly generated neurons migrate to the damaged region thereby integrating into the neural network [127–129]. These studies suggest that endogenous neurogenesis contributes to brain repair and spontaneous recovery after ischemia. As we already discussed, normally quiescent astrocytes become reactive and resume proliferation after ischemia. This implies that reactive astrocytes might be in an immature state due to a certain degree of dedifferentiation induced by ischemia. Studies have demonstrated that proliferating astrocytes in specific brain regions act as adult neural stem cells under normal conditions [130, 131]. In addition to many changes in gene profile, reactive astrocytes also exhibit stem cell-like properties [132–136]. These studies suggest that reactive astrocytes may attempt to reconstitute neurons after ischemia. Indeed, they express neural stem-cell-related proteins such as nestin and Sox2 [136], doublecortin (DCX), an immature neural stem cell marker [101, 137], and oligo 2, a transcription factor that regulates neuroglia fate decision [138]. It shows that a sonic hedgehog (SHH) signal can directly act on the astrocytes and is necessary and sufficient to elicit the stem cell response both in vitro and in vivo [133]. Stroke also elicits a latent neurogenic program in striatal astrocytes in a mouse model [101, 139]. It is reported that Notch 1 signaling is reduced in reactive astrocytes after stroke and attenuated Notch 1 signaling is necessary for neurogenesis by striatal astrocytes [101]; furthermore, blocking Notch 1 signaling triggers astrocytes in the striatum and the medial cortex to enter a neurogenic program even in the absence of a stroke, indicating that attenuated Notch 1 signaling is necessary for neurogenesis by striatal astrocytes. After a prolonged time (e.g., 13–16 weeks) following ischemia, reactive astrocyte-derived neurons in striatum not only express mature neuronal markers but also are able to fire the action potential and exhibit synaptic activity, suggesting the integration into their local neural network [139]. These studies provide a molecular base for how reactive astrocytes acquire stem cell lineage and become neurons after a stroke.

Reactive astrocytes also affect the migration of neural progenitor cells after ischemia. Young et al. [140] reported for the first time that ischemic stroke causes substantial reactive astrogliosis in SVZ and the hypertrophic reactive astrocytes and their tortuous processes disrupt the neuroblast migratory scaffold and cause SVZ reorganization after a stroke [140]; thus, reactive astrocytes in SVZ might also play a modulating role in neurogenesis thereby affecting stroke recovery.

2.7 Direct Astrocyte-to-Neuron Conversion After Ischemic Stroke

The fact that reactive astrocytes express neural stem markers and can be transformed into mature neurons after a prolonged time following ischemia indicate that reactive astrocytes have the intrinsic potential to generate neurons and reactive astrocytes-to-neuron conversion can be facilitated under permissive conditions such as overexpression of neurogenic factors. It has been recently reported that astrocytes indeed can be converted into neuroblasts and neurons in vitro and in vivo by forced expression of a single transcriptional factor such as Sox2 [141, 142], Neurog-2 [143–145], NeuroD1 [146], Ascl1 [147], or a combination of multiple transcriptional factors such as Ascl1, Lmx1B, Nurr1, Oct4, Sox2, or Nanog [148, 149]. Astrocyte-converted neurons express mature neuronal markers such as Map2 and NeuN [141, 146]. Electrophysiological recording showed that cultured mouse astrocyte-converted neurons by NeuroD1 are glutamatergic; furthermore, astrocyte-converted neurons by NeuroD1 in mouse brain can exhibit synaptic currents and integrate into neural circuit in vivo [146]. In vitro direct reprogramming study of postnatal astrocytes showed that astrocyte-to-neuron conversion is swift, but different proneural factors can elicit distinct transcriptional programs. Both Neurog-2 and Ascl1 rapidly elicited neurogenic programs to induce glutamatergic and GABAergic neuronal conversion but followed distinct paths with a few common genes in the neurogenic cascades [145]. The study of shared target genes will allow identifying a particularly important subset of downstream targets capable of directly converting astrocytes into functional neurons.

Although, reactive astrocyte-to-neuron conversion by forced expression of transcription factors has not been reported in the context of strokes, these studies suggest that recruiting reactive astrocytes and directly converting them into neurons might be a promising strategy for brain repair after ischemic stroke. Since reactive astrocytes exhibit different proliferating rates at different times after FIS [68], there might be an optimal timing to convert them into neurons using genetic manipulation techniques such as astrocyte-specific inducible transgenic mice and viral transduction. Because stroke causes a large damage in the cortex, it is also interesting to study whether reactive astrocytes in the cortex have the similar capacity of neuronal transformation to striatal astrocytes. Future study is also needed to explore whether any small molecular drugs targeting the signaling pathway of endogenous neuronal

differentiation will facilitate astrocyte-to-neuron conversion [150, 151] and to determine whether astrocyte-derived neurons contribute to brain recovery and improvement of stroke outcomes.

3 Concluding Remarks

Despite tremendous investments in translational research, there are still limited treatment options for stroke therapy. Tissue plasminogen activator (tPA) is the only available FDA-approved drug for acute stroke treatment, but it is effective within a narrow therapeutic window of a few hours after the onset of a stroke. Treatment of a stroke is primarily dependent on supportive care, secondary prevention, and rehabilitation after the acute phase. tPA treatment is also associated with intracranial bleeding. Although various mechanisms showing how ischemia leads to neuronal death and brain damage have been identified using animal models, current neuron-centric strategies have not resulted in major breakthroughs in stroke therapy; therefore, alternative strategies to target astrocytes could be an important direction for stroke therapy. Since astrocytes have intimate contact with neurons and vasculature and undergo dynamic changes in Ca^{2+} signaling, morphology, proliferation, and gene expression after ischemia, targeting reactive astrocytes is a promising strategy for brain damage and repair after stroke. Knowledge regarding the dynamic changes of function and signaling pathways that trigger reactive astrogliosis and facilitate neuronal differentiation should provide information for effective strategies for ischemic brain therapy. Studies on genome-wide transcriptome profiles further provide information for astrocyte-targeted therapeutic strategies for brain protection and repair. However, as astrocytes exhibit heterogeneity in different regions, a comprehensive understanding of reactive astrogliosis is extremely vital to the designing strategies to modulate reactive astrogliosis for brain repair. On the other hand, using the stem-cell-like properties of reactive astrocytes, it is feasible to directly convert them into neurons by expressing transcriptional factors. Thus, targeting astrocytes for stroke therapy is an emerging and promising area.

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Microglia: A Double-Sided Sword in Stroke

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Abbreviations

CD200R	CD200 receptor
CNS	Central nervous system
CX3CR1	CX3C chemokine receptor 1

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DCs	Dendritic cells
Gals	Galectins
HMGB1	High-mobility group box 1
HSP	Heat shock protein
IFN- γ	Interferon-gamma
IgSF	Immunoglobulin superfamily
iNOS	Inducible nitric oxide synthase
KO	Knockout
NO	Nitric oxide
Nox	NADPH oxidase
NSC	Neural stem cells
OPC	Oligodendrocyte progenitor cell
pro-MMP-9	pro-matrix metalloproteinase-9
RAGE	Receptor for advanced glycation endproducts
ROS	Reactive oxygen species
SGZ	Subgranular zone
SVZ	Subventricular zone
TGF- β	Transforming growth factor- β
TIMP-1	Tissue inhibitor of metalloproteinases-1
TNF- α	Tumor necrosis factor- α
TREM	Triggering receptors expressed on myeloid cells
VEGF	Vascular endothelial growth factor

1 Introduction

Microglia were first described by Pio del Rio Hortega as early as in 1919 [1]. Although the precise origin of microglia is still in debate up to date, it is commonly accepted that the microglial progenitors are derived from the yolk sac. They migrate into the central nervous system (CNS) during early embryogenesis and give rise to microglia throughout the brain parenchyma [2, 3].

Microglia have multiple vital functions in the CNS. Under physiological state, microglia display a ramified phenotype, characterized by a small cell body and numerous branched processes. These so-called “resting microglia” actively survey the surrounding environment and respond promptly to even subtle changes in the CNS. When a microvascular or an isolated neuron is impaired, the surveying microglia may offer structural and trophic support to them so that the normal functions could be preserved. Microglia also remove the sick individual neurons or other CNS cells. These daily functions of microglia are called housekeeping activity, which contributes to the maintenance of CNS homeostasis [1, 4–6]. Upon the noxious cues such as brain infection or injury, microglia expand their cell bodies and retract their processes, becoming reactive microglia. These reactive microglia are the first line of defense in the compromised brain. Due to the importance of microglia

in brain homeostasis and brain injuries, their activities are well-regulated by a large group of surface receptors [7].

Activation of microglia occurs within 1 h after stroke, much earlier than the infiltration of other immune cells from the periphery [8]. Interestingly, the activated microglia appear to be double-edged swords in the battle of brain injuries. The beneficial effects of microglia include cell debris clearance and trophic factor production [4, 9–11]. In contrast, overactivated microglia can expand tissue damage and hinder CNS repair by releasing large amount of inflammatory mediators or free radicals [11–13]. These seemingly contradictory functions of microglia might be at least partially due to their diverse phenotypes in response to different microenvironmental cues. Actually, microglia exhibit a spectrum of different but overlapping functional phenotypes upon activation. In a conceptual framework, M1 (classically activated) and M2 (alternatively activated) dichotomy have been used to represent two extreme activation states of microglia. Although being oversimplified, the M1 and M2 classification is a useful concept to understand microglia functional status in the progress of injury/repair and to explore therapeutic strategies targeting microglia responses (Fig. 1) [14].

2 Surface Receptors That Modulate Microglial Responses in the Ischemic Brain

Microglia make contact with other cells via recognizing the signals released after CNS insults. Two types of signals, the so-called “Off” and “On” signals are identified by different microglial surface receptors (Table 1). The “Off” signals occur constantly under normal physiological conditions, maintaining microglia at the quiescent surveillant state. In contrast, the “On” signals are released under various pathological conditions, leading to microglial activation and mobilization. Several receptors have been identified to recognize the “On” or “Off” signals released in the ischemic brain and play an important role in stroke pathology. In this section, we will discuss some of these microglial receptors, including the “Off” receptors TREM2, CD200, CX3CR1, as well as the “On” receptors RAGE and galectins.

2.1 Triggering Receptors Expressed on Myeloid Cells

Triggering receptors expressed on myeloid cells (TREM) belong to the immunoglobulin superfamily (IgSF), which plays an important role in regulating immunity responses and medicating cell surface recognition. It is expressed in a variety of immune cells, including microglia, dendritic cells (DCs), monocytes, and neutrophils [15]. Among these cells, TREM2 is the most relevant to microglia due to its constitutive expression.

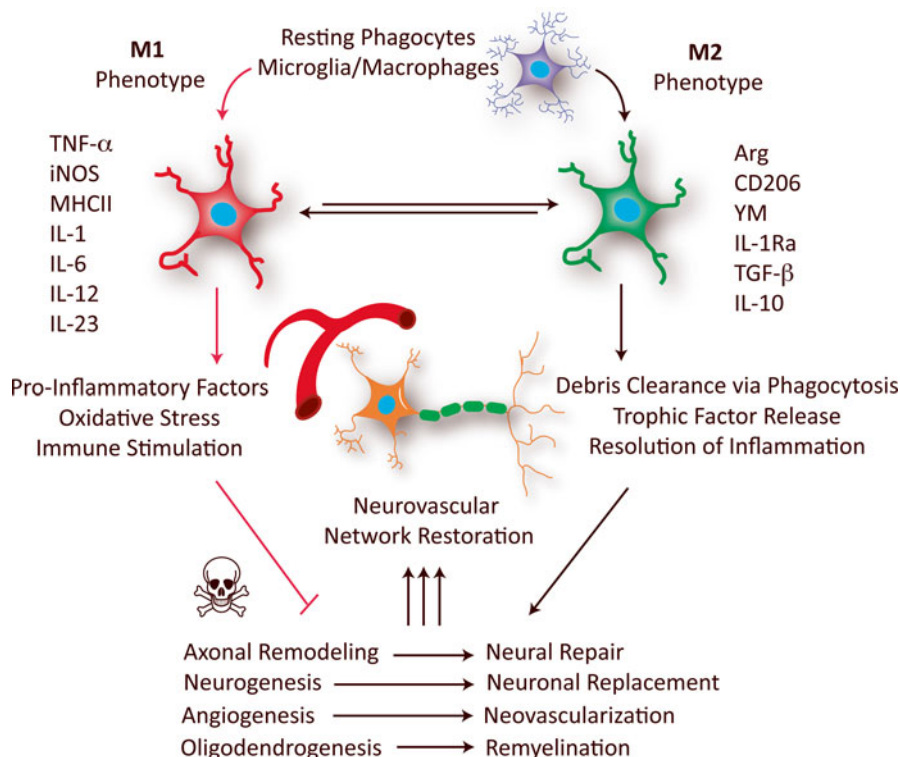


Fig. 1 Microglia response to different stimuli with either M1 or M2 phenotype. M1 microglia are characterized by production and secretion of proinflammation factors such as $TNF\alpha$, iNOS, MHCII, IL-1, IL-6, IL-2, IL-23, etc., leading to exacerbated tissue damage. In contrast, M2 microglia polarization are characterized with the expression of Arg, CD206, YM, IL-1Ra, IL-10, and TGF- β , etc. M2 microglia improve brain repair/regeneration by enhancing phagocytosis, releasing trophic factors, and resolving cerebral inflammation. Arg: arginase; IL: interleukin; IL-1Ra: IL-1 receptor antagonist; iNOS: inducible nitric oxide synthase; MHCII: major histocompatibility complex II; TGF- β , transforming growth factor- β ; $TNF\alpha$: tumor necrosis factor- α . (Adapted from Hu et al., *Nat Rev Neurol*. 2015; 11(1):56–64.)

TREM2 migrates to the cell surface from the intracellular pool upon stimulation by ionomycin or interferon-gamma (IFN- γ) [16, 17], promoting microglial anti-inflammatory responses and decreasing the production of $TNF-\alpha$ or IL-6 [18, 19]. Loss of a single copy of TREM2 alters the morphological phenotype of microglia and induces neurotoxicity [20]. Studies also reveal increased microglial expression of TREM2 after ischemic stroke. In *in vitro* ischemic model, TREM2 deficiency in microglia ameliorates microglial activation and inhibits the phagocytosis of injured neurons. In a mouse model of stroke, TREM2 knockout (KO) results in reduced phagocytosis and impaired infarct brain tissue resorption. As a result, TREM2 KO mice exhibited worsened neurological recovery and increased brain tissue loss in

Table 1 Microglial surface receptors in stroke

Function	Receptor type	Receptor name	References
“Off”	Immunoglobulin superfamily	TREM2	[21, 22, 105]
	Chemokine receptors	CX3CR1	[34, 106–109]
	Immunoglobulin superfamily	CD200R	[26, 110, 111]
“On”	Immunoglobulin superfamily	RAGE	[44, 112–114]
	Chemokine receptors	CCR2	[115–119]
		CCR5	[120–123]
	Purinergic receptors	A2A receptor	[124–126]
		A3 receptor	[127, 128]
		P2X4	[129, 130]
		P2X7	[131–133]
	Carbohydrate-binding receptor	Galectins	[48, 51]
		Selectins	[134–136]
	Pattern recognition receptors	TLRs	[137]
	Scavenger receptor	CD36	[138–140]
	Phosphatidylserine receptors		[141–143]

the ipsilateral hemisphere [21, 22]. In contrast, TREM2 overexpression promotes microglial migration, lipid catabolism, and the phagocytosis of neuronal cell debris in stroke as well as in Alzheimer’s disease (AD) [17, 23].

The ligands of TREM2 have not yet been identified, which prevents the understanding of microglial activation through this receptor. Multiple types of cultured neuronal cells have been shown to bind to TREM2-Fc fusion protein but not TREM1-Fc, suggesting that TREM2 ligand(s) are constitutively expressed on healthy neurons [24]. The identity of TREM2 ligand(s), however, remains elusive. Interestingly enough, a recent study showed that TREM2-Fc fusion protein pulled down nucleic acids from ischemic brain lysate, suggesting that nucleic acids might be a potential ligand for TREM2 in brain ischemia [21]. Another study has shown that heat shock protein (HSP) 60 expresses on neuron and astrocytes surface and might be a TREM2-binding protein [25]. Accordingly, treatment with HSP60 can stimulate microglial phagocytosis effectively in a TREM2-dependent manner.

2.2 CD200 Receptor

CD200 receptor (CD200R) is another member of the immunoglobulin receptor family. CD200R expression is restricted to cells of the myeloid lineage, including CNS microglia [26]. CD200, the sole ligand to CD200R, is expressed by a variety of cells including neurons. The CD200-CD200R signaling plays a critical role in keeping microglia at the physiological “resting” state. Microglia in CD200-deficient

mice display spontaneous activation, characterized by morphological changes and upregulation of CD11b and CD45 [27]. In experimental autoimmune encephalomyelitis (EAE) and experimental Parkinson's disease (PD), disruption of the CD200-CD200R binding results in enhanced microglial activation and worsens the outcome [28]. Shortly after ischemic injury, CD200 mRNA is decreased, correlating with enhanced microglial proinflammatory activation [29]. These evidence strongly support the role of CD200R signaling in the maintenance of immune homeostasis in the CNS.

2.3 CX3CR1

CX3CR1 is a G-protein-coupled chemokine receptor expressed mainly in microglia in the CNS. Its unique ligand CX3CL1 is highly expressed in neurons, CX3CR1 has been shown to play an important role in neuron–microglia crosstalk and the maintenance of CNS homeostasis. CX3CR1 participates in microglia-mediated synaptic pruning [30]. In addition, CX3CR1 regulates a variety of microglial functions, including migration, proliferation, adhesion, phagocytosis, and inflammatory cytokine production [31, 32]. The role of CX3CR1 signaling in CNS disorders is controversial. On the one hand, CX3CR1-deficient microglia produce less TNF- α , NO and superoxide in AD [33]. In models of ischemic brain injuries, inhibition of CX3CR1 signaling has been shown to be neuroprotective by reducing activation of resident microglia and inhibiting recruitment of peripheral macrophages [34, 35]. On the other hand, some studies found that microglia without CX3CR1 are neurotoxic due to the loss of inhibitory signaling [36]. The role of the CX3CR1-CX3CL1 signaling may be context-dependent under different pathological conditions and in different stages of disease progression.

2.4 Receptor for Advanced Glycation Endproducts

Receptor for advanced glycation endproducts (RAGE) belongs to the immunoglobulin superfamily and is highly expressed in immune cells [37]. RAGE exaggerates tissue injury under various pathological conditions by potentiating host immune responses [38]. Microglial RAGE recognizes ligands such as high-mobility group box 1 (HMGB1), S100 [39, 40], A β , and AGE [41, 42]. The role of microglial RAGE in brain injuries has been reported. The expression of RAGE in macrophages/microglia increased in active lesions while its ligand, S100 β expression was strikingly upregulated in reactive astrocytes in a demyelination model [40]. Another study showed that HMGB1 is released from injured neurons after ischemia and contributes to tissue injury through its interaction with RAGE [43]. HMGB1-RAGE signaling results in functional exhaustion of lymphocytes and mature monocytes

after extensive ischemia, which is a hallmark of immune suppression [43, 44]. These studies suggest that HMGB1-RAGE signaling is critical for microglia/macrophage activation upon neuronal necrosis.

2.5 Galectins

Galectins (Gals) are a family of β -galactoside-binding lectins. They bind to cell surface to form multivalent complexes and induce intracellular signals regulating cell survival and differentiation [45, 46]. Among all the Gals, Galectin-3 (Gal-3) has been shown to be important in stroke pathology. Gal-3 has been shown to be expressed on microglia and acts as an endogenous ligand of TLR4. Depletion of Gal-3 induced an anti-inflammatory and neuroprotective effects after global brain ischemia, suggesting a detrimental role of this receptor in ischemic stroke [47]. However, other studies argued that Gal-3 expression might have beneficial effects in the ischemic brain. Gal-3 gene deletion prevented VEGF upregulation after MCAO, which inhibited increases in endothelial proliferation and angiogenesis in the striatum [48, 49]. Another study proved that Gal-3 mediated angiogenesis and microglia migration [50]. In addition, Gal-3 modulates microglial response by upregulating phagocytic receptor TREM-2b during cuprizone-induced demyelination which can help the myelination process by enhancing oligodendrocyte differentiation [51, 52].

Recent research have identified more and more surface receptors on microglia mediating microglial recognition of “On” or “Off” signals in the ischemic brain. Integrated actions of these receptors result in fine-tuned microglial functions including phagocytosis, cell migration, inflammatory mediation and trophic factor release. These tightly regulated microglia play diverse and dynamic functions in the progress of stroke and in the process of brain repair.

3 Dual Roles of Microglia in the Ischemic Brain

Detrimental effects of activated microglia in the ischemic brain. It is well known that excessive microglial activation may lead to expansion of ischemic brain damage and deterioration of neurological outcomes [53, 54]. The toxicity of microglia, especially the M1 microglia, is mediated by the release of a variety of harmful substances, including nitric oxide (NO), reactive oxygen species (ROS), and proinflammatory cytokines [55]. In addition, overactivated microglia impair post-stroke neurogenesis [12, 13], prevent axon regeneration [56], and limit the efficacy of thrombolytic therapy [57].

Proinflammatory cytokines/chemokine-mediated neurotoxicity. Acute brain injuries lead to rapid damage to CNS cells including endothelial cells, astrocytes

and neurons. Those damaged cells release a set of activation signals such as ATP, glutamate, HMBG1 et al. In response to these alarming signals, microglia may release large amount of proinflammatory cytokines including IL-1, IL-6, IL-12, and TNF- α . The accumulation of these cytokines may exacerbate brain damage. TNF- α is a typical and potent proinflammatory cytokine released from activated microglia. It plays a crucial role in cerebral ischemia and brain inflammation. TNF- α deficiency is associated with reduced microglial population size and Toll-like receptor 2 expression, which can influence the neuronal response to injury [58]. TNF- α has also been demonstrated to induce cell death via caspase-8-mediated apoptosis and RIPK1-mediated necroptosis [59, 60]. Moreover, microglial TNF- α has been reported to block both neurogenesis and oligodendrogenesis [61].

In addition to the secretion of proinflammation cytokines, microglia also release chemokines to recruit peripheral leukocytes. For example, microglia secretes CCL2 when triggered by the proinflammatory stimuli such as IFN- γ , TNF- α , and IL-1 β [62]. CCL2 is a potent chemokine for peripheral monocyte recruitment [63]. Once reaching the injured site, monocytes further differentiated into activated macrophages, and contribute to tissue damage. CCL2 or CCR2 knockout mice exhibited reduced number of phagocytic monocytes and smaller infarcts after middle cerebral artery occlusion. Therefore, microglia-released CCL2 may contribute to the recruitment of monocytes and the exacerbation of tissue damage [64].

Free radical-mediated neurotoxicity. Oxidative stress is important in ischemic/reperfusion injury following stroke [65]. Activated microglia, together with intracellular organelles of injured cells and infiltrating immune cells, are main sources of ROS in the ischemic brain [66, 67]. NADPH oxidase (Nox) is the major enzyme that catalyzes the production of ROS. Nox generates the highly reactive free radical, superoxide, via its Nox2 catalytic subunit [68]. The innate immune cells including microglia have been reported to express this enzyme. Apocynin, a known antioxidant and inhibitor of Nox2 NADPH, significantly reduced infarct volume when applied 1 h prior to stroke or 24 or 48 h after stroke [69].

Activated microglia also upregulate inducible nitric oxide synthase (iNOS) and produce NO. ROS can react with NO to form peroxynitrite, a more toxic product that cause cell damage. Iadecola C et al. have demonstrated that inhibiting iNOS was able to ameliorate brain damage after stroke [70, 71], suggesting the importance of NO and derivatives in the ischemic brain injury.

Microglial inhibition of axonal regeneration. It is noted that the loss of neurological function after ischemic brain injury might be attributed to the interruption of axonal connectivity rather than the neuronal loss. Activated microglia/macrophages have been shown to hinder axonal regeneration by physical cell–cell interactions with neurons [72]. M1 Microglia activated by LPS inhibited neurite outgrowth and induced growth cone collapse of cortical neurons in vitro, a pattern that was only observed when there was direct contact between microglia and neurons [73]. These studies suggest that microglia/macrophages play a role in the failure of axonal regeneration in the CNS.

3.1 *Beneficial Effects of Activated Microglia*

Resolution of local inflammation While the M1 microglia promote local inflammation, the M2 microglia have been proposed to play an anti-inflammatory role. In the normal CNS, microglia have been documented to produce anti-inflammation cytokines such as IL-4 and TGF- β 1 [74]. Zhou et al. also demonstrated that IL-4-induced M2 microglia increase the expression and secretion of TGF- β 2 [75], which may subsequently involve in the resolution of local inflammation. In addition to direct secretion of anti-inflammation cytokines, microglial clearance of cell debris through phagocytosis also helps to mitigate successive inflammation. Phagocytosis of apoptotic cells are mediated by the interaction of engulfment receptors on microglia and the “eat me” signal from apoptotic cells [76–78].

Activated microglia in oligodendrogenesis. Oligodendrocytes are myelinating glial cells that ensheath axons, ensuring fast and saltatory impulse propagation. Oligodendrocytes are differentiated from the oligodendrocyte progenitor cells (OPCs), which disperse throughout the adult brain. OPCs continue to proliferate in healthy adult brain and generate oligodendrocytes. Recent study demonstrated that those adult-born OLs could remodel existing myelin and play a role in myelin homeostasis [79]. After stroke, OPCs are activated and migrate toward the infarct core, accounting for the recovery of oligodendrocytes and remyelination around the peri-infarct core [80].

Accumulating studies have highlighted the importance of microglia-released trophic factors in the process of remyelination after brain injury. IGF-1 secreted from activated microglia has been reported to induce hippocampus-derived neural progenitor cells to differentiate into oligodendrocytes in adult rats, and protects immature oligodendrocytes from glutamate-mediated apoptosis [81, 82]. Moreover, IL-1 β -deficiency mice devoid of microglial IGF-1 displayed impaired OPC differentiation into mature oligodendrocytes, leading to the remyelination failure [83]. Lalive and colleagues demonstrated that HGF released from microglia is both a chemotactic and differentiation factor for OPCs in injured mature CNS [84]. In addition, microglial CNTF has also been reported to provide a protective role in the remyelination phase [85, 86]. A recent study with specific microglial phenotype depletion further confirmed that M2 microglia promote, while M1 microglia impair OPC differentiation and remyelination.

Microglia in neurogenesis. Microglia play an important role in neurogenesis during brain development. Microglial cells display special amoeboid morphology in early developing brain. These amoeboid microglia possess high phagocytic capacity and actively clear the apoptotic neurons to ensure correct cytoarchitecture [87]. In addition, microglia actively secrete pro-apoptotic factors to induce neuron death during development. For example, NGF secreted from microglia has been reported to promote the death of neurotrophin receptor p75-positive neurons in retina [88], and microglial TNF- α -induced cell death was demonstrated to control the number of hippocampal progenitor cells [89]. Moreover, microglia release high levels of superoxide ions to promote neuronal death in the cerebellum [90].

Ischemic brain injury triggers neurogenesis in the subgranular zone (SGZ) in the dentate gyrus of the hippocampus and the subventricular zone (SVZ) lining the lateral ventricles [91]. Microglia has been demonstrated to actively promote post-stroke neurogenesis. In rat, treatment of minocycline immediately after transient cerebral ischemia suppressed microglial activation, leading to significantly decreased neurogenesis [92]. In vitro study by Walton et al. confirmed that coculture with microglia or their conditioned medium greatly enhanced neurogenesis of neural stem cells (NSC) [93].

Microglia secrete an array of factors to promote neurogenesis. Liao et al. reported that microglial cells can interact with tenascin-R and secrete NGF, BDNF, and TGF β to promote NSC proliferation and differentiation into neurons [94]. Battista et al. have also reported that activated microglia secrete TGF β and promote neurogenesis in the dentate gyrus of the hippocampus in adrenalectomized animals [95]. Moreover, microglial IGF-1 is known to suppress apoptosis and increase proliferation and differentiation of neural stem cells [9]. Taken together, those studies suggested an active role of microglia in the recovery course after stroke by enhancing neurogenesis.

Microglia in angiogenesis. In the developing CNS, microglia progenitors derived from the yolk sac invade brain parenchyma before vascularization takes place, and has been proposed to promote the formation of brain blood vessels [96]. An evidence for the function of microglia in angiogenesis comes from the PU.1-deficient mice. PU.1 is a transcription factor that is indispensable for microglia development. Mice with PU.1 deficiency display impaired microglial development, which is accompanied by reduced vascular branching and complexity [97]. Kubota et al. showed that CSF1 op/op mice with inactive CSF1 displayed reduced retinal microglia and a significant decrease in branching of the primary vascular plexus in developing retina [98]. Similarly, depletion of retinal microglia by intravitreal clodronate liposome application led to reduced number of retinal microglia associated with decreased retinal vascular density [99]. Those studies strongly suggested an angiogenic role of microglia in the developing brain. The function of microglia in angiogenesis in the ischemic brain, however, has not been well-addressed.

To date, the molecular mechanisms underlying microglial promotion of angiogenesis remain elusive. Rymo et al. showed that coculture of microglia with aortic ring stimulated vessel sprouting, and this effect was dependent on soluble factors released from microglia rather than direct cell–cell interaction [100]. Recently, Li et al. demonstrated that microglial TNF- α upregulate the production of ephrin-A3 and ephrin-A4 in brain microvascular endothelium cells, resulting in the formation of capillary-like tube in vitro [101]. M2 microglia/macrophages produce robust pro-angiogenic factors such as vascular endothelial growth factor (VEGF) [102] and IL-8 [103]. M2 macrophages also release pro-matrix metalloproteinase-9 (pro-MMP-9) in a unique, tissue inhibitor of metalloproteinases-1 (TIMP-1)-free form to enhance their angiogenic capacity [104]. Those studies suggested that microglia may promote brain angiogenesis through an array of soluble factors. Further studies to identify those factors are warranted.

4 Conclusions

Microglia play diverse roles in the injured brain. Our current understanding of microglial functions in the ischemic brain is far from complete. Multiple lines of evidence suggest dynamic changes in microglia after stroke, with a trend of phenotype switch from protective M2-like polarity to toxic M1-like polarity [12, 55]. Such transition not only explains the dual-faced nature of microglia, but also provides targets for immunomodulatory therapies after stroke. Further identification of the microglial receptors and/or signaling pathways that are in charge of phenotype switch is essential for the phenotypic research in the stroke field.

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Crosstalk Between Cerebral Endothelium and Oligodendrocyte After Stroke

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1 Introduction

The concept of the neurovascular unit is now well-accepted in the understanding of brain physiology and pathophysiology. The “neurovascular unit” provides a conceptual framework that emphasizes cell–cell interactions between neuronal, glial, and vascular elements [1–7]. This concept primarily guides research in neuron-related cell–cell interaction mechanisms in gray matter. But cell–cell signaling between non-neuronal cells is also critical for white matter and gray matter function. As shown in the diagram in Fig. 1, cells in cerebral white matter are closely related to each other. For example, oligodendrocytes enwrap axons for efficient conduction of electrical impulses, and the oligodendrocyte–axon interaction is one of the most well-documented aspects of neurovascular unit in white matter. In fact, oligodendrocyte-derived trophic factors such as insulin-like growth factor-1 and glial cell line-derived neurotrophic factor increase axonal length and support neuron survival [9]. Another example of the importance of cell–cell interactions in white matter may be found in the roles of oligodendrocyte precursor cells (OPCs).

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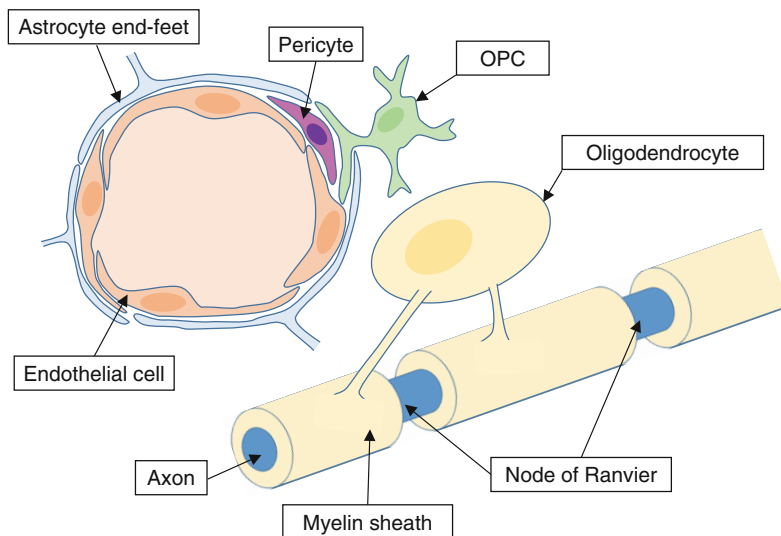


Fig. 1 Schematic of the neurovascular unit in white matter. Neuronal axons, astrocytes, cerebral endothelial cells, pericytes, and oligodendrocytes/OPCs comprise the neurovascular unit in white matter. The concept of the neurovascular unit emphasizes the functional aspects of cell–cell signaling, and in fact, the components are closely located and presumably exchange signals to support their function with each other. This figure was drawn from the authors' previous review article [8]

are an important component in the neurovascular unit in cerebral white matter, and are situated directly next to astrocytes. Recent reports suggest that OPCs are also closely located to cerebral endothelial cells [10, 11]. Although OPCs are most active during development, when they mature into oligodendrocytes to form myelin sheath, a number of OPCs resides in adult white matter to maintain white matter homeostasis. Importantly, OPCs receive support from their neighboring cells to proliferate and differentiate into mature oligodendrocytes in brain.

Another important aspect within the neurovascular unit is the distinct contribution of cerebral endothelium-to-brain function. The cerebral vascular system is one of the major constituents of the brain, traditionally viewed as a passive conduit for blood. However, recent research has shown that this system plays more active roles in maintaining the central nervous system (CNS) homeostasis. For example, cerebral endothelial cells, along with astrocytes and pericytes, form the blood–brain barrier (BBB). The BBB constitutes an anatomical, physiochemical, and biochemical barrier that controls the exchange of materials between blood, brain, and cerebrospinal fluid. BBB breakdown due to endothelial dysfunction is frequently associated with a myriad of neurological pathologies, including chronic CNS diseases [12–14]. Another important function of the cerebral vascular system is that of providing trophic support to neighboring neurons. By releasing trophic factors, cerebral endothelial cells guide developing axons [15], protect neurons against stress [16, 17], and provide a niche for supporting neural stem/progenitor cells (NSPCs) [18]. NSPCs were shown to have direct coupling with cerebral endothelial

cells [19], and in this so-called “neurovascular niche,” cell–cell signaling between cerebral endothelial cells and neuronal precursor cells help mediate and sustain pockets of ongoing neurogenesis and angiogenesis in adult brain [18, 20]. Even under the remodeling phase after brain injury, these close relationships are maintained, and both neurogenesis and angiogenesis occur in the neurovascular niche to promote repair of the brain. Indeed, angiogenic stimulation enhances neurogenesis after stroke [18, 21]. In turn, neuroblasts migrate along perivascular routes and the promotion of neurogenesis enhances vascular re-growth [22].

Much of the research into mechanisms of trophic coupling in the neurovascular unit has focused on endothelium–neuron and endothelium–astrocyte interactions. However, cell–cell interactions between endothelial cells and oligodendrocyte lineage cells are also important in maintaining brain function, especially in white matter. In cerebral white matter, OPCs play essential roles in white matter remodeling/repairing after stroke. Under pathological conditions associated with demyelination, OPCs are activated, recruited to the site of injury, and differentiated into mature oligodendrocytes. These OPC activities proceed in a coordinated fashion in response to complex signaling events involving numerous mediators [23–25]. In the first step, OPCs are activated by phenotypically switching from a mitotically dormant inactive state to a proliferative active state [26, 27]. The next step is OPC recruitment, which is a process involving proliferation and migration. In the final step, the differentiation phase, recruited OPCs exit the mitotic cell cycle and change their phenotype to mature oligodendrocytes, and then, newly created oligodendrocytes establish contact with unmyelinated axons and form functional myelin sheaths around them [25, 28, 29]. The mechanisms underlying this complex process have been the subject of several recent studies, in which the role of cell–cell interaction in remyelination has been increasingly recognized [30, 31]. As noted, some populations of oligodendrocyte lineage cells are located closely to cerebral endothelial cells, and these cells may communicate with each other via secreting soluble factors [10, 11]. In this chapter, we attempt to overview key findings for the crosstalk between cerebral endothelial cells and oligodendrocyte lineage cells, focusing on oligodendrocyte damage and repair after stroke.

2 OPC Differentiation to Oligodendrocytes Under Normal and Pathological Conditions

Oligodendrocytes are one of the major glial cells in the CNS, and produce a lipid-rich membrane called myelin. Each oligodendrocyte can enwrap up to 60 axonal segments, thereby enabling fast and salutatory nerve impulse conduction [32]. During development, OPCs are first generated in the germinal zones, where they will proliferate. They then migrate to both gray and white matter areas, where most will differentiate into mature oligodendrocytes and form myelin sheaths.

Although myelinated tracts are formed early in life, renewal of myelin/oligodendrocyte continues throughout adult life [33–35]. Myelin in the adult CNS maintains some plasticity in response to changes in neural activity [36] and brain injury [37].

Under normal conditions in the adult brain, most subventricular zone (SVZ) progenitor cells give rise to neuronal lineage cells. They migrate along the rostral migratory stream (RMS) to the olfactory bulbs, where they terminate and differentiate into mature interneurons [38–40]. Oligodendrocytes can also be generated from SVZ cells in the adult brain, and newly generated OPCs migrate towards the corpus callosum and the white matter tracts of striatum and fimbria fornix [41]. However, the proportion of SVZ progenitor cells that differentiate into oligodendrocyte lineage cells decrease after the early postnatal period [39]. Interestingly, in the SVZ, neuronal and oligodendroglial progenies constitute separate lineages under physiological conditions. Using continuous live imaging and single-cell tracking of NSPCs, Ortega et al. have demonstrated that a single NSPC and its offsprings in the SVZ would not result in both neuronal and oligodendroglial progenies [42]. Furthermore, the adult SVZ is highly regionalized. The neuronal progeny with distinct identity is generated in different areas along the dorsoventral and rostrocaudal axes [43, 44]. In addition, clones fated to generate oligodendrocytes are prevalent in NSPCs isolated from dorsolateral SVZ. On the contrary, ventrolateral SVZ regions consist of both neuronal and astroglial progenies with few oligodendroglial progeny [42, 45, 46].

SVZ progenitor cells in the adult brain show some lineage plasticity under pathological conditions. After CNS damage, a number of progenitors migrate out of the RMS to the injured site. The fate of these progenitor cells can be dynamically altered according to the disease type. The fate of SVZ progenitor cells can shift from NSPCs to OPCs after demyelination, and these newly generated OPCs proliferate and migrate to the lesion areas [39, 47–49]. In a model of experimental autoimmune encephalomyelitis (EAE), enhanced proliferation and migration of SVZ NSPCs are observed, and these mobilized cells give rise to oligodendrocytes and astrocytes without neurons in the injured white matter [49]. In addition, demyelination would change the fate of glutamic acid decarboxylase 65 (GAD65)/doublecortin (Dcx)-expressing NSPCs derived from the adult SVZ to generate oligodendrocytes, rather than neurons, in corpus callosum [47]. This process may restore developmental myelination to some extent; NSPCs that generate oligodendrocytes migrate from SVZ to developing white matter, where they stop dividing to differentiate and myelinate axons [47, 50].

Although the most studied disease with oligodendrogenesis after white matter injury is multiple sclerosis (MS), there are some findings reported for research studies using stroke models. After brain ischemia, immature oligodendrocytes proliferate in the regions surrounding the lateral ventricles [51] and the infarction site (peri-infarct areas) [52], with a delayed increase in the number of mature oligodendrocytes in peri-infarct areas [51]. As discussed, under normal conditions, oligodendrocytes can be generated in the SVZ and migrate to white matter tracts of the corpus callosum, fimbria fornix, and striatum [41]. After demyelination, SVZ-generated OPCs proliferate and migrate to peri-lesional areas, attempting to differentiate into mature oligodendrocyte and remyelinate axons in response to numerous mediators [37, 47]. This finding has been reported in rodent models of ischemic brain injury, in which OPCs are generated by SVZ neuronal progenitor cells and later become mature myelinating oligodendrocytes, which help restore damaged

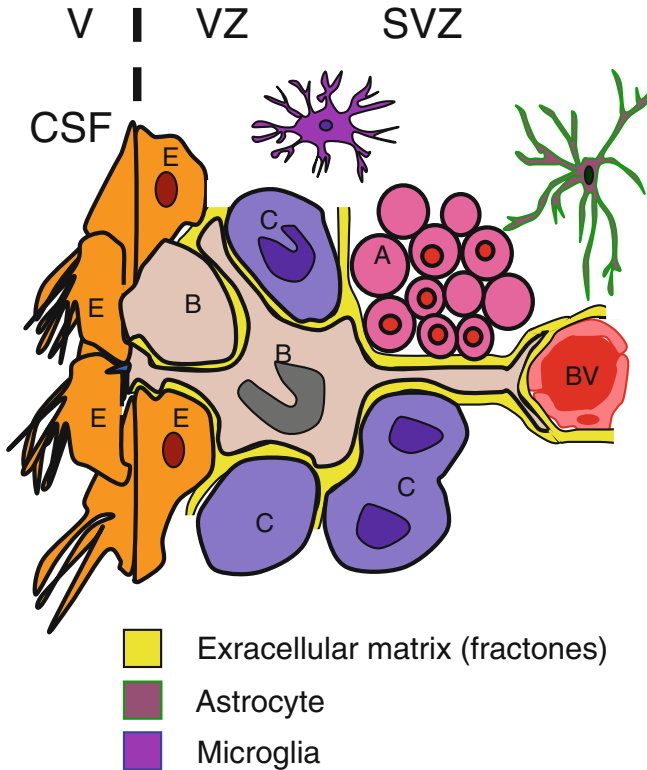


Fig. 2 Schematic of interplay for SVZ cells. The subventricular zone (SVZ) and ventricular zone (VZ) line the lateral ventricles (V) in the brain. Type B cells (B) contact the ventricle (V) containing cerebrospinal fluid (CSF) through specialized apical processes. The processes contain a single primary cilium, which is surrounded by a rosette of ependymal cells (E) with large apical surfaces forming pinwheel-like structures. On the other side, the type B cells have long basal processes with specialized endings that frequently contact blood vessels (BV). The type B cells also contact their progeny, i.e., type C cells (C) and the chains of migrating type A neuroblasts (A). The V-SVZ includes extracellular matrix (fractones) that contacts all the cell types including blood vessels, microglia, and astrocytes in this region. This figure was drawn from the authors’ previous review article [59]

white matter [53–56]. Thus, increasing the recruitment of immature oligodendrocytes to the injured site and promoting their progression to mature oligodendrocytes will enhance white matter repair after injury. The process of oligodendrogenesis is influenced by many intrinsic and extrinsic factors from various types of cells, thus offering a number of pathways for potential therapeutic interventions [57].

NSPCs in the SVZ display diverse interactions with their neighboring environments [38, 58] (Fig. 2). On one side of SVZ, type B cells are surrounded by multi-ciliated non-dividing ependymal cells, which form pinwheel-like structures on the ventricular surface. These cells are in direct contact with the cerebrospinal fluid (CSF) through a short non-motile primary cilium that extends towards the

ventricle. On the other side, type B cells interact with the extensive network of blood vessels with a long basal process. Type B cells also attach to type C cells and chains of young neurons (type A cells) by the extracellular matrix. Proliferating type C cells are closely located to their progenitors, and are also often in close proximity to blood vessels [60]. Type B cells interact with one another by gap and adherens junctions, the same as in ependymal cells [61]. Furthermore, the adult SVZ possesses a highly organized basement membrane, which is absent in other areas of the brain. Overall, the SVZ is poised to receive informational inputs via cell–cell and cell–matrix contacts. The integration of these multifaceted external cues (e.g., extracellular signals from the vasculature, extracellular matrix, and the cerebrospinal fluid) to intrinsic factors leads to the determination of the fate and behavior of each cell lineage. In the next section, we will discuss how cerebral endothelial cells (and blood vessels) support oligodendrocyte function, including OPC generation from NSPCs.

3 Roles of Cerebral Endothelium on NSPC/OPC Function

The vasculature is an integral component of the SVZ stem cell niche that possesses specialized properties in regulating stem cell proliferation and regeneration [60, 62]. Endothelial cells secrete factors that contribute to stem cell self-renewal or proliferation. Co-culture of endothelial cells with NPSCs enhance the *in vitro* neurosphere generation from embryonic progenitors [63]. NSPCs were shown to have direct coupling with cerebral endothelial cells [19], and various kinds of perivascular regulators, including growth factors, purinergic signaling, nitric oxide signaling, and chemokines, contribute to cell genesis and fate determination in the SVZ [64].

Dividing progenitor cells (type B cells) and their transit-amplifying type C cells lie adjacent to the extensive planar vascular plexus in the SVZ. Approximately 47% of dividing type B cells and 46% of type C cells are found within 5 microns of the vasculature. During homeostasis and regeneration, type B cells and type C cells directly contact SVZ blood vessel sites devoid of astrocyte end-feet and pericyte coverage [60, 62]. Most dividing type B and type C cells are close to these sites, highlighting the importance of vasculature in supporting progenitor cell function. By contrast, most migrating neuroblasts are more distal to the vasculature (only 14% are within 5 μm) compared to type B and type C cells, even though blood vessels run parallel to the aggregates of migrating neuroblast chains in the dorsal aspect of the SVZ and in the RMS. However, it still remains to be understood whether neuronal differentiation occurs in response to leaving the vascular bed or whether cells leave the vasculature after they are differentiated [62].

Blood vessels in the SVZ region also serve as a scaffold for long-distance migration of neuroblasts from SVZ to the olfactory bulb, potentially through the release of chemoattractant (e.g., brain-derived neurotrophic factor (BDNF), vascular endothelial growth factor (VEGF)), and chemorepulsive factors (e.g., semaphorins, ephrins) [65–68]. Migrating neuroblasts are ensheathed by a layer of astrocyte

processes and use each other as guides in the migration process toward the olfactory bulb. Similarly, in animal stroke models [18] and human stroke patients [69], long-distance migration of newly born immature neurons from SVZ to peri-infarct cortex is observed. Stromal cell-derived factor 1 (SDF-1)/C-X-C chemokine receptor type 4 (CXCR4) signaling assists blood vessel- and astrocyte-associated migration of adult SVZ progenitors after cortical injury [70]. Recent studies have identified that SDF-1/CXCR4-mediated signaling is a critical homing factor in the SVZ niche. CXCR4 is expressed by all progenitor cells in the SVZ. SDF-1 is expressed in the SVZ blood vessels, but the ependymal cells that line the lateral ventricles express higher levels of SDF-1 to create a concentration gradient. SDF-1 increases integrin $\alpha 6$ and epidermal growth factor receptor expression in activated type B and type C cells, enhancing their activated states and ability to bind laminin in the vascular niche [20]. SDF-1 also increases the motility of type A neuroblasts. These type A cells express lower levels of integrin $\alpha 6$, which might promote evacuating from the vascular niche.

As noted, cerebral endothelial cells within the cerebral blood vessels support neighboring cells by secreting trophic factors. Recent studies confirmed that cerebral endothelial cells regulate the function of oligodendrocyte lineage cells. In a co-culture system of endothelium with NSPCs, the chemokine CCL2/MCP-1 mediates the interaction between endothelium and neural precursor cells to promote the differentiation of NSPCs into oligodendrocytes [71]. Another study used the *in vitro* media-transfer system to show that conditioned medium from endothelial cells promotes the differentiation of NSPCs into oligodendrocyte lineage cells [72]. In addition, cerebral endothelial cells and OPCs may provide an oligovascular niche to promote the proliferation and migration of OPCs [73–75]. This endothelium-to-OPC supportive signaling would be attenuated by excessive oxidative stress [76], supporting the idea that oligodendrocyte/myelin maintenance and renewal is disturbed under the acute phase of stroke conditions. The detailed mechanism (e.g., mediators, etc.) from endothelial cells to OPCs will be discussed in the Sect. 5.1 and 5.5 below.

4 Roles of Oligodendrocyte Lineage Cells in Vascular Remodeling After Stroke

Oligodendrocyte lineage cells are essential for oligodendrogenesis and remyelination after white matter damage. However, the role of oligodendrocyte lineage cells in post-stroke repair is not restricted to restoration of damaged myelin. Oligodendrocyte lineage cells may give support to other cell types during the recovery process after ischemic injury. While there are only a limited number of studies describing this process, we will briefly introduce how oligodendrocyte lineage cells contribute to vascular remodeling after brain damage including ischemic conditions.

Cell–cell signaling between cerebral endothelium and OPCs contribute to vascular remodeling after stroke, in processes mediated by secretion of trophic factors and extracellular matrix proteins such as matrix metalloproteinases (MMPs) [77]. BDNF is one such trophic factor, which promotes endothelial cell survival and angiogenesis [78]. It has been suggested that BDNF derived from oligodendrocyte lineage cells do support angiogenesis, since BDNF mRNA is present in oligodendrocytes, and the physical proximity of oligodendrocytes to cerebral endothelial cells would allow oligodendrocyte-secreted BDNF to interact with the endothelial cells [77, 79]. Transforming growth factor (TGF- β) is another factor that is essential for vascular development and maturation. OPCs are one of the cell types in the neurovascular unit secreting TGF- β 1 and this OPC-derived TGF- β 1 has a role in maintaining BBB integrity [11]. Treating endothelial cells with OPC-conditioned media increased tight junction proteins in endothelial cells; while OPC-specific TGF- β 1 knock-out mice exhibited cerebral hemorrhage, indicating loss of BBB function [11]. Recent research examining OPC contribution to postnatal angiogenesis showed that OPCs regulate angiogenesis within the white matter through OPC-encoded hypoxia-inducible factor (HIF) signaling. OPC-specific HIF stabilization resulted in increased expression of the proangiogenic genes *Wnt7a/7b*, and promoted angiogenesis and endothelial proliferation in vivo [80]. Further studies are needed to determine whether this OPC-specific HIF pathway plays a role in recovery after CNS injury. In addition to OPCs, mature oligodendrocytes would promote vascular remodeling after white matter damage. Cultured rat oligodendrocytes secrete MMP-9 in response to inflammatory responses. Importantly, conditioned media from IL-1 β -stimulated oligodendrocytes would promote in vitro angiogenesis compared to normal oligodendrocyte-conditioned-media [81]. Similarly, in a mouse model of focal stroke, oligodendrocytes were shown to produce MMP-9 during the recovery phase, which may lead to vascular remodeling in white matter [81].

5 Possible Mediators for Endothelium–Oligodendrocyte Interaction

Crosstalk between the vascular and neuronal compartments in the neurovascular niche is mediated by an exchange of soluble signals [3, 6, 7, 17, 63, 82–84]. Many of these trophic factors from cerebral endothelial cells may also affect oligodendrocyte lineage cells. Moreover, a growing literature suggests that similar to cerebral endothelium, oligodendrocyte lineage cells could work as a “bank” for trophic factors. Therefore, it would be reasonable to hypothesize that the endothelium–oligodendrocyte interaction is also extensively mediated by secreting factors. Although the precise regulatory mechanisms that underlie angiogenesis and oligodendrocytes in the oligovascular niche still require further research to be elucidated, we will summarize several candidates of mediators for the crosstalk between cerebral endothelium and oligodendrocytes in this section.

5.1 BDNF

BDNF is a member of the “neurotrophin” family of growth factors, which are related to the canonical nerve growth factor (NGF). BDNF is well-known to act on neurons to support the survival of existing neurons and enhance the growth and differentiation of new neurons and synapses [85, 86]. BDNF can also modulate non-neuronal cell types. Irrefutably, BDNF is one of the major mediators for endothelium–oligodendrocyte interaction.

BDNF is known to regulate oligodendrocyte function in several ways. Firstly, BDNF promotes OPC proliferation and differentiation into mature oligodendrocytes. Cell culture and *in vivo* animal studies suggested that Trk-B receptor and ERK pathway mediated the supportive effects of BDNF [87–90]. Secondly, even under pathological conditions, BDNF can be supportive for oligodendrocyte lineage cells. In *in vivo* white matter injury models (spinal cord injury or cuprizone-treated model), BDNF was shown to play an important role in regulating the number of oligodendrocyte lineage cells after demyelination [91, 92]. Thirdly, BDNF may also work on NSPCs to trigger their differentiation into oligodendrocyte lineage cells. Chen et al. reported that BDNF enhanced the cell commitment of NSPCs to neuronal and oligodendrocytic fates by activating Wnt/ β -catenin signaling pathway *in vitro* [93]. Since cerebral endothelial cells are the major BDNF-producing cell types, cerebral endothelial cells may support oligodendrocyte lineage cells via BDNF signaling. As introduced above, endothelial-derived BDNF was shown to promote the proliferation of OPCs *in vitro* [94]. But importantly, stressed endothelial cells secrete less BDNF and can no longer support OPCs [94], indicating that endothelial–oligodendrocyte interaction is highly dependent on their cellular conditions.

BDNF also promotes endothelial cell survival and induces angiogenesis in the brain [95]. While cell–cell trophic interactions are generally considered as “two-way,” there have been still no direct proofs that BDNF mediates signals from oligodendrocytes to endothelial cells. Nevertheless, several lines of evidence strongly support the idea that oligodendrocyte lineage cells may produce BDNF to support endothelial cells. *In situ* hybridization and immunocytochemical studies identified expressions of BDNF mRNA/protein in cultured basal forebrain oligodendrocytes [79, 96]. The physiological relevance of these *in vitro* studies is supported by detection of BDNF in oligodendrocytes *in vivo*. For example, BDNF mRNAs are localized in subpopulations of myelin-basic-protein-positive mature oligodendrocytes in the basal forebrain, cingulate cortex, and corpus callosum of postnatal day 7 rats [79]. Similarly, BDNF mRNA and protein are also expressed in subsets of APC (adenomatous polyposis coli)-positive oligodendrocytes of adult spinal cords [97, 98]. The oligodendrocytic BDNF is bioactive because conditioned media from oligodendrocyte cultures increased cholinergic neuronal function and the effect was partially blocked by co-treatment with anti-BDNF neutralizing antibody [79]. Considering the fact that oligodendrocyte lineage cells are often located close to cerebral endothelial cells, it would be reasonable to think that oligodendrocytic BDNF may interact to some extent with the cerebral vascular system.

5.2 *Fibroblast Growth Factor-2*

Similar to BDNF, fibroblast growth factor-2 (FGF-2/bFGF) can be proposed as an important mediator for endothelium–oligodendrocyte interactions. FGF-2 is a potent stimulator of endothelial cell migration, proliferation, sprouting, and tube formation. In addition, past studies have substantially revealed the effects of FGF-2 on oligodendrocyte function. FGF-2 by itself stimulates proliferation of late-stage OPCs and blocks their terminal differentiation into mature oligodendrocytes *in vitro*. Upon removal of FGF-2 from the cell culture medium of OPCs, the cells readily enter the terminal differentiation [99–101]. In addition, FGF-2 cooperates with platelet-derived growth factor (PDGF) to upregulate the expression of PDGF-receptor-alpha for OPC proliferation [101, 102]. As for OPC migration, the importance of FGF signaling in early-stage OPCs has been evaluated both *in vivo* and *in vitro*. Using an *in vivo* transplantation approach, Osterhout et al. demonstrated that OPCs with dominant-negative FGF receptor 1 failed to migrate [103]. Subsequent *in vitro* study showed that in response to FGF-2 stimulation, OPCs growing in an agarose drop successfully move out from the drop [104]. Another study suggested that FGF-2 also enhanced the migration of pre-OPCs from oligospheres, but might not induce the migration of late-stage OPCs [105]. While still controversial, FGF-2 may affect OPC survival. FGF-2 can prevent OPCs from apoptotic stress [106], but this protective effect was not observed in optic nerve OPCs [107]. Since the cellular localization of FGF-2 is observed in cerebral endothelium both in normal and pathological conditions [108], future studies are warranted to examine how FGF-2 modulates the dynamics of endothelium–oligodendrocyte crosstalk under normal conditions as well as in the acute and chronic phases after brain injury.

5.3 *TGF- β*

TGF- β is a prototypic member of a large family of pleiotropic cytokines, and may also be a potent modulator in the oligovascular niche. Many cells have been reported to produce TGF- β , and TGF- β is known to modulate a diverse array of cellular function [109]. In terms of its involvement in endothelium–oligodendrocyte interactions, oligodendrocytes are reported to express TGF- β *in vitro* in cell culture systems [110] and *in vivo* in spinal cords [111]. TGF- β contributes to angiogenesis by stabilizing newly formed capillary sprouts [112]. Thus far, many studies in mouse and human have demonstrated its pivotal roles in modulating angiogenesis after brain injury, such as stroke [113–118]. In addition, dysregulation of TGF- β signaling may cause hereditary vascular disorders [109]. For example, mutations in TGF- β receptors lead to hereditary hemorrhagic telangiectasia [119], and in the cerebral white matter, accumulation of TGF- β 1 due to HtrA1 mutation is associated with a hereditary disorder CRASIL (cerebral autosomal recessive arteriopathy with

subcortical infarcts and leukoencephalopathy) [120]. In contrast, the roles of TGF- β on oligodendrocytic function are still understudied. An early report suggests that TGF- β affects the migration of cultured OPCs [121]. Using the oligodendrocyte precursor cell line OLI-neu, TGF- β was shown to upregulate a chondroitin sulfate proteoglycan DSD-1-PG on the cell surface [122], indicating that TGF- β may modulate the cell adhesion property of OPCs to change cell motility.

5.4 *Adrenomedullin*

Adrenomedullin (AM) was originally isolated from pheochromocytoma cells [123]. AM has a variety of actions on the vascular systems, such as endothelial survival/proliferation, vasodilatation, regulation of BBB permeability, and modulation of oxidative stress levels in endothelium [124, 125]. AM is secreted from various organs, and in the CNS, AM is mainly expressed in neurons and cerebral endothelium [123, 126]. AM has been shown to reduce infarct volumes in transient stroke models [127, 128]. Under stroke conditions, AM expression is markedly increased via the hypoxia-inducible factor-1 signaling [129], indicating that AM might support vascular remodeling/repairing after brain injury. Interestingly, in the mouse model of chronic cerebral hypoperfusion (i.e., vascular dementia model), AM was demonstrated to be protective toward cerebral white matter [130], where most oligodendrocytes are populated. In vitro experiments also confirmed that AM promoted OPC maturation under prolonged hypoxic conditions [131]. These reports suggest that AM may be important in oligodendrocyte regeneration after white matter damage. On the contrary, AM is released by oligodendrocyte lineage cells as well as cerebral endothelial cells. Uezono et al. detected AM mRNAs in human oligodendroglial cell line, and showed the potential effects of AM in modulating oligodendrocyte function through AM receptors in oligodendrocytes [132]. Importantly, past studies have demonstrated that AM increases several growth factors such as VEGF and FGF [133–136]. Taken together, AM might work as a “master” modulator for growth factors in the oligovascular niche.

5.5 *VEGF*

VEGF is a primary regulator of angiogenesis by stimulating endothelial cell proliferation, migration, and tube formation [83]. But it is now well recognized that VEGF is not solely an endothelial mediator, rather it may represent one of the major mediators that signal to more than one type of cell in the neurovascular unit [137–139]. For example, VEGF signaling plays a key role in neuronal migration and CNS development [140]. In addition, VEGF may also affect oligodendrocyte lineage cells. OPCs express the VEGF receptor, Flk-1 (also known as KDR or VEGF-receptor-2) [74], which is primarily responsible for VEGF-induced angiogenesis.

The same study suggested that VEGF-A significantly accelerated the motility of OPCs through Flk-1, and this effect was partly mediated by ROS production [74]. Notably, conditioned media from cerebral endothelial cells promoted both OPC proliferation and migration. However, endothelial-derived VEGF-A may participate in the OPC migration but not proliferation [141]. In addition to VEGF-A, other VEGF families and VEGF receptors may also be involved in oligodendrocyte function. Le Bras et al. demonstrated that VEGF-C promoted OPC proliferation through VEGF-receptor-3 [142]. Taken together, these findings indicate that VEGFs/VEGF receptors may play a central role in the endothelial–oligodendrocyte trophic coupling.

If VEGF is an important modulator for the oligovascular niche, we may need to consider the “biphasic actions” of VEGF to comprehend the dynamics of trophic coupling between cerebral endothelium and oligodendrocytes. VEGF is known to act as a deleterious factor during the acute phase of stroke. For example, VEGF administration during ischemic insults worsens BBB leakage [143, 144]. On the other hand, VEGF can be a primary regulator of angiogenesis. VEGF would trigger remodeling responses in endothelial cells (i.e., accelerating angiogenesis) after brain damage, such as stroke. In fact, infusing VEGF into the lateral ventricles stimulated angiogenesis and decreased infarct volumes in rodent models of focal cerebral ischemia [145]. An increase in angiogenesis by VEGF in rats was also associated with reduced neurological deficits after stroke [146]. Similar effects were also reported in neonatal focal rodent stroke [147]. Moreover, in transgenic mice that overexpressed human VEGF165, brain microvessel density was significantly elevated compared to wild-type mice before ischemia, and the microvessel density was higher 3 days after stroke onset [148]. Therefore, depending on the context of the microenvironment in the oligovascular niche, VEGF may exert different effects on endothelium–oligodendrocyte interactions that maintain and remodel white matter function.

5.6 MMPs

MMPs comprise a family of zinc endopeptidases, and their activities are tightly regulated by tissue inhibitor of metalloproteinases (TIMPs). This large MMP network plays major roles in the physiology and pathology of the mammalian CNS, including stroke (MMP-2/3/7/9/13, TIMP-1/2 [149]), vascular dementia (MMP-2, 3, 9 [150, 151]), and CADASIL (cerebral autosomal-dominant arteriopathy with subcortical infarcts and leukoencephalopathy: TIMP-3 [152]). Similar to VEGF, MMPs are well-known biphasic modulators in the CNS. During the acute phase of stroke, MMP-2/3/9/13 are deleterious, i.e., they degrade the extracellular matrix that comprises the basal lamina, which could directly damage BBB [153]. In addition, proteolysis of the neurovascular matrix by MMP-9 after stroke can trigger anoikis-like neuronal death [154]. However, during delayed phases of stroke recovery, some of these proteinases (MMP-2/3/9) may play beneficial roles [149, 153, 155]. In a mouse stroke model, peri-infarct cortical areas demonstrate a secondary

elevation in MMP-9 in endothelial and glial cells within networks of regrowing microvessels [156]. Inhibition of MMPs during this delayed phase disrupted brain repairing with the induction of hemorrhagic and malformed blood vessels. Moreover, signals of MMP-9 were observed in the migrating neuroblasts from the subventricular zone after brain injury, and inhibition of these MMPs also blocked the movement of these neuroblasts originally headed towards damaged brain [157].

As MMPs play multiple roles in the neurovascular unit, MMPs may mediate the crosstalk between cerebral endothelium and oligodendrocyte. Thus far, several studies imply the essential roles of MMP network in oligodendrocytic function. For example, MMP-9 mediated the extension of oligodendrocytic processes in cell culture [158], and MMP-9 expression level was increased during myelin formation in the optic nerve in vivo [159]. Moreover, after white matter injury, MMP-9 removed injury-induced deposition of inhibitory NG2 proteoglycan, which is an essential step for OPCs to differentiate into mature oligodendrocytes for remyelination [160]. In addition, MMP-12 was shown to cause demyelination, macrophage infiltration, and motor deficits in a mouse model of virus-induced multiple sclerosis [161]. Tissue inhibitor of metalloproteinases might be also involved in cellular function/survival of oligodendrocyte lineage cells. In a mouse model of focal stroke ischemia, mice with TIMP-3 deficiency had a higher number of immature oligodendrocytes after injury [162]. In addition to being the recipient cell for MMP actions, oligodendrocyte lineage cells may in turn produce MMPs to send signals to cerebral endothelial cells. As noted, OPCs respond quickly to a stress stimulation after white matter damage and release MMP-9 which induces early BBB leakage in a mouse model of prolonged hypoperfusion model [10]. On the other hand, during the chronic phase of white matter injury, oligodendrocytic MMP-9 may promote vascular remodeling [81]. Hence, as in the neurovascular unit, MMPs contribute different effects to the dynamics of cellular interactions in the oligovascular niche due to their biphasic property.

6 Conclusion Remarks

The concept of neurovascular unit emphasizes that cell–cell interaction is critical to maintain normal brain function as well as brain remodeling after injury. Within the conceptual framework of neurovascular unit, cerebral endothelial cells are particularly important in releasing soluble factors to nourish neighboring cells, such as astrocytes or neurons. As we discussed in this chapter, cerebral endothelial cells may also support oligodendrocyte lineage cells in the white matter. On the other hand, oligodendrocyte-derived factors can in turn modulate cerebral vascular systems. In the so-called oligovascular niche (i.e., microenvironment between cerebral endothelium and oligodendrocytes), cerebral endothelium and oligodendrocyte lineage cells may cooperate in maintaining white matter homeostasis. Although some key mediators for endothelium–oligodendrocyte crosstalk were briefly discussed here, the precise underlying mechanisms still remain to be elucidated.

Importantly, there may be overlap between factors in the oligovascular niche and the well-established neurovascular niche. Therefore, a deeper analysis, perhaps using subtractive approaches, may be required in order to rigorously define the regulatory signals that are truly unique to the oligovascular niche. As white matter injury is a key part of most CNS diseases, understanding the cell–cell interaction between cerebral endothelium and oligodendrocytes may lead to effective therapeutic approaches for white matter-related diseases, such as stroke and vascular dementia.

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Part III
Peripheral Immune Cells in Stroke

The Peripheral Immune Response to Stroke

Josef Anrather

1 Introduction

Historically, the central nervous system (CNS) has been viewed as a place with limited immune surveillance and capacity to mount immune responses. This has led to attribute “immunological privilege” to the CNS, a term first coined by Billingham and Boswell [1]. Already in the nineteenth century the Dutch ophthalmologist van Dooremaal [2] described the phenomenon of immune privilege by demonstrating the long-term survival of mouse skin grafts placed in the anterior chamber of a dog’s eye. However, within the brain, immune privilege does not apply to all structures. Murphy and Sturm [3] observed that allogeneic tumor transplants were readily rejected when they came in contact with the cerebral ventricles while transplants that were embedded within the brain parenchyma did not elicit an immune response and showed prolonged survival times. These studies were later extended by Medawar [4] to show that allogeneic skin grafts transplanted into the anterior chamber of the eye or the brain of rabbits were efficiently rejected when the host was previously immunized with donor-derived cells and that this rejection was dependent on vascularization of the graft. The resistance of the CNS to mount an effective immune response also extends to the innate immune system. Bacterial endotoxin injected into the brain parenchyma triggers less inflammatory response as measured by immune cell infiltration, than injection of the same endotoxin dose into the skin [5]. Similar to observations with the adaptive immune response, the reduction in endotoxin-mediated inflammation was not found when endotoxin was injected into the ventricles. These studies collectively suggest that the CNS is not immune privileged per se, but there are mechanisms in place to suppress both the innate and

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adaptive immune responses. These include the blood–brain barrier (BBB) and the glia limitans, which minimize the exchange of potentially antigenic macromolecules and pro-inflammatory mediators (e.g., cytokines, chemokines, DAMP) and pose a barrier for easy immune cell entry from the circulation. In addition, the brain features a generally anti-inflammatory milieu under homeostatic conditions sustained by the expression by highly immunosuppressive cytokines such as transforming growth factor- β (TGF- β) and insulin-like growth factor-1 (IGF-1). Once these structural and molecular barriers are removed, as it is the case during ischemic brain injury, inflammation can proceed “as planned” and contribute to the outcome of stroke. This chapter will briefly discuss immune cells participating in the response to cerebral ischemia, deliberate potential entry points of these cells to gain access to the ischemic tissue, and outline the role of the ischemic brain in shaping the peripheral immune response.

2 Participating Immune Cells

Post-ischemic inflammation is characterized by activation of brain-resident leukocytes, microglia, astroglia, and endothelium, and the orchestrated recruitment of various blood-borne immune cells.

Microglia, monocytes/macrophages, and dendritic cells—Historically it has been difficult to separate the relative contribution of resident (microglia) and blood (monocyte)-derived macrophages to ischemic brain injury. Because these cells are not readily distinguishable by morphology or marker gene expression, most results derived from in vivo studies could either implicate microglia or macrophages or both. Only recently, multi-label flow cytometry of brain immune cells and bone marrow chimeric animals that express discernable markers (fluorescent proteins or CD45 alloantigen) in have allowed a clear distinction of microglia and blood-derived myeloid cells. A recently developed mouse model, that expresses myeloid-restricted tamoxifen-inducible Cre-recombinase, opens the possibility to generate microglia-selective gene deletions to further dissect the contribution of brain resident and hematogenous myeloid cells in stroke [6]. Development of mice expressing fluorescent proteins in different classes or functionally distinct subsets of hematogenous immune cells should also benefit studies that investigate origin, trafficking, and fate of immune cells after cerebral ischemia. Analysis of microglia/macrophages in brain ischemia is further complicated by the fact that the cellular and functional phenotype of these cells in the post-ischemic microenvironment is not static but fluent as exemplified by the rapid transformation of microglia after cerebral ischemia. Microglial activation occurs before the appearance of neuronal cell death [7] and this early response is characterized by increased arborisation and exploratory behavior giving way to de-arborisation and amoeboid transformation within the first 24 h after stroke while the cells remain stationary [8, 9]. In a study in bone marrow chimeric mice expressing green fluorescent protein (GFP) Schilling et al. [10] found that microglia contributed to increased numbers of macrophage-like

cells during the first days after cerebral ischemia while GFP⁺ blood-borne cells appeared in robust numbers only after day 2. While pro-inflammatory and cytotoxic activities of microglia have been demonstrated in vitro, eliminating proliferating microglia/macrophages by ganciclovir treatment in mice that express herpes simplex virus-1-thymidine kinase in the myeloid lineage increased ischemic brain injury and reduced expression of the neurotrophic cytokine IGF-1 [11]. Similar to microglia, monocyte-derived macrophages can exhibit various phenotypes. Blood monocytes exist as two functionally distinct subpopulations (inflammatory and patrolling) best characterized by their expression of C-C chemokine receptor 2 (CCR2) and C-X3-C chemokine receptor 1 (CX3CR1), both of which are important for monocyte entry into the ischemic brain [12–14], and the relative abundance of these subsets in the blood has been linked to clinical stroke outcome [15, 16]. Initially, infiltrating monocytes are of the “inflammatory” subtype while “patrolling” monocytes are prevalent at later time points [17]. The pathophysiological importance of this finding, however, has yet to be determined. In experimental stroke, monocytes/macrophages infiltrate the brain parenchyma 48–72 h after ischemia and can persist for several weeks [10]. There is also evidence that inhibition of adhesion molecules involved in lymphocyte and monocyte trafficking, such as the very late activation antigen-4 (VLA-4) or the lymphocyte function antigen-1 (LIF-1), can reduce infarct volumes in stroke models [18–21]. These studies suggest that lymphocytes and monocytes, in addition to neutrophils, participate in post-ischemic brain inflammation [22]. However, the precise role of monocyte-derived macrophages in the deleterious effects of post-ischemic inflammation remains to be defined. In addition, depending on molecular cues encountered in the post-ischemic brain parenchyma, monocytes/macrophages can undergo classical (M1, pro-inflammatory) or alternative (M2, anti-inflammatory in most settings) activation, thus contributing to the resolution of inflammation [8, 23]. Phagocytosis, which is mainly seen in microglia and not macrophages and is tied to MER proto-oncogene tyrosine kinase (MerTK), milk fat globule-EGF factor 8 protein (MFG-E8) and triggering receptor expressed on myeloid cells-2 (TREM2) expression [24, 25], can contribute to delayed neuronal cell death after ischemia [24, 26]. A subset of microglia and infiltrating macrophages express the dendritic cell (DC) marker CD11c and MHC class-II molecules [27]. Using chimeric animals that express yellow fluorescent protein (YFP) under the control of the CD11c promoter in bone marrow-derived cells, the same study showed that DCs originate from brain-resident myeloid cells early after ischemia, while peripheral immune cells contribute the majority of CD11c⁺/YFP⁺ cells 3 days after stroke [27]. Although brain DCs have the capacity to present antigen and induce T cell proliferation in vitro [28], it remains to be shown whether these cells engage in antigen presentation after brain ischemia as predicted by some studies [29].

Neutrophils—Although intravascular adhesion of neutrophils is a relatively early post-ischemic event, parenchymal accumulation is generally observed later. Nevertheless, neutrophils are among the first hematogenous immune cells found in the brain after experimental stroke peaking at 48–72 h in most models and declining rapidly afterwards. Although it is not clear whether they enter the brain parenchyma

under all circumstances [30], there is evidence that neutrophils contribute to post-ischemic inflammation by limiting tissue perfusion due to intravascular clogging [31, 32], destabilizing the BBB by releasing matrix-metalloproteinases (MMP) [33, 34], and by generating reactive oxygen and nitrogen species [35, 36]. However, a cause-and-effect relationship between the extent of neutrophil trafficking and the severity of ischemic damage has not been firmly established [37]. Attesting to the complex role of neutrophils in cerebral ischemic injury, a recent study postulates a protective role of neutrophils, that have undergone N2 polarization as characterized by Ym1 (chitinase 3-like 3, *Chi3l3*) expression, in stroke pathology [38].

Mast cells—Are brain resident immune cells located in the perivascular space surrounding brain parenchymal vessel and in meninges that are rich in peptidases and vasoactive molecules. Mast cells are activated early after cerebral ischemia and contribute to the BBB breakdown and brain edema by releasing gelatinase and vasoactive mediators [39, 40].

Lymphocytes—T cells are detrimental in the early phase of ischemia and lymphocyte-deficient mice are protected in models of focal ischemia [41, 42]. The mechanism does not involve classical antigen-mediated T cell activation and the cytotoxic activity might be tied to innate T cell functions [42]. Accordingly, IL-17 secreting $\gamma\delta$ T cells, that do not undergo classical antigen-dependent T cell activation, have been shown to contribute to ischemic injury [43, 44]. While effector T lymphocytes may contribute to focal ischemic injury [41, 45], regulatory T cells (Treg) could have a protective effect by downregulating post-ischemic inflammation. Tregs appear in the ischemic tissue after the acute phase and confer neuroprotection by IL-10 secretion, an effect that might be antigen-independent [46–49]. Similarly, regulatory B cells (Breg) confer neuroprotection through an IL-10 dependent mechanism, but do not enter the ischemic brain [50–52]. Recent work that has extended the range of stroke-related inflammation beyond innate immune mechanisms to include adaptive immunity [53, 54]. Disruption of the blood–brain barrier during acute stroke releases novel CNS antigens that are normally sequestered in the brain and exposes them to the systemic immune system. Meningeal lymphatic vessels that run along the venous sinus and terminate in the deep cervical lymph nodes [55], are likely to be involved in the cranial export of antigenic macromolecules and antigen-presenting cells (APC) after ischemic brain injury (Fig. 1). Evidence for antigen-specific T cell reactivity has been found in animal models of stroke. There is an ever increasing list of CNS-derived peptides that can induce a peripheral T cell response after stroke including—but not limited to—peptides derived from myelin basic protein (MBP), neuron-specific enolase (NSE), proteolipid protein (PLP), NMDA receptor 2A (NR2A), and microtubule-associated protein (MAP) [54, 56]. Peptide-reactive B and T cells can be found in cervical lymph nodes and spleen as early as 4 days after transient focal ischemia in mice [56]. Among the T cells, the response was similar in memory (CD4⁺) and effector (CD8⁺) T cells. In human stroke, APCs loaded with CNS-derived peptides have been found in T cell zones of cervical lymph nodes and palate tonsils [57]. Association studies of this small patient cohort indicated that increased reactivity to neuronal-derived antigens was correlated with smaller infarct size and better long-term outcome,

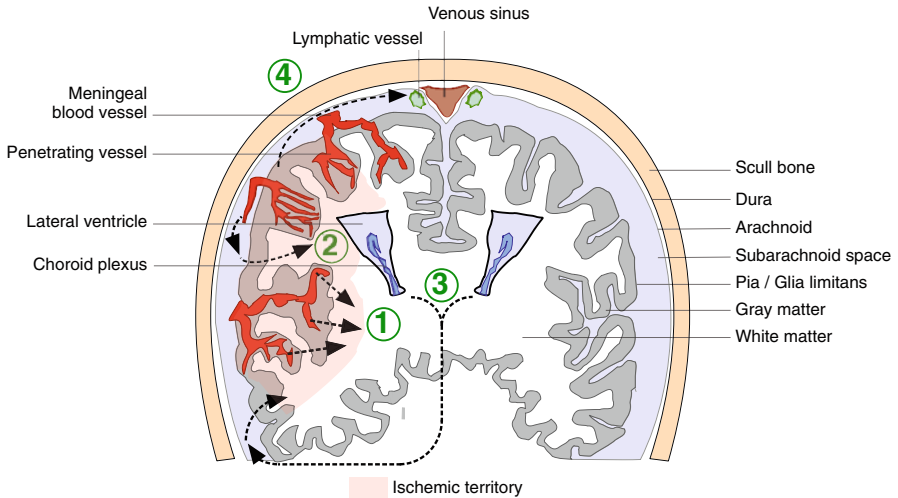


Fig. 1 Possible immune cell entry points after ischemic brain injury. (1) Circulating immune cells adhere to the activated endothelium within the ischemic territory and migrate into the ischemic tissue across the damaged BBB. (2) Blood leukocytes can extravasate through meningeal vessel to enter the subarachnoid space. From there they might infiltrate the brain parenchyma along chemo-tactic gradients produced within the ischemic territory by activated endothelial cells, glia, and neurons. Impairment of the surface glia limitans covering the ischemic tissue facilitates the exchange of such inflammatory mediators between the brain parenchyma and the subarachnoid space. (3) Immune cells can gain access to the brain by entering the ventricular space through the choroid plexus, where adhesion molecules are upregulated in response to ischemic brain injury. Once in the CSF they can enter the brain parenchyma by penetrating the epithelial cell layer of the ventricles or migrate from the ventricles into the subarachnoid space via the foramina of Luschka and of Magendie located in the fourth ventricle. (4) Antigenic macromolecules or activated antigen-presenting cells can exit the cranium through meningeal lymphatic vessel terminating at deep cervical lymph nodes

whereas greater reactivity to MBP was correlated with larger infarcts and worse outcome. This dichotomy might be an indication that the adaptive immune response to ischemic brain injury can be skewed towards reactive (Th1/Th17) or tolerogenic (Th2) phenotypes. This interpretation is supported by studies in rodent stroke models that indicate a beneficial effect of a Th2 immune response on stroke outcome [58, 59]. Future studies will have to address whether a tolerogenic immune response is linked to favorable stroke outcome in humans.

3 Immune Cell Entry Points

The forebrain with exception of the olfactory bulb is covered by a continuous layer of astrocyte cell bodies in rodents or by cytoplasmic processes of marginal astrocytes in primates known as the glia limitans [60]. The astrocytes are in close contact

with the basal lamina which is in intimate contact with the pia mater, the innermost leptomeningeal membrane. This structure forms a tight barrier to the subarachnoid space and limits the exchange of solutes and cells between the brain parenchyma and the cerebrospinal fluid (CSF). The glia limitans extends along penetrating blood vessels and together with pericytes, the basement membrane and interendothelial tight junctions form the BBB that renders the blood vessel of the cerebral vasculature impermeable to macromolecules and circulating cells. In the course of inflammation these barriers are altered and leukocytes penetrate into the CNS via multiple routes including (1) blood-to-parenchyma through a weakened BBB, (2) blood-to-CSF via the choroid plexus, (3) blood-to-CSF via leptomeningeal vessel, and (4) CSF-to-parenchyma across the glia limitans (Fig. 1).

Parenchymal entry—In the course of cerebral ischemia the BBB undergoes several changes that are reversible at first but become permanent at later time points of the ischemic injury. The first opening of the BBB is due to loss of endothelial tight junctions with concomitant transient downregulation or redistribution of junction proteins such as occludins, claudin, and zonula occludens-1. While the impairment of the BBB caused by endothelial tight-junction dysfunction occurs within hours after cerebral ischemia, a second opening of the BBB occurs in later phases of the ischemic insult and is characterized by loss of vascular cells, including endothelial cells and pericytes, proteolytic degradation of the basal membrane and retraction of astrocytic endfeet from the gliovascular unit [61]. It is during this phase that substantial infiltration of peripheral immune cells is observed in the ischemic territory. Within the reperfused territory leukocytes adhere to the endothelium of arterioles and post-capillary venules via endothelial upregulation of selectin-family adhesion molecules (P- and E-selectin) which results in a slow-down and rolling of intravascular leukocytes [62]. Firm attachment to endothelial cells is achieved through integrin-mediated interaction with endothelial-expressed intercellular adhesion molecule-1 (ICAM-1) and vascular cell adhesion molecule-1 (VCAM-1), among others. Interestingly, the interaction with the vasculature does not result in leukocyte extravasation and adherence is a relatively short-lived event that, depending on the experimental model, reverts back to pre-ischemia levels within hours after reperfusion [63, 64]. Leukocyte clogging at the time of reperfusion is thought to contribute to “no-reflow” phenomena [32], but leukocyte trapping in capillaries has not been consistently found in cerebral ischemia [63, 65]. Eventually leukocytes extravasate from the circulation into the brain parenchyma although this event is not synchronized with endothelial expression of adhesion molecules and generation of parenchymal chemokines, which in most transient focal cerebral ischemia models in rodents peak well before sustained numbers of parenchymal leukocytes are observed. The reasons for this remain elusive, but the existence of a double barrier—endothelial and glial basement membrane—could contribute to this effect [66]. As a result, blood-borne immune cells, specifically neutrophils, are often found in the ischemic territory forming cuffs that surround the vessel, without entering the neuropil. This has led to the assumption that the contribution of peripheral immune cells to ischemic damage might be marginal. However, “injury at a distance” due to release of neurotoxic factors from perivascular leukocytes has to be

considered as a contributing factor during the early stages of ischemia-reperfusion. Activated leukocytes express several proteases, which are released extracellularly and matrix metalloproteases (MMPs) have been consistently linked to ischemic brain injury. Increased levels of such proteases should aid the degradation of the extracellular matrix including the basement membranes and expose the leukocytes to parenchymal chemokine gradients that enhance leukocyte infiltration. Overall, although immune cell entry from the vasculature within the ischemic parenchyma is likely, hard evidence for this route of entry is still missing.

Choroid plexus—The choroid plexus is a highly vascularized structure that attaches to the ependymal cells lining the cerebral ventricles and separates the ventricles from the subarachnoid space. In contrast to the brain parenchymal and meningeal blood vessel, the endothelial cells of the choroid plexus are fenestrated allowing solutes and intravascular cells to cross the endothelial cell layer. Instead, the choroid epithelial cells are interconnected by continuous tight junctions similar to the ones found in endothelial cells of parenchymal blood vessels. The highly polarized epithelium forms villous structures, which increase its ventricular surface area, and is able to actively produce large quantities of CSF. The stroma of the choroid plexus contains a sizable number of tissue-resident macrophages and dendritic cells that express MHC class-II molecules and may present antigens to T cells entering the CSF [67]. In the healthy brain the choroid plexus is the entry site for patrolling lymphocytes—mostly CD4⁺ central memory T cells [68]. To facilitate leukocyte trafficking through the choroid plexus, the choroid plexus epithelium expresses constitutively ICAM-1 and VCAM-1, which together with mucosal vascular-addressing cell adhesion molecule-1 (MAdCAM-1), are upregulated under inflammatory conditions [69, 70]. The choroid plexus epithelial cells also constitutively express the chemokine CCL20 that acts on the CCR6 receptor on the surface of IL-17 secreting lymphocytes to promote their CSF entry [71]. Given that IL-17 $\gamma\delta$ T cells have been implicated in stroke pathophysiology, the selective recruitment of IL-17⁺ T cells across the choroid plexus might be relevant for the outcome of ischemic brain injury. CD73, an ecto-ATPase expressed on choroid plexus epithelial cells and lymphocytes, has been shown to mediate monocyte/macrophage migration across the choroid plexus in a model of spinal cord injury [72]. Deletion of CD73 exacerbated ischemic injury in mice undergoing focal ischemia even when wild-type leukocytes were engrafted in the CD73^{-/-} host after bone marrow ablation [73]. The results indicate that immune cells with an anti-inflammatory phenotype or immune cells contributing to tissue repair [72] may use this route of entry. Taken together, more studies are needed to address the likely role of the choroid plexus as an entry point for leukocytes after stroke.

Meninges—The brain and the spinal cord are enclosed by three membranes. The fibrous dura mater, the highly vascularized arachnoid, that with its trabecular structure spans the subarachnoid space filled with CSF, and the pia mater, which covers the brain parenchyma and is in close contact with astrocytes forming the glia limitans at the surface of the brain. The arachnoid is highly vascularized and cerebral vessels are embedded within the arachnoid trabeculae before they enter the brain parenchyma. Similarly to the choroid plexus, the meninges are populated by bone

marrow-derived perivascular myeloid cells that express the microglia/macrophage marker Iba1 and MHC class-II molecules while a subpopulation of meningeal macrophages expresses the dendritic cell marker CD11c [67]. Little is known about the physiological role of meningeal macrophages but given their proximity to blood vessels and their possible interaction with CSF lymphocytes, it is predicted that these cells play a role in the CNS immune surveillance. The other constitutive meningeal immune cell population identified in humans and rodents are mast cells primarily located in the dura mater [74]. Because mast cells contain granules with vasoactive molecules and proteases, they have been implicated in BBB disruption and in promoting neutrophil extravasation in the course of cerebral ischemia [74, 75]. Several studies have addressed the role of leptomeningeal vessel as a source of blood-borne immune cells after stroke. Consistent with a meningeal origin, neutrophils are found on the abluminal site of leptomeningeal vessel within hours after stroke in permanent and transient ischemia models in rodents. A strong association of neutrophils with leptomeningeal vessel has also been observed in tissue samples from human stroke victims [76]. Whether neutrophils that extravasated into the subarachnoid space go on to infiltrate the ischemic territory remains to be established, but the fact that accumulation in the meninges precedes the appearance of neutrophils in the brain parenchyma supports such a scenario [30, 76, 77].

4 Alterations of the Peripheral Immune System After Stroke

While peripheral immune cells contribute to ischemic brain injury, the ischemic brain in turn has a profound effect on the composition and behavior of the peripheral immune system. This effect is mediated by the systemic release of danger-associated molecular pattern (DAMP) molecules from the ischemic territory, humoral signals generated by hypothalamic stress response centers, and neural signals of the autonomous nervous system. The response is characterized by an early state of hyperinflammation, followed by a phase of immunosuppression with increased susceptibility to infection. Parts of this response are not specific to ischemic brain injury and are also observed in other conditions associated with severe tissue injury such as burns and traumatic organ damage [78, 79]. Of note, in a mouse model of transient focal ischemia, anesthesia and surgical trauma itself led to early changes in the composition of blood immune cell populations independent of actual middle cerebral artery occlusion [80]. Therefore, observed changes in the peripheral immune response after experimental stroke have to be interpreted in the context of the occlusion model, anesthesia, surgical manipulation, and time points of analysis. Moreover, many of the changes to the peripheral immune system discussed below are only observed in stroke models that result in large ischemic injuries (>30–40% of the hemispheric volume), a correlation also observed in human stroke [81]. Accordingly, distal occlusion of the middle cerebral artery in rodents, which produces infarcts <20% of the hemispheric volume, does not elicit a peripheral immune response and does not cause immunosuppression. In addition, stroke-induced changes in the

peripheral immune system show lateralization [82, 83] and the net immunomodulatory autonomic output after ischemia might be dependent on brain structures damaged [84]. Consequently, electrolytic lesion experiments showed that lesions in the hypothalamus were immunosuppressive while similar lesions in the hippocampus and amygdala resulted in immunostimulation [85, 86].

Hyperinflammation—the early response to ischemic brain injury. In experimental stroke, the earliest peripheral immune response to ischemic brain injury is characterized by elevated serum cytokine levels (IL-6, interferon- γ , CXCL1) and increased production of inflammatory mediators in circulating and splenic immune cells (TNF, IL-6, IL-2, CCL2, and CXCL2) within hours after ischemia [87, 88]. The response is generally transient and most parameters return to baseline levels 24 h after stroke. Comparable changes can be observed in human stroke patients. Tumor necrosis factor (TNF) and IL-6 are increased in patients at stroke onset (<24 h) [89] and IL-6 serum levels are positively correlated with stroke severity and unfavorable outcomes [90]. However, this early activation of the immune system is superseded by a state of systemic immunodepression that predisposes to post-stroke infections [91, 92]. Accordingly, complications from pulmonary or urinary tract infections have been observed in ~20 % of stroke patients [93].

Stroke-induced immunodeficiency syndrome—Early studies on the immune status of stroke patients found prolonged peripheral lymphopenia and reduced T cell responsiveness [82, 94]. Ischemic brain injury leads to sustained decrease in blood and splenic B, T, and NK cells in mice undergoing transient middle cerebral artery occlusion [92, 95, 96]. These changes are observed 12 h after ischemia and persist for several weeks. Importantly, the immunosuppression leads to increased susceptibility to nosocomial bacterial infection and to higher mortality rate [92]. The decrease in lymphocytes was correlated with increased splenocyte apoptosis, spleen atrophy, and Treg expansion [92, 95]. The sympathetic nervous system was fundamentally involved in this response. Treatment of animals with the β -adrenergic receptor antagonist propranolol or chemical ablation of sympathetic noradrenergic terminals by 6-hydroxydopamine was sufficient to lower bacteremia and bacterial colonization of the lungs and increased significantly survival rates along with preservation of splenic and blood lymphocyte populations [92]. Interestingly, the glucocorticoid receptor antagonist RU-486 reversed the lymphopenia in the blood, but not in the spleen, and did not impact bacterial colonization or survival rates after stroke. Whether the splenic response to cerebral ischemia is induced by similar mechanisms in all stroke models and in humans remains to be determined. Studies in rats concluded that the loss of splenic lymphocytes was not due to increased apoptosis but due to activation of α -adrenergic receptors on trabecular and capsular smooth muscle cells that leads to spleen contraction and expels immune cells into circulation [97] (Fig. 2). The study found that although spleen size was decreased 1 day after stroke, spleen volume was restored 3 days thereafter arguing against long-lasting effects of cerebral ischemia on spleen physiology. As in rats, splenic size loss in humans is transient and there is a tendency of increased splenic volume 4 days after stroke [98]. The immunosuppressive effects of ischemic brain injury are not limited to the spleen. In the bone marrow, tyrosine hydroxylase and

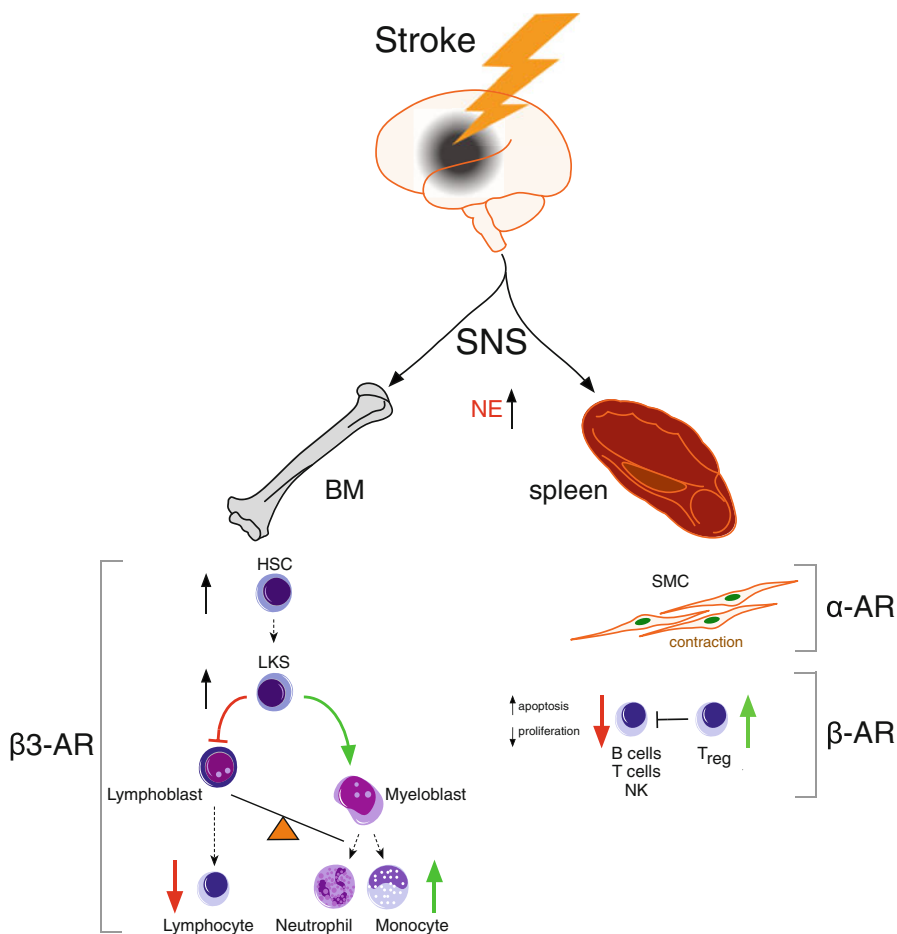


Fig. 2 Stroke-induced immunodeficiency syndrome. Stroke increases activity of the sympathetic nervous system (SNS) resulting in the release of norepinephrine (NE) from sympathetic terminals in bone marrow (BM) and spleen. In the spleen activation of α -adrenergic receptors (α -AR) on capsular and trabecular smooth muscle cells (SMC) results in contraction of the splenic pulp, decrease in organ size, and reduction in cellular density. Furthermore, stimulation of β -adrenergic receptors (β -AR) on splenic lymphocytes results in Treg expansion and B cell, T cell, and NK cell apoptosis. In the BM, NE activates β -adrenergic receptors (β -AR) located on mesenchymal stromal cells, which trigger the downregulation of homeostatic factors including CXCL12 (SDF-1), resulting in increased hematopoietic stem cell (HSC) and Lin⁻/c-Kit⁺/Sca-1⁺ (LKS) stem cell proliferation. LKS stem cells are located immediately before the commitment to lymphoid or myeloid lineages. Adrenergic signaling skews LKS cells to the myeloid lineage resulting in increased neutrophil and monocyte production while decreasing the generation of lymphocytes

norepinephrine (NE) levels increase 1 day after transient middle cerebral artery occlusion in mice [99]. This triggers a response in mesenchymal stromal cells, probably through activation of β -adrenergic receptors (β -AR), resulting in the reduction of homeostatic and cell retention factors such as interleukin-7, C-X-C

motif chemokine 12 (stromal cell-derived factor 1), VCAM-1, stem cell factor, and angiopoietin-1. Downregulation of these factors increases hematopoietic stem cell (HSC) proliferation (Fig. 2). This proliferative response, however, does not profit all arms of leukocyte lineages equally [99]. The hematopoietic system becomes skewed towards the myeloid lineage. This switch is accomplished by increased expression of transcription factors associated with myeloid lineage progression such as the ETS-domain transcription factor PU.1 and CCAAT/enhancer-binding protein β (C/EBP β). Collectively the data suggest that the brain is a strong regulator of the peripheral immune system by regulating development and homeostasis of splenic and BM immune cell populations and that increased sympathetic output after stroke is the main efferent branch responsible for these effects.

5 Epilog

Bidirectional interactions between the injured brain and the peripheral immune system are not only important for the development of the ischemic injury but also strongly affect the immune status of the organism as a whole. Whereas the initial immune response triggered by stroke is largely pro-inflammatory, stroke-induced immunodeficiency syndrome, while being desirable for limiting the deleterious effects of post-ischemic inflammation, poses a severe risk for bacterial infections and unfavorable outcome. Because inflammation and peripheral immune cells that infiltrate the ischemic brain might also be involved in repair processes, it will be of importance to determine how the functionality of these systems is affected by stroke. In conclusion, immunomodulation by enhancing the activity of reparatory neutrophils and macrophages or by skewing the adaptive immune system to a “tolerized” state might constitute a more promising approach than indiscriminate anti-inflammatory therapies to limit the deleterious effects of post-stroke inflammation.

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The Role of Spleen-Derived Immune Cells in Ischemic Brain Injury

Heng Zhao

1 Introduction

Despite numerous extensive studies on stroke therapy in the past decades, few have successfully translated to the clinic and many limitations and potential problems remain. One major limitation is that the brain has been studied in isolation, as if tissue injury in the brain is unrelated to what occurs in peripheral organs. We now realize that stroke not only injures the brain, but it also affects multiple organs, including the skeletal muscle [1–3], heart [4], liver [5], lung [6], and peripheral immune system, including the spleen [7–9]. As a result, it has been speculated that modulating peripheral organs may protect against brain injury. And in fact, we have reported that repetitive ischemia performed in the hind limbs of rats reduces brain infarction after focal ischemia—a phenomenon defined as remote preconditioning [10]. Other important facts are that the spleen interacts with the ischemic brain, as stroke results in spleen atrophy, while spleen removal (splenectomy) performed before stroke onset robustly reduces post-stroke brain injury [11, 12]. Therefore, the spleen may be an alternative avenue or target for stroke treatment. In this Chapter, we discuss how the spleen and its immune cells contribute to brain injury induced by stroke.

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2 The Structure and Function of the Spleen

The spleen is a unique organ belonging to the super immune system, which consists of many lymphatic organs and multiple immune cells [13–15]. The immune system includes primary and secondary lymphoid organs. The primary lymphoid organs mainly include the bone marrow and thymus, which are the sites of lymphocyte development and maturation. The secondary lymphoid tissues include the spleen, lymph nodes, and mucosa- and skin-associated lymphoid tissues, within which mature lymphocytes exert their immune response, such as antigen recognition and activation, clonal selection and proliferation, as well as phagocytosis [16–19].

The spleen, as the largest secondary lymphoid organ, plays important roles in the immune response. It is composed of red and white pulp with distinct morphologies and functions [16–19]. Blood circulation in the spleen is supplied by a single artery called the splenic artery, which branches into a network of smaller arteries that travel through the spleen, forming arterioles, and connecting with sinuses [14]. The small arterioles travel to and end in a venous sinusoidal system in the red pulp, which serves as a blood filter by trapping old or damaged red blood cells that are phagocytosed by red pulp macrophages. Another major function of the red pulp is for iron recycling [14]. White pulp is a lymphoid tissue surrounding the small branches of arterials, and is composed of T cell, B cell (follicle), and marginal zones, where the immune response occurs [16–19].

Taken together, the primary function of the spleen is to monitor circulating blood, filter blood by trapping damaged or old red blood cells, and initiate the immune response against pathogens found in the circulation. Both innate and adaptive immune responses can be easily mounted in the spleen, as the unique pattern of blood supply and structure or organization of the immune compartments allow various immune responses to occur. However, the spleen also releases lymphocytes into the blood circulation in response to stress and inflammation, which travel to other organs, including the brain. Therefore, the spleen is also an important organ for modulating brain functions and neuroinflammation.

3 The Spleen Is Involved in Brain Infarction Induced by Stroke

It is well known that bidirectional effects occur between the injured brain and peripheral immune system after stroke, and the spleen has a unique role in such bidirectional effects. On one hand, stroke causes peripheral immune suppression [9], which results in post-stroke infection and mortality [20–23]. The immune suppression is manifested by a reduction in lymphocyte activation and spleen atrophy after stroke [9]. As a major secondary lymphocyte organ, the spleen functions to maintain lymphocyte and erythrocyte populations. Splenic atrophy is a typical

marker of the immune suppression induced by stroke [9]. On the other hand, the peripheral immune system increases local brain inflammation via recruitment and infiltration of circulating neutrophils [24–27], monocytes/macrophages [28, 29], and T cells [30–35], thus exacerbating ischemic injury.

The most direct evidence that spleen is involved in brain injury is shown in splenectomy studies. Ajmo and colleagues showed in their first study that splenectomy performed 2 weeks before stroke reduced infarction by 80% [11, 12]. This study is supported by Li et al., who found that splenectomy performed immediately before traumatic brain injury decreases animal mortality and improves cognitive function in rats [36, 37]. In addition, Ostrowski et al. reported that acute splenic irradiation, which results in lymphocyte death in the spleen, reduces brain injury in rat focal cerebral ischemia [38]. They found that the therapeutic time window extends to 3–4 h after stroke onset, suggesting that spleen modulation post-stroke can provide neuroprotection [38]. More recently, Seifert et al. and Zhang et al. also confirmed the neuroprotective effect of splenectomy against stroke [39–41]. In contrast, Kim et al. failed to see any neuroprotective effect of splenectomy immediately before stroke on reducing infarction, although splenectomy did reduce the accumulation of inflammatory macrophages in the ischemic brain [42]. The explanation for this controversial result is unknown. Although the authors argue that the timing of splenectomy immediately before stroke is a factor [42], others have shown that spleen irradiation performed even 3–4 h after stroke can generate protection [38].

In addition to the protective effects of splenectomy against stroke, evidence exists to suggest that the spleen modulates brain injury induced by stroke. First, there is a correlation between reductions in spleen size after stroke and the extent of infarction in animal studies, as well as between spleen size and brain injury in stroke patients [43]. Second, a number of studies suggest that neuroprotectants attenuate the reduction in spleen size and splenocyte numbers. As we reported, moderate hypothermia attenuates the reduction in spleen size and lymphocyte numbers, and this correlates with the robust protective effects of hypothermia on brain infarction [44]. In addition, cord blood injection inhibited brain injury, which was associated with less reduced spleen sizes [45]. Taken together, the spleen contributes to brain injury induced by stroke.

Nevertheless, the underlying mechanisms involved in the cross talk between the spleen and post-stroke ischemic brain, and how immune cells derived from the spleen modulate ischemic brain injury are unclear. In the next two sections I will identify potential mechanisms based on previous studies about spleen function, as well as the possible roles of splenic immune cells in neuroinflammation.

4 Communication Between the Spleen and Brain

Bidirectional communication between the spleen and brain occurs via the immune system and central nervous system (CNS). The peripheral lymphoid organs, including the spleen, are hardwired to the autonomous nervous system [46]. In addition,

lymphocytes in the spleen and other peripheral lymphoid organs express receptors for neurotransmitters released from the nervous system [23]. Other sensors, such as cytokine receptors in the CNS, relay information from the CNS to peripheral organs. This information is mainly processed in the brain by the frontal premotor cortex, hypothalamus, pituitary, and brain stem, which forms three major pathways: the hypothalamus–pituitary–adrenal (HPA) axis, the sympathetic nervous system (SNS), and the parasympathetic nervous system (PNS) [23]. These three major systems bridge the peripheral lymphoid organs, including the spleen, with the brain.

The SNS innervates almost all lymphoid organs, including the bone marrow, thymus, lymphoid nodes, and spleen [23]. In addition, almost all leukocytes express adrenergic receptors, which react with the neurotransmitter catecholamine released from the SNS [47–51]. Activation of the SNS due to inflammatory stimulation in the brain results in a large release of catecholamine, which influences immune functions in the peripheral lymphoid organs [23]. It is known that catecholamine released from the SNS results in a transient and rapid release of leukocytes from the lymphoid organs, including the spleen [52, 53]. This results in a rapid increase of leukocytes into the circulation, which migrate to the brain and exacerbate the brain's inflammatory response [52]. Indeed, Ajmo and colleagues demonstrated that treatment with either prazosin, an α_1 receptor blocker, or carvedilol, a pan adrenergic receptor blocker, prevented the reduction in spleen size, and carvedilol significantly reduced infarct volume, suggesting that SNS-mediated catecholamines regulate the splenic response to stroke through the activation of adrenergic receptors [12].

In addition to the SNS, PNS and HPA are also involved in the crosstalk between the CNS and peripheral lymphoid organs [23]. PNS activity modulated by the release of acetylcholine promotes the anti-inflammatory response, and reduces cytokine production, such as IL-1 β and TNF α , in the peripheral organs [54]. In addition, HPA activity leads to the release of glucocorticoids, which are also anti-inflammatory and immune-suppressive [55, 56]. Nevertheless, whether or not PNS and HPA activity results in leukocyte release from the spleen is not clear.

Although it is clear that the spleen contributes to brain injury induced by stroke, and communication between the spleen and CNS is modulated by the three pathways previously discussed, the role of the SNS, PNS, and HPA in the worsening effects of spleen on brain injury has not been well studied. It is likely that stroke results in neuroinflammation in the ischemic brain, which produces various inflammatory cytokines, such as IL-1 β , IL-6, and TNF α [23]. These released cytokines may secrete into the CSF or diffuse within the ischemic brain, thus stimulating the SNS and resulting in the release of leukocytes from the spleen. Another possibility is that inflammatory cytokines are released from the ischemic brain into the circulation, and these cytokines act directly on the spleen via the circulating blood [23]. The released leukocytes could then infiltrate into the ischemic brain, and result in a larger infarction.

5 Neuroinflammation and Spleen Immune-Cell Trafficking to the Brain After Stroke

Inflammation plays several critical roles in stroke and stroke-induced brain injury. For example, systemic infection-induced inflammation correlates with stroke [57–63], and surgery-induced inflammation also increases the risk of stroke [64–66]. Second, after ischemic stroke there is an immediate onset of neuroinflammation, which involves multiple facets, including brain damage, tissue clearance, and functional recovery [67]. The spleen may be involved in these aforementioned neuroinflammation and stroke, although there is no clear evidence indicating how spleen immune cells modulate stroke, and the role of splenocytes on post-stroke brain recovery has not been studied.

Upon stroke, cerebral blood vessel(s) is occluded and become hypoxic, causing reduced nitric oxide and further constriction of blood vessels, which increases production of reactive oxygen species and platelet activity [67]. These intravascular changes cause the adhesion of leukocytes to the blood vessel walls, and increases in blood–brain barrier permeability and leukocyte infiltrate to the ischemic brain. Therefore, in addition to resident microglia in the brain, which are activated and transformed into macrophages, other leukocytes from circulating blood, including neutrophils, monocytes/macrophages, B cells, T cells, NK, and NKT cells, play important roles in neuroinflammation and brain injury induced by stroke [67].

It is known that T cells modulate brain injury [31, 32, 67]. T cells include the subsets of CD4 T cells, CD8 T cells, and $\gamma\delta$ T cells. Functionally, activated CD4 T cells during immune responses can be further differentiated into Th1, Th2, and Th17 subsets. It has been reported that the lack of total T cells, or a subset of CD4, CD8, or $\gamma\delta$ T cells, results in a smaller infarction compared with immune intact animals [68]. In addition, the lack of Th1 cells results in neuroprotection, while the lack of Th2 cells exacerbates brain injury [31]. We also observed that the lack of Th17 was neuroprotective against stroke (unpublished observation). Th17 cells function by secreting IL-17. A previous study has shown that neutralization of IL-17 attenuates neuroinflammation [69]. Among these cell types, CD4+CD25+ regulatory T cells (Treg) have been the most widely studied, with results showing that Treg cells have anti-inflammatory functions and their activity attenuates delayed brain injury induced by stroke [68, 70, 71]. B cells have been less studied in stroke. The first study on B cells suggested that B cells do not have an effect on brain injury induced by stroke [33]. More recently, regulatory B cells were found to inhibit brain injury [72].

Monocytes/macrophages are also important for brain injury induced by stroke. Macrophages can be derived from both brain resident microglia and peripheral blood monocytes. The latter can be released from bone marrow and other secondary lymphoid organs, including the spleen. Functionally, macrophages are polarized into M1 and M2 phenotypes, with pro- and anti-inflammatory roles, respectively [73, 74].

It has been reported that the M1 phenotype is detrimental, while M2 phenotype is beneficial, in ischemic stroke [75–77]. In particular, M2 macrophages may play a critical role in promoting brain repair and recovery after stroke [75].

As we discussed, the spleen contains most of the cell types in the immune system, including T cells, B cells, monocytes, and macrophages, and stroke results in the contraction of the spleen, which leads to the release of leukocytes into the blood circulation. The released leukocytes then migrate to the brain modulating acute brain injury. Nevertheless, as discussed, leukocytes recruited into the ischemic brain include those released not only from the spleen and other secondary lymphoid nodes, but also from the primary lymphoid organs, such as bone marrow. It still remains unclear how much the spleen contributes to or modulates brain injury induced by stroke among these various lymphoid organs.

Although the spleen contributes to brain injury induced by stroke, few studies have examined the exact roles of the individual cell types that make up splenocytes or the underlying mechanisms of spleen-induced brain injury. Nevertheless, several lines of evidence from previous studies indeed suggest that the involvement of spleen in ischemic brain injury is associated with splenocytes and neuroinflammation. First, one recent study suggests that spleen contraction induced by stroke correlates with reduced cell numbers of monocytes in the spleen, including pro-inflammatory Ly6C^{hi} and anti-inflammatory Ly6C^{low} monocytes [42]. The study further showed that the deployment of these monocyte subsets coincided with respective increases in the ischemic brain [42]. In contrast, splenectomy reduced leukocyte infiltrations into the ischemic brain [42], suggesting that monocytes are released from the spleen and migrate to the ischemic brain. Second, there is additional direct evidence that splenocytes infiltrate into the ischemic brain. In one study, splenocytes were labeled with CFSE, and it was found that CFSE-positive cells were released into the blood from the spleen, including T cells, neutrophils, and monocytes. The presence of CFSE-positive monocytes and NK cells in the cerebral blood vessels of the ischemic brain [39] suggests that splenocytes migrate into the ischemic brain. Third, splenocyte infiltrations are associated with the expression of inflammatory factors. Splenectomy results in reductions in T cells, neutrophils, and macrophages in the ischemic brain [41], and this is associated with decreases in pro-inflammatory cytokines, such as IL-1 β and TNF α , and with increases in anti-inflammatory cytokines, including IL-10, in the brain [41]. Another study suggests that the protective effect of splenectomy is associated with IFN γ , as IFN γ was found to be increased in the spleen in early stroke followed by increases in the brain [40]. Fourth, the protective effects of neuroprotectants are linked to spleen functions. For instance, agmatine treatment reduced the contraction of white pulp, and inhibited the accumulation of CD11b macrophages and Treg cells in the spleen [78]. In addition, MFX treatment attenuated Ly6C expression in pro-inflammatory macrophage subsets and CCR2 expression in the spleen tissues [79]. Furthermore, the injection of cord blood reversed the reduction in spleen size and concomitant reductions in CD8 T cells after stroke, as well as increased IL-10 while inhibiting IFN-gamma [45].

6 Problems and Future Research Directions

It is well established that the spleen contributes to acute brain injury after stroke, but there are many questions raised from previous studies. First, the underlying mechanisms of the spleen's contribution to brain injury are not understood. Future studies can address the following questions: How are the nerve pathways, including SNS, PNS, and HPA involved in the interaction between the spleen and brain after stroke? How are immune cells released from the spleen? Which splenocytes migrate to the ischemic brain, and which cell types play the most important roles? Second, the spleen is only one of the immune organs, and how much it contributes to ischemic brain injury is unknown. Therefore, one may ask, what is the relative contribution of the spleen, in comparison to the other lymphoid organs, to brain injury? Third, previous studies have focused on studying the relationship between acute brain injury and the spleen; whether the spleen plays an important role in brain repair and recovery has not been studied. More scientific issues include: Does the spleen play a critical role in brain recovery after stroke? If so, what is the underlying mechanism?

7 Conclusions

Stroke results in the release of inflammatory factors that stimulate nerve tissue sensors inside and outside of the brain as well as sensors expressed on peripheral organs, such as the spleen. The contraction of the spleen after stroke leads to the release of splenocytes, which migrate to the ischemic brain and exacerbate brain injury. Taken together, the spleen plays an important role in acute brain injury induced by stroke, but the underlying mechanisms require more study, and the importance of the spleen to brain repair and recovery is not clear.

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Regulatory T Cells in Ischemic Brain Injury

Arthur Liesz

1 Introduction

Post-stroke neuroinflammation has come into the focus of current preclinical stroke research [1]. Among the pathophysiological mechanisms of microglial activation, brain leukocyte invasion and secretion of pro-inflammatory factors, lymphocytes have been uncovered as the key leukocyte subpopulation determining the neuroinflammatory outcome. Several studies have shown that pro-inflammatory lymphocytes such as TH1, TH17, and $\gamma\delta$ -T cells worsen stroke outcome and that blocking their brain invasion is neuroprotective [2–5]. Contrary to pro-inflammatory lymphocytes, regulatory T cells have been characterized in primary inflammatory diseases as disease-limiting protective cells [6]. Finding this key role of regulatory T cells in other T cell-driven pathologies has initiated productive research efforts on the role of regulatory T cells also in ischemic brain injury over the past years. Due to the complex function of regulatory cells in immune homeostasis and disease as well as partially divergent findings using different stroke models, uncertainty has emerged about the pathophysiological function of regulatory lymphocytes in stroke.

2 Regulatory T Cells in Post-Stroke Neuroinflammation

The immune system has evolved several regulatory mechanisms to inhibit an overshooting immune reaction to tissue damage including cell depletion, anergy, and unresponsiveness to self-antigens. The presence of regulatory T cells (Treg)

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suppressing actively (self-reactive) immunity is one of the key mechanisms of preserving immune homeostasis and limiting inflammatory collateral damage [7]. For example, depletion of CD25+ CD4+ Treg cells naturally arising in the immune system induces autoimmune diseases and reconstitution of this cell population prevents disease development [8]. Lack of Treg has been shown to be a primary cause of autoimmune diseases in humans [9]. In addition to sustaining self-tolerance, Treg are also intricate in suppressive control of a broad spectrum of immune responses including those against autologous tumor cells [10], allergens [11] and organ transplantation [12]. Although Treg might be simplistically defined as immunosuppressive T cells, several phenotypically and functionally distinct Treg subpopulations have been defined, such as induced Treg populations, Tr1, TH3, and various others [13]. Yet, the best investigated population of regulatory lymphocytes are CD4+ CD25+ Foxp3+ naturally occurring regulatory T cells, which are physiologically produced in the thymus as a mature and functional cell population [14]. While it is still unknown at which specific site Treg act as immunomodulators after stroke, their kinetics and magnitude of brain infiltration have been characterized in several studies using models of experimental middle cerebral artery occlusion (MCAO). One of the first studies systematically analyzing brain leukocyte invasion after transient brain ischemia (filament-induced MCAO resulting in large hemispheric lesions) by flow cytometry was performed by Gelderblom and colleagues [15]. In this study only a very low number of CD25+ Foxp3+ Treg cells at a frequency of less than 5 % of all CD4+ T cells was observed within the first week after transient ischemia. In contrast, using a permanent MCA occlusion model we detected substantial T cell and Treg counts in the ischemic hemisphere (Fig. 1) with Foxp3+ Treg cells constituting approx. 20 % of all CD4+ cells [16]. Correspondingly, it was previously shown that distal, permanent occlusion induces a significantly stronger neuroinflammatory reaction with manifold higher T cell counts in the ischemic hemisphere than in proximal, transient occlusion models [17]. Also in contrast to the initial study by Gelderblom et al., a more recent work by Stubbe et al. has investigated cerebral Treg counts also at later stages after large hemispheric lesions and detected only negligible Foxp3+ cells in ischemic hemispheres early after MCAO

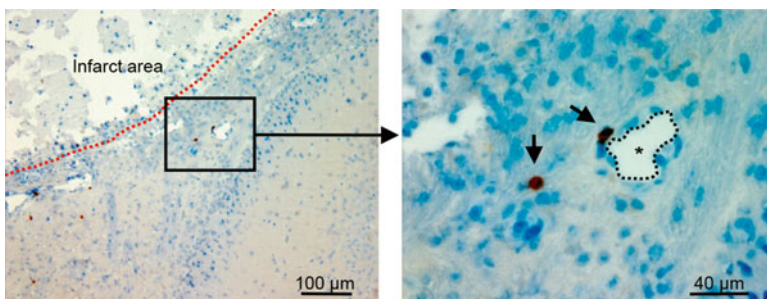


Fig. 1 Regulatory T cells invade the ischemic brain. Immunohistochemical sections showing Treg cells predominantly in the peri-infarct area. Arrows indicate Foxp3+ Treg cells; the asterisk marks a blood vessel (Liesz et al., *Nature Medicine*, 2009)

but substantial brain Treg invasion at 14days and 30days after MCAO. Additionally, the percentage of Treg cells within the T_{helper} cell population was increased in brains after stroke compared to peripheral lymphatic organs [18].

Apart from the evidence of variable Treg cell invasion depending of the used stroke model, it is feasible to assume that Treg cells might execute their function on target cells in the peripheral immune system. The majority of studies has detected the effector function of Treg cells in depletion or therapeutic paradigms (see below) already within the first days after brain injury before a considerable amount of Treg have even invaded the brain. Likewise, delayed deletion of Treg cells by antibodies or in inducible Foxp3-KO mice at 3days after MCAO was not effective in altering stroke outcome [18, 19]. Hence, although the definite site of action for Treg after stroke is still unsolved, currently available data suggests that Treg fulfill their immunomodulatory function within the first 3days after stroke and most likely outside the CNS. Potential peripheral target sites of Treg cells might include suppression of peripheral effector T cell activation, inhibition of autoantigen-specific clonal expansion or priming of transendothelial effector T cell-migration via currently unknown mechanisms.

3 Treg Depletion in Experimental Stroke

Depletion of the regulatory T cell population has been used as an experimental paradigm to investigate the functional role of Treg in brain ischemia in several publications (Table 1). Methodologically, two different approaches have been used for Treg depletion, antibody-mediated cell lysis using CD25-specific antibodies or the use of transgenic mice with a diphtheria toxin receptor (DTR) transgene under the control of the Foxp3 promotor for inducible Treg depletion. Using Treg-depletion paradigms about half of the experiments performed revealed an increased infarct volume while the other half did not detect any effect on stroke outcome and one study even observed a reduction of infarct size in Treg-deficient mice (Table 1). This discrepancy of Treg depletion on stroke outcome led to an unsolved debate on their role and denomination of Treg as a “double-edged sword” in acute brain injury [20–22]. This discrepancy could not be attributed to the used depletion paradigm, since genetic deletion models can be found on both sides of the “efficacy spectrum”. Notably, the three publications using inducible Foxp3-KO mice used in each case a different mouse strain: Ren et al. [23] used the original Foxp3^{DTR} mouse [24], Liesz et al. [19] the Foxp3.LuciDTR.4 mouse line [25], and Kleinschnitz et al. [26] the Dereg mouse [24]. All three of these DTR-transgene-mediated Foxp3-deletion models are using different transgene constructs, are variable in Treg-depletion efficacy and the used DT treatment protocol. Nevertheless, a systematic bias by the used depletion paradigm did not become evident for the authors. In contrast, it is apparent that the utilized stroke models, and more precisely the resulting volume of the ischemic lesion, do indeed predict a deterioration of outcome versus no effect or improved lesion size after Treg depletion (Table 1). In this line, both studies from our group

Table 1 Summary of studies investigating Treg depletion paradigms [31]

Study	Model	Lesion (% hemisphere)	Latest endpoint	Depletion model	Depletion efficacy	Outcome on infarct vol.
Liesz et al. (2009) [16]	Distal, permanent	≈10 %	7days	1. Anti-CD25 2. Adoptive cell transfer	90 % n.a.	Increased
Liesz et al. (2013) [19]	Distal, permanent	≈10 %	7days	Foxp3-KO	85 %	Increased
	30 min transient	≈10 %	7days	Anti-CD25	90 %	Increased
Xie et al. [27]	Rats: 90 min transient	≈40 %	3days	Anti-CD25	65 %	Increased
Liesz et al. (2009) [16]	90 min transient	≈55 %	7days	Anti-CD25	90 %	No effect
Ren et al. [23]	60 min transient	≈55 %	3days	Foxp3-KO	90 %	No effect
Li et al. (2013) [28]	60 min transient	≈50 %	3days	Anti-CD25	90 %	No effect
Stubbe et al. [63] ^a	30 min transient	≈50 %	27days	Anti-CD25	75 %	No effect
Liesz et al. (2013) [19]	60 min transient	≈50 %	3days	Foxp3-KO	85 %	No effect
Kleinschmitz et al. [26]	60 min transient	≈55 %	7days	Foxp3-KO	90 %	Reduced

^aDepletion 3days after MCAO induction; in all other studies: pretreatment or acute depletion

Table 2 Effect of Treg depletion on cerebral neuroinflammation

Study	Outcome on infarct volume	Leukocyte invasion	Cerebral cytokines	Microglia
Liesz et al. (2009)	Increased	↑	↑	(↑)
Liesz et al. (2013)	Increased	NA	↑	NA
Xie et al.	Increased	↑	↑	No effect
Liesz et al. (2009)	No effect	NA	NA	NA
Ren et al.	No effect	NA	NA	NA
Li et al. (2013)	No effect	NA	NA	NA
Stubbe et al.	No effect	No effect	No effect	No effect
Liesz et al. (2013)	No effect	NA	NA	NA
Kleinschnitz et al.	Reduced	NA	NA	NA

have detected an increase of stroke volume after Treg depletion only in small, permanent ischemia lesion models but not after extensive infarction induced by transient MCAO [16, 19]. A recent study by Xie et al. has also detected an increase of infarct volume after antibody-mediated Treg depletion in a rat model of moderate brain ischemia with rapamycin pretreatment [27]. In contrast, all studies investigating effects of Treg depletion without detection of an effect on stroke outcome [16, 18, 19, 23, 28] or even an increase of lesion volume [26] were performed in transient mechanical occlusion models with extensive brain lesions.

The immunological effects of Treg depletion on the neuroinflammatory outcome after stroke have unfortunately only been investigated in a small fraction of the studies (Table 2). All three studies detecting an exacerbation of stroke outcome after Treg depletion also measured an associated increase in neuroinflammatory biomarkers [16, 19, 27]. The most robust findings were an increase in leukocyte brain invasion, particularly of lymphocyte subpopulations, and an increase in pro-inflammatory cytokine secretion such as TNF- α , IL-1 β , IL-12, and IFN- γ . Among the studies using models of extensive brain injury with no effect of Treg depletion on stroke outcome, only one study [18] has investigated neuroimmunological readouts. Correspondingly to the missing effect on lesion volume, also inflammatory markers were not altered by the antibody-mediated Treg depletion in this study.

4 Treg-Therapies for Ischemic Stroke

Despite the obvious and still unresolved discrepancies arising from studies investigating effects of Treg depletion, a rapidly increasing number of reports have analyzed different strategies to increase the Treg number and/or function for experimental stroke therapy (Table 3). Out of 15 independent experiments reported in 13 studies, 13 have detected an improvement of stroke outcome while two have demonstrated an increase in infarct volume when using two independent Treg-targeted therapies. This discrepancy could not be attributed to the used

Table 3 Studies investigating Treg-targeted therapeutic approaches (by treatment and outcome)

Study	Model	Lesion (% hemisphere)	Latest endpoint	Treatment paradigm	Outcome on infarct volume
Chen et al. [64]	Rats: distal, permanent	≈20 %	48 h	Mucosal immunization	Reduced
Gee et al. [65]	Rats: 3 h transient	Infarct vol. NA ^a	28days	Mucosal immunization	Reduced (behavior)
Ishibashi et al. [41]	Rats: distal, permanent	≈20 %	28days	Mucosal immunization	Reduced
Li et al. (2013) [28]	Rats: 2 h transient	≈40 %	3days	Adoptive transfer	Reduced
Li et al. (2013) [46]	Mouse: 60 min transient	≈35 %	3days	Adoptive transfer	Reduced
Li et al. (2014) [66]	Mouse: 60 min transient	≈35 %	3days	Adoptive transfer	Reduced
Brea et al. [67]	Rat: transient	≈30 %	28days	Adoptive transfer	Reduced
Liesz et al. (2013) [19]	Mouse: distal, permanent	≈10 %	7days	HDACi	Reduced
Brea et al. [67]	Rat: transient	≈30 %	10days	CD28SA	Reduced
Na et al. [29]	Mouse: distal, permanent	≈10 %	7days	CD28SA	Reduced
Xie et al. [27]	Mouse: 60 min transient	≈50 %	3days	CD28SA	Reduced
Kleinschmitz et al. [26]	Rats: 90 min transient	≈45 %	28days	mTOR Inh.	Reduced
Schuhmann et al. [30]	Mouse: 60 min transient	≈55 %	3days	Adoptive transfer	Increased
	Mouse: 60 min transient	≈40 %	3days	CD28SA	Increased

^aThis study analyzed the behavioral deficits as the primary outcome and did not report infarct volume

therapeutic paradigm since both studies showing an exacerbation of lesion volume (adoptive Treg transfer and CD28SA) have been respectively tested in three or more other experiments that have been showing an improved outcome (Table 3). Again, it seems that the lesion size induced by the used stroke model might be a predictor of the neuroprotective effect of Tregs: out of the 13 experiments detecting a benefit of Treg-therapy, only two have reported infarct volumes in the control group of more than 40 % of the hemisphere, while the other studies deployed models inducing small to moderate-sized lesions. Yet, particularly two studies [29, 30] using both the CD28SA treatment in transient ischemia models with similar lesion size have reported contradicting results, indicating the presence of other confounding factors independent of lesion volume. We have performed in a recent review a Meta-analysis in order to assess an approximation of the efficacy of Treg-targeted therapeutic approaches in experimental stroke models [31]. For this, all studies having investigated interventions with the aim to modulate Treg (defined at least by CD25 or Foxp3 expressing CD4+ T cells) number and/or efficacy in models of experimental brain ischemia were included in the analysis. The overall effect size estimation revealed an odds ratio favoring Treg-targeted interventions [31]. However, a large heterogeneity between the included studies regarding study design, experimental model and species (rat and mouse) has to be taken into account. Therefore, the number of studies investigating Treg interventions has to be increased to more robustly estimate a realistic effect size. Also the publication of negative results has to be encouraged to avoid a publication bias in such an emerging research field as Treg-targeted cell therapies for acute brain injuries.

5 Mechanisms of Treg Function in Post-Stroke Neuroinflammation

Basic immunological research on Treg function has identified several mechanisms by which Treg cells suppress an immune reaction [32]. These can be mainly divided into mechanisms acting on lymphocyte activation and mechanisms inhibiting antigen-presenting cells of the innate immune system. While most of these mechanisms can be recognized using *in vitro* model systems, the contribution of individual mechanisms such as different anti-inflammatory cytokines or cell–cell-contact dependent suppression is highly diverse in between *in vivo* disease models. The most prominent mechanisms of Treg function *in vivo* are the secretion of anti-inflammatory cytokines (IL-10, TGF- β), expression of immunosuppressant molecules (CTLA-4, CD39, PD, consumption of vital cytokines (IL-1, IL-2) and the secretion of cytolytic molecules (granzymes, perforin) [6].

In the model of experimental brain ischemia, several studies have demonstrated that IL-10 is a critical neuroprotective cytokine regulating post-stroke neuroinflammation [33–36]. The main sources of cerebral IL-10 are regulatory lymphocytes

(T and B cells) and microglia/monocytes. The key role of IL-10 as a primary effector mechanism of Treg cells in tissue injury [37] was verified in some of the experimental stroke studies [16, 19, 38]. Accordingly, strategies increasing lymphocyte-derived IL-10 production [19, 29, 35, 39] or therapeutic IL-10 administration [33, 36, 40] have been shown to be beneficial for stroke outcome. In addition, alternative Treg mechanisms have been identified also in brain ischemia models such as the expression of TGF- β and IL-35 [27] as well as a role of Treg cells in the PD-1/PD-L1 pathway. Pro-inflammatory lymphocytes and brain-resident microglia/macrophages have been characterized as the target cells for the immunosuppressive effect of Treg cells after brain injury. Studies that have identified a beneficial effect of Treg detected an inhibition of T cell brain invasion [16, 28], suppression of effector T cell proliferation [19], reduced production of lymphocytic and monocytic cerebral cytokines [27, 28, 41], suppression of microglia/monocyte activation [16] or a priming towards a M2-like microglial phenotype [27]. In contrast, studies detecting an exacerbation of stroke burden by Treg-targeted therapies [26, 30] have found an association of the increased lesion volume with amplified cerebral immune cell accumulation [30], however, it is assumable that the deleterious effects observed by Treg in these two studies might be independent of an immunological function but rather attributable to an impact on secondary microthrombosis [26].

6 Circulating Treg Cells After Stroke

Acute brain ischemia induces profound alterations of the peripheral immune reaction, encompassing peripheral immune activation in the acute phase after brain injury [42] and an immunosuppressive syndrome in the later phase [43]. In addition to the effect of Treg in modulating central neuroinflammation, their functional role has also been detected in alterations of the peripheral immune system after stroke. A consistent finding in the subacute phase after extensive experimental stroke is that cellular immunosuppression and splenic atrophy is accompanied by a relative expansion of Treg cells in spleen and blood [44, 45]. Interestingly, a recent report suggested that the adoptive transfer of Treg cells reduced on one side the systemic inflammatory reaction in the acute phase after stroke and on the other side also ameliorated the extent of immunosuppression (i.e., lymphopenia) in the later subacute phase [46]. This finding of a dualistic role of Treg in systemic immunity comparing the acute and subacute phases after stroke is well in line with the concept that the initial overactivation of the peripheral immune system might result in the later immunosuppression syndrome [47]. Thereby, the potent homeostatic function of Treg might suppress the initial systemic inflammation, attenuating the subsequent immune disturbances such as lymphopenia due to activation-induced lymphopenia cell death and exhaustion of antigen-presenting cells.

7 Regulatory T Cells in Stroke Patients

Only very few studies have investigated regulatory T cells in stroke patients. Understandably, these studies have focused on peripheral immune effects after stroke by investigating blood samples of patients with different stroke entities. The first study specifically analyzing Treg function after stroke by Hug et al. has found that Treg function is preserved in the context of post-stroke immunosuppression in contrast to the dysfunction of effector cell populations such as circulating monocytes or helper T cells [48]. In contrast, a second study has found reduced suppressive capacities of post-stroke Treg in female but not male stroke patients, proposing sex-specific effects on post-stroke peripheral immunity [49]. This difference between the studies could be explained by differing patient characteristics in terms of comorbidities and stroke severity. While the latter study has detected a robust increase of Treg cell counts after stroke in accordance with experimental studies, another report has shown the opposite. Li et al. reported a significant reduction of peripheral Treg in stroke patients [50]. In contrast to the previous studies investigating post-stroke effects on peripheral Treg function, a report by Wigren et al., analyzed the association of Treg counts in a cohort of 700 participants of the Malmö Diet and Cancer study with the prospective incidence of stroke. While low Treg counts at baseline were associated with an increased risk of myocardial infarction, this association was not present for stroke [51]. Overall, clinical data is supporting the experimental finding of substantial peripheral immunomodulation after stroke, which is also affecting the Treg population, yet, specific changes might depend on stroke entity, severity, and patient characteristics.

8 Potential Causes for Differential Effects of Treg Function in Post-Stroke Neuroinflammation

As previously discussed, considerable discrepancy on the role of Treg in stroke can be observed in the literature. Based on our current knowledge of mechanisms involved in post-stroke secondary neuroinflammation as well as the impact of acute brain injuries on the peripheral immune system [1, 43, 52], the most probable reasons for this phenomenon shall be discussed in the following paragraph.

8.1 *Thromboinflammation and Secondary Microthrombosis*

The commonly used transient MCA occlusion models by intraluminal filaments which were also used in the two studies detecting a deleterious role of Treg in stroke [26, 30] can be described as a transient mechanical vascular occlusion model (TMVO) with prompt revascularization. One prominent and distinct feature of this model is

the occurrence of a delayed neuronal damage due to microthrombosis [53]. Notably, the occurrence of microthrombosis is not common to human stroke and animal models of permanent or gradual reperfusion [54] and might rather represent a subpopulation of stroke patients receiving endovascular reperfusion therapy with immediate recanalization of a proximal artery. In experimental models, the presence of such microthrombosis might be determined by the extent of endothelial damage induced by the filament insertion, speed of filament retraction, and occlusion time. Finally, these at first-sight minor methodological differences might have a major pathophysiological impact by the presence or absence of secondary thrombosis [54, 55].

8.2 *The Impact of Lesion Volume on Stroke-Immunology*

As noted above, it seems that lesion size might be a confounder independent of reperfusion. It is well acknowledged that the phenomenon of post-stroke peripheral immunosuppression occurs in stroke patients and animal stroke models only after extensive brain tissue injury [43, 45, 52, 56]. While mice and men with substantial brain lesions develop substantial lymphopenia and alterations of the monocyte population, small ischemic injuries induce only a minor immunomodulation but no immunosuppressive syndrome. While the impact of peripheral immunosuppression after major stroke on secondary neuroinflammation was to our knowledge not systematically investigated, it is conceivable that peripheral immune alterations might affect also central neuroinflammation. Indeed, a recent previous study performing a face-to-face comparison of the extent of cerebral leukocyte invasion, microglial activation, and cytokine secretion in three common models of brain ischemia of differing lesion size detected crucial difference in between models with manifold stronger inflammatory reaction in small permanent ischemia models than in extensive hemispheric lesions after transient MCA occlusion [17]. Therefore, it is plausible that regulatory T cells have an inferior role in stroke models with only minor bystander inflammation compared to the critical role of Treg cells in lesion models with an overshooting immune reaction (Fig. 2).

8.3 *Milieu-Dependent Treg Function*

Moreover, the above stated methodological difference including the presence of microthrombosis/thromboinflammation, induction of peripheral immune alterations, and the extent of neuroinflammation might have a direct impact on the functional properties of natural Treg cells. It is known from other disease paradigms that Treg function in vivo might be particularly dependent of the immunological milieu. For example in cancer research this phenomenon—very much alike to stroke-immunology—has been termed as the “Janus-faced function of Treg” [57, 58]. In addition, the currently predominant perception of neuroinflammation after acute brain injury and particularly stroke research as *too much of a bad thing* should be revisited. Several elegant reports by the groups of Michal Schwartz and Jonathan

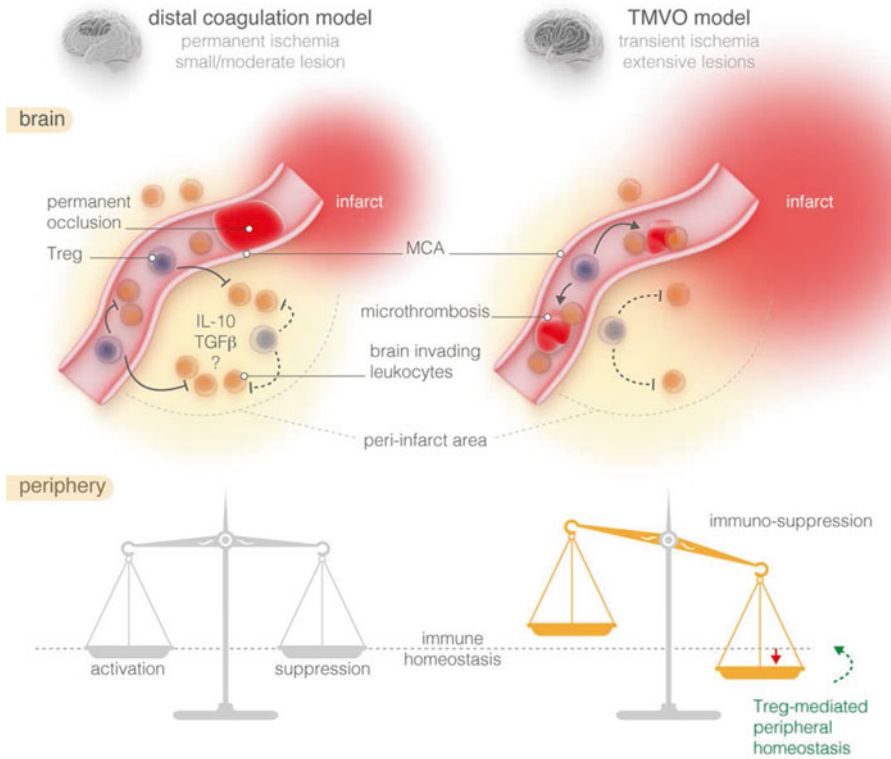


Fig. 2 Differential role of regulatory T cells in experimental stroke models. The schematic diagram illustrates the discrepancy of the observed role of Treg in the brain (*upper panels*) versus peripheral immune system (*lower panels*) in models of permanent ischemia with small- to moderate-sized lesions or after extensive brain lesions in TMVO models. Permanent occlusion of the distal middle cerebral artery (MCA; *left panels*) induces a strong neuroinflammatory reaction, but preserves peripheral immune homeostasis with only minor immunomodulation. In this context, Treg have a primary role in inhibiting an overshooting inflammatory reaction mediated by pro-inflammatory leukocytes. It is assumed that this immunosuppressive function of Treg takes place in the periphery even before brain invasion. In contrast, TMVO models with extensive brain injuries (*right panels*) have less pronounced neuroinflammation, but induce an immunosuppressive phenotype of the peripheral immune system. Here, Treg have only a minor function in suppressing the neuroinflammatory response, and might even have a non-immunologic function in the manifestation of secondary microthrombosis and thromboinflammation. The principal role of Treg in TMVO models seems rather to be in ameliorating immune disturbances by inhibiting initial immunologic activation or overactivation and later immunosuppression, thereby preserving homeostatic systemic immune function. *IL-10*, interleukin 10; *TGFβ*, tumor growth factor β (Reproduced from Liesz et al., *Stroke*, 2015)

Kipnis have established the concept of “protective autoimmunity” (see Schwartz and Raposo [59] for a recent review of the concept). This concept ascribes secondary inflammation as a generally physiological and protective mechanisms in which too much of immune activation as well as immunosuppression might be deleterious. Accordingly, in certain situations of acute brain lesions and neurodegeneration

models—determined by the time-dependent inflammatory milieu of the brain during disease progression—Treg depletion as well as its augmentation might negatively affect the outcome [60–62]. Therefore, the very specific properties of the used stroke model and targeted mechanism of Treg function have to be carefully investigated before considering Treg in a potentially oversimplified model as *good or bad* immune cells after stroke.

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B-Cells in Stroke and Preconditioning-Induced Protection Against Stroke

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Abbreviations

APCs	Antigen-presenting cells
BAFF	B-cell-activating factor
BBB	Blood–brain barrier
B cells	B-lymphocytes
BCR	B-cell receptor
BDNF	Brain-derived neurotrophic factor
Bregs	Regulatory B-cells
CCL2	Chemokine (C–C motif) ligand 2
CLN	Cervical lymph node
CNS	Central nervous system
CSF	Cerebrospinal fluid
CSPGs	Chondroitin sulfate proteoglycans
CXCL13	Chemokine (C–X–C motif) ligand 13
DAMPs	Danger-associated molecular pattern molecules
EAE	Experimental autoimmune encephalomyelitis
ICAM-1	Intracellular adhesion molecule
Ig	Immunoglobulin
IL-	Interleukin
IFN γ	Interferon γ
LPS	Lipopolysaccharide
LTP	Long-term potentiation

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MAP2	Microtubule-associated protein 2
MBP	Myelin basic protein
MCAo	Middle cerebral artery occlusion
MHC	Major histocompatibility class
MOG	Myelin oligodendrocyte glycoprotein
MS	Multiple sclerosis
μ MT ^{-/-}	Transmembrane exon of the IgM heavy chain
MZ B-cells	Marginal zone B-cells
NA	Noradrenaline
nIgM	Natural IgM
NR2A	<i>N</i> -methyl-D-aspartate (NMDA) receptor subunit 2A
RA	Rheumatoid Arthritis
RF	Rheumatoid factor
RHP	Repetitive hypoxic preconditioning
ROS	Reactive oxygen species
Sema3a	Semaphorin IIIa
SHP	Single-exposure hypoxic preconditioning
SLE	Systemic lupus erythematosus
TAC1	Transmembrane activator and calcium modulator and cyclophilin ligand interactor
TGF- β	Transforming growth factor β
T _h cells	T helper cells
TLR	Toll-like receptor
TNF α	Tumor necrosis factor α

1 Introduction

Within the last decade, research into the role of the immune system following stroke onset has revealed a complex inflammatory response that contributes to both stroke-induced injury, as well as central nervous system (CNS) repair. It is now generally accepted that most leukocyte subsets, including both the innate and adaptive arms of the immune system, are activated within the brain parenchyma following stroke. What is less understood is how the expression, timing, and duration of accumulating leukocytes affect functional recovery in the injured CNS. In fact, it may not be that all inflammation is inherently detrimental to recovery, particularly with regard to regulatory lymphocytes, including T-regulatory cells (Chap. 11). B-lymphocytes, or B-cells, also exhibit a potential impact on recovery mechanisms, particularly with regard to neuroprotective preconditioning treatments. This chapter will begin with an overview of B-cell function, and the role of B-cells in other CNS and systemic inflammatory diseases, with relevance to the clinical stroke populations. The chapter will then focus on the role of B-cells in stroke pathology, as well as preconditioning-induced protection from stroke, including potential mechanisms derived from both animal and clinical preconditioning studies.

2 An Overview of B-Cell Function

2.1 Antigen Processing and Presentation by B-Cells

B-cells require antigen exposure to become fully activated [1–3]. After their initial development in the bone marrow, B-cells migrate to the spleen and differentiate into mature, naïve B-cells. These naïve B-cells enter the bloodstream, circulate, and localize to secondary lymphoid organs, including lymph nodes and spleen. These environments aid B-cells in encountering antigens, which are either soluble and enter through the afferent lymphatics, or large particulate antigens presented on the nearby cell surface of antigen-presenting cells (APCs). This results in an immunological synapse forming between the B-cell and the APC, which can include macrophages (Chap. 10) and follicular dendritic cells. This immunological synapse consists of the B-cell receptor (BCR) on B-cell surface, which recognizes the antigen, surrounded by a ring of adhesion molecules, which improves adherence to the APCs [4, 5]. After the external antigens are recognized and captured by B-cells through their BCR, they undergo processing in the internal compartments of the B-cell for subsequent presentation to CD4⁺ T-cells (i.e., T helper cells; T_h cells). Specifically, antigen is broken into peptide fragments bound to major histocompatibility class (MHC) II molecules on the B-cell surface, which shifts the role of the B-cell into that of an APC for CD4 T-cell activation [6].

2.2 Activation of B-Cells Through CD4 T-Cell Interactions

Thus, one of the primary functions of B-cells is to directly activate T-cells during an adaptive immune response. After encountering an antigen, B-cells require accessory signals directly from microbial constituents or from an activated CD4 T-cell to form germinal centers for activation and proliferation [7, 8]. This necessity for accessory signals creates a tight control over the differentiation of naïve B-cells into memory, and/or antibody-producing plasma cell populations [9]. The process of antigen recognition is highly complex, and antigens that activate T-cells can be completely distinct from the surface epitopes recognized by B-cells. In fact, T-cells and B-cells need not even necessarily recognize the same protein before directly interacting, which could lead to quick activation following a massive release of antigen, such as occurs following stroke.

B cells interact with CD4 T-cells in “T-cell zones” within the spleen and lymph nodes [10]. When B-cells enter the lymphoid tissue through permissive systemic venules, they migrate through the T-cell zones into the B-cell zone. Here, those B-cells bound to antigens are trapped through activation of various adhesion molecules and engagement of chemokine receptors, thereby optimizing the rate of interaction with an activated CD4 T-cell at the border of T- and B-cell zones. The specific B-cells, activated by CD4 T-cells, clonally expand and both B-cells and T-cells

proliferate for several days [11]. This is effectively the first phase of primary humoral immune response, and has previously been considered to only occur in the CNS during rare pathology [12]. Recent work by Louveau and colleagues [13], however, identified lymphatic vessels within both murine and human brains, which challenges the dogma that the CNS is predominantly immune-privileged and lacks the lymphatic system found in the periphery. Thus, future studies will need to determine if germinal centers develop, either within the parenchyma or adjacent cortical lymphatics, under pathological states that involve antigen presentation and immune activation, as occurs following stroke [14].

2.2.1 B-Cell Differentiation and Relevant Subsets

Activated B-cells in germinal centers, which are continually refined during the entire germinal center response, differentiate into antibody-secreting plasma cells or memory B-cells [15]. B-cells differentiate into plasmablasts secreting high-affinity antibody at rapid rates, but lose MHC II expression and thus the ability to activate CD4 T-cells [16, 17]. Plasma and memory cells exhibit a wide range of lifespan, with some longer-lived plasmablasts accounting for a persistent, long-lasting source of high-affinity antibody [18]. As reviewed below, these long-lasting B-cell populations may have prolonged effects in clinical stroke recovery [19, 20].

2.2.2 Innate-Like B-Cells

Apart from the B-cells described above which stem from the bone marrow (i.e., B-2 cells), there are heterogeneous populations of unconventional B-cells (e.g., B-1 cells, marginal zone (MZ) B-cells) which play major roles in innate sensing and rapid immune response [21, 22]. B-1 cells produce more mature naive cells in peripheral lymphoid tissues, in contrast to conventional B-2 cells, which only divide upon antigen exposure. B-1 cells secrete antibodies which are polyreactive, low-affinity antibodies at steady state with broad reactivity [22]. When they get activated, they can rapidly acquire immune regulatory activities through the secretion of natural immunoglobulin (Ig)M and interleukin (IL)-10. Thus, these innate-like B-cells constitute an important source of IL-10-producing regulatory B-cells (Bregs), which have been shown to play critical roles in autoimmunity, inflammation, and infection following stroke, as described below.

2.2.3 B-Regulatory Cells and IL-10

Multiple murine disease models for both autoimmune [23–25] and inflammatory diseases [26] confirm the potential for B-cells to downregulate pro-inflammatory immune responses. In particular, Bregs exert immunosuppressive activity through

both direct and indirect mechanisms, including production of cytokines (e.g., IL-10, IL-35, and transforming growth factor β (TGF- β) [27, 28], and expression of inhibitory molecules to downregulate the expansion of pathogenic lymphocytes in a cell contact-dependent manner [28–30]. Regulatory T-cells are classically identified by the expression of Foxp3 marker [31, 32]. However, there is no known master transcription factor for Bregs, which are instead defined based on their functional specificities [31, 32]. Similar to regulatory T-cells, Bregs need to be activated before exerting their functions, with different inflammatory environments inducing different types of IL-10-producing regulatory B-cells [31, 33, 34]. Breg populations are highly heterogeneous, though it has been proposed to classify Bregs as “innate type” and “adaptive type” [29].

“Innate” Bregs are derived from innate-like B-cells, and are characterized by the capacity to rapidly produce high amounts of IL-10 and IgM antibodies upon activation [35, 36]. These “innate” Bregs are activated following toll-like receptor (TLR) or microbial stimulation. Some TLR agonists, like lipopolysaccharide (LPS) from gram-negative bacteria, induce IL-10-producing Bregs and reduce disease severity in a murine model of autoimmune disease [37]. Interestingly, systemic LPS preconditioning can induce tolerance to stroke injury [38, 39], though a link between LPS-induced “innate” Bregs and neuroprotection has yet to be established. On the other hand, “adaptive” Bregs demonstrate antigen specificity and are generated via BCR and CD40 responses, and possibly TLR signaling. In the murine autoimmune disease experimental autoimmune encephalomyelitis (EAE) model (described below), B-cells re-stimulated with auto-antigen produced IL-10 through CD40, and loss of Breg induction exacerbated disease [23].

3 Understanding the Contribution of B-Cells to Other Disease States

B-cells are mainly considered to function through antibody production, with effects limited to downstream events including antibodies coating antigen, activation of the complement pathway, and antibody-dependent cellular cytotoxicity [40]. Though antibodies do play a major role in defense against foreign pathogens, antibodies directed to self-antigen (i.e., autoreactive) can directly induce inflammatory conditions resulting in tissue damage. For example, B-cells are present in airway mucosa, and IgE antibodies are involved in allergic respiratory diseases such as asthma [41, 42]. But apart from antibody production, B-cells can also function as antigen-presenting cells, produce inflammatory and regulatory cytokines, as well as anti-inflammatory cytokines. These latter functions, in particular, may contribute to B-cell-mediated protection in the injured CNS. Unfortunately, there is a paucity of B-cell research compared to investigations of post-stroke monocyte, neutrophil, and T-cell populations. As such, it is first necessary to review the role of B-cells established through studies in other inflammatory disease states.

3.1 The Detrimental Role of B-Cells in Autoimmune Disease

The primary function of immune system is to differentiate between self and foreign antigens, and produce effector responses toward the foreign pathogens, whilst avoiding destructive self-targeting [43]. Autoimmune diseases generally involve abnormal recognition of self-antigens, resulting in severe inflammatory conditions and subsequent tissue damage. This occurs due to a combination of both environmental and genetic factors, which leads to a loss of tolerance toward self-antigens. B-cells are usually considered to be detrimental with respect to autoimmune diseases because of the secretion of autoantibodies, which damage tissue and bind receptors, acting as agonists to either activate or inactivate function [44, 45]. Furthermore, B-cells activate the complement system, leading to the release of anaphylatoxins which increase the secretion of pro-inflammatory cytokines and attract other effector cells to the site of injury [46, 47]. Antibody-independent pathogenic functions include providing accessory signals to T-cells to activate an autoreactive T-cell response [48–50]. B-cells secrete pro-inflammatory cytokines like tumor necrosis factor (TNF) α and interferon (IFN) γ [51], and areas of chronic inflammation form ectopic germinal centers [52]. These germinal centers are similar to the germinal centers of the secondary lymphoid organs described above, though the plasma cells in these structures secrete autoantibodies [53]. While at first these diseases may seem irrelevant to stroke, there is a call for the medical community to re-evaluate the role of chronic inflammatory diseases in the development of atherosclerosis, a predominant risk factor for stroke [54].

3.2 Multiple Sclerosis and EAE

Multiple sclerosis (MS) is an inflammatory and demyelinating disease of the human CNS. Most patients with MS exhibit lesions comprised of immune cell infiltrates, demyelination plaques, and axonal damage, with major heterogeneity of the pathological features in patients [55, 56]. EAE is an induced autoimmune disease in mice to model MS, though it is predominantly considered T-cell-mediated, as the adoptive transfer of myelin-specific autoreactive T-cells are sufficient for disease induction [57, 58]. CD4 T-cells specific for CNS antigens, such as myelin oligodendrocyte glycoprotein (MOG), induce inflammation and result in demyelination of the CNS, though IL-10-producing T-cells ameliorate disease [59–61]. The importance of the role of B-cells in MS was first stressed by the reports of a presence of high levels of immunoglobulins in the cerebrospinal fluid (CSF) MS patients [62]. While EAE induction does not require autoantibody production, they can exacerbate the existing inflammatory environment [63, 64]. Studies involving B-cell depletion using anti-IgM antibodies found that B-cell depleted mice were more resistant to the induction of EAE via recombinant MOG protein [64–66]. On the contrary, B-cell depletion during the later parts of the disease resulted in reducing the disease symptoms [57], possibly as B-cell antigen presentation and activation affects the entry of autoreactive T-cells into the CNS [67, 68].

3.3 Systemic Lupus Erythematosus

Systemic lupus erythematosus (SLE) is a prototypical autoimmune disorder characterized by a significant increase in autoantibodies against a variety of predominantly nuclear self-antigens, (e.g., DNA, ribonucleoproteins, and phospholipids) [69]. SLE patients exhibit diverse symptoms because of the heterogeneous impact of the disease on different tissues and organs, including kidney [70–73] and brain [74], resulting in a challenge to diagnosis the disease. B-cell depletion in SLE mouse models resulted in the absence of nephritis and vasculitis [75], and decreased activation of T-cells [76]. Considering the significance of B-cells in the initiation and progression of SLE, treatments targeting B-cell survival and differentiation for B-cell depletion therapy are being evaluated for clinical applications [77].

3.4 Rheumatoid Arthritis

Rheumatoid arthritis (RA) is an autoimmune disease which results in inflamed synovial tissue due to infiltration of inflammatory cells into the affected joints [78]. The synovial membrane contains numerous activated CD4 T-cells, which are important for the pathogenesis of RA. B-cells function as APCs to prime autoreactive T-cells in experimental RA [79]. B-cells surround the T-cell aggregate, with activation dependent on the presence of B-cells [78]. Experiments where inflamed tissue samples from RA patients were transplanted into mice lacking lymphocytes confirmed the requirement of B-cells for both disease initiation and direct sustainment of chronic inflammation [80, 81]. B-cells also produce rheumatoid factor (RF) antibody production [82], with the presence of RF antibody correlated with increased mortality and morbidity in RA patients [83].

3.5 Current FDA-Approved B-Cell Therapeutics

CD20 is a B-cell surface marker expressed in different stages of B-cell development. Patients with autoimmune diseases, including MS, SLE, and RA, positively respond to B-cell depletion treatments using an anti-CD20 antibody, commercially known as Rituximab [77, 84–86]. In contrast, B-cell depletion exacerbates disease severity in patients with certain T-cell-mediated autoimmune conditions, including ulcerative colitis and psoriasis [87, 88]. Determination of efficacy of B-cell depletion is limited to changes in B-cell populations in the peripheral circulation. Therefore, the full effect on B-cells population in secondary lymphoid tissues outside of bone marrow, including the CNS, remains unknown [89].

Other B-cell-specific therapies are also under development, including the targeting of B-cell receptors involved in proliferation. A double-blind phase-II study with

Belimumab, a monoclonal antibody affecting the function of the B-cell receptor through the B-cell-activating factor (BAFF) [90]. BAFF is essential for B-cell survival and maturation [91, 92], with Belimumab significantly lowering B-cell counts for active RA patients [90]. Atacicept (i.e., antibodies against transmembrane activator and calcium modulator and cyclophilin ligand interactor (TACI), another BAFF receptor required for B-cell maintenance and survival) also decreased B-cells in preclinical studies [93], with a Phase-Ib study with patients having moderate to severe RA, demonstrating efficacy [93, 94]. Finally, BR3-Fc, a recombinant protein that blocks BAFF signaling, is under evaluation in clinical trials [95].

Currently, most B-cell therapeutics are focused on B-cell depletion and antibody efficiency [96]. Induction of Bregs, however, also offers a potential therapeutic strategy, though further studies are required to establish the functional stability of Bregs [44]. However, there are some clinical trial results which support this avenue of intervention. Anti-IL-6R (Tocilizumab) treatment for RA increased TGF- β expression in CD25^{high} B-cells, and could potentially function by regulating Bregs [97]. Also Laquinimod, a drug under development for MS and in phase-III clinical trials for SLE, increases the percentage of CD25^{high} B-cells and their IL-10 expression, thereby expanding subpopulations of Bregs [98, 99]. Finally, Copaxone could prove efficacious in upregulating Bregs after stroke, as its effects include promoting the conversion of inflammatory CD4 T-cells to Tregs [100], the upregulation of B-cell-derived IL10 [101, 102], and the neuronal production of growth factors [103], though efficacy in experimental stroke models has been mixed (see Sect. 3.2).

4 B-Cells During Stroke Injury and Recovery

Stroke occurs when there is a loss of blood supply to the brain, either due to a vessel blockage (ischemic stroke) or vessel rupture (hemorrhagic stroke), and is a major cause of death and disability in the global adult population [104–106]. Despite significant improvements in the understanding and identification of factors contributing to post-stroke CNS tissue damage, therapeutic options for stroke are highly limited. Apart from the primary tissue damage due to lack of blood supply, there is a secondary ischemia-reperfusion injury due to the inflammatory cascade. Following the occurrence of either ischemic or hemorrhagic stroke, there is disruption of the blood–brain barrier (BBB), allowing endogenous CNS antigens access to activated lymphocytes [107]. Shortly after experimental stroke onset, endothelial cells within the BBB are activated to recruit peripherally circulating leukocytes to infiltrate into the brain parenchyma. This early CNS sterile inflammation induces an immediate and strong antigen-nonspecific innate immune response against the exposed CNS molecules, resulting in brain tissue damage and functional impairment, which later results in an antigen-specific adaptive immune response [108]. However, not all of the post-stroke immune responses are inherently detrimental to functional recovery. This section will review post-stroke inflammation, with a focus on B-cell-mediated mechanisms of both injury and repair in the CNS.

4.1 Post-Stroke Immune Responses Within the CNS

After stroke, there is a loss of BBB integrity, while reactive oxygen species (ROS) and other inflammatory mediators activate endothelial cells and leukocytes to increase expression of adhesion molecules [109, 110]. This results in the migration and recruitment of peripherally circulating leukocytes to the ischemic cortex, allowing the immune cells to access novel CNS antigens. While this is the first step of the inflammatory cascade, the acute inflammatory response consists of both infiltrating leukocytes and tissue-resident immune cells, including microglia (Chap. 8) [111]. Resident microglia are activated immediately after stroke and secrete pro-inflammatory mediators (e.g., IL-1 β , TNF- α). Neutrophils are one of the first leukocytes to infiltrate the ischemic tissue, entering within a few hours after experimental stroke [112, 113]. Neutrophils secrete granules containing pro-inflammatory factors, elastases, myeloperoxidases, and multiple matrix metalloproteinases. The macrophages and mast cells residing in the perivascular space also release multiple pro-inflammatory factors, like TNF- α , IL-8, and IL-1 β , as well as vasoactive mediators like histamine and proteases [114–117]. This highly enhances the infiltration of peripheral immune cells into the infarct region in a feed-forward mechanism that collectively constitutes the acute phase of the post-stroke innate immune response.

Lymphocytes, including T-cells and B-cells, enter the brain parenchyma early after onset as well, though T-cells function in the acute phase without any antigen specificity through secretion of cytokines and ROS generation [112, 113]. The dead and damaged cells in the infarct release an array of molecular signals called danger-associated molecular pattern molecules (DAMPs) [118]. These bind to and activate TLRs and scavenger receptors present on APCs to prime them for antigen presentation. This results in the CNS antigens being presented to B-cells and T-cells [14], and the subsequent development of cellular and humoral immunity against the CNS antigens. Data concerning antigen-dependent post-stroke adaptive immune responses are contradictory, as both beneficial and harmful effects to the brain have been shown in experimental stroke models [119–121].

Ortega and colleagues recently characterized the time course of autoreactivity to myelin-derived antigens (e.g., MOG, myelin basic protein (MBP)) and neuronal-derived antigens (e.g., microtubule-associated protein 2 (MAP2), *N*-methyl-D-aspartate (NMDA) receptor subunit, NR2A) [14]. Experimental stroke induced a high autoreactivity to CNS antigen as early as 4 days after onset for both B- and T-cell populations isolated from the spleen and cervical lymph node (CLN). The CLNs are draining lymph nodes for brain, but the rapid response may reflect a previously unstudied APC activation of B- and T-cells occurring in newly identified lymphatic vessels within the CNS [13]. High autoreactive responses to MAP2 and myelin occurred in the mice with the smallest infarct volumes, suggesting autoimmune responses after stroke may not be solely detrimental to recovery. This corroborates a clinical study in stroke patients, wherein Planas and colleagues demonstrated that increased APC presentation of MAP2 and NR2A correlated with smaller infarct volumes and better long-term outcome [122]. This study also found that higher reactivity for MBP correlated with larger infarctions and greater stroke

severity [122], though T-regulatory cells specific for MBP [103], and MBP antigen presentation before stroke [123] improved long-term recovery in animals [14]. Thus, further studies are required to completely elucidate this complex and ambiguous adaptive immune response to understand the mechanisms related to CNS recovery that could be harnessed as potential therapeutic targets.

4.2 B-Cell Potential to Limit Detrimental Acute Post-Stroke Inflammation

As mentioned previously, regulatory T- and B-cells produce anti-inflammatory cytokines that suppress the function of infiltrating pro-inflammatory cells. In fact, IL-10 expression is elevated during major CNS diseases [124], and regulatory lymphocytes play a protective role, limiting the CNS damage due to inflammation in autoimmune diseases such as MS and SLE [44, 61, 125]. Therefore, it is hardly surprising that in the absence of B-cells, there is increased infiltration of multiple leukocyte subsets into the ischemic parenchyma, including neutrophils, monocytes, macrophages, and T-cells, resulting in larger infarct volumes [124]. Confirmatory studies used transgenic mice that have a nonsense mutation introduced into the transmembrane exon of the IgM heavy chain ($\mu\text{MT}^{-/-}$), which results in the total deletion of B-cells [126]. These mice developed larger infarct volumes, more severe functional deficits, and had a higher mortality rate following middle cerebral artery occlusion (MCAo), a common experimental model of stroke [127]. The complete absence of B-cells also resulted in higher numbers of activated T-cells, macrophages, microglial cells, and neutrophils in the ischemic brain of $\mu\text{MT}^{-/-}$ mice following MCAo [127].

This phenotype was rescued after the adoptive transfer of highly enriched populations of wild-type B-cells to B-cell-deficient $\mu\text{MT}^{-/-}$ recipient mice before MCAo [127]. The neuroprotective, anti-inflammatory effect was not observed, however, when IL-10-deficient B-cells were adoptively transferred prior to stroke. Mice receiving B-cells as a neurotherapeutic exhibited higher numbers of regulatory cells in the periphery concomitant with lower numbers of activated inflammatory T-cells [128]. After LPS stimulation, purified B-cells produced significant amounts of IL-10. Adoptive transfer of these cells prior to stroke decreased infarct volumes, confirming the beneficial role of IL-10-producing B-cell subsets on stroke outcome in the acute phase of recovery [124]. These B-cell subsets also increased the regulatory T-cell numbers, co-inhibitory receptor PD-1 expression, and significantly reduced peripheral pro-inflammatory immune cells [124].

As mentioned previously, Copaxone as a therapeutic can induce IL-10 from Bregs [102]. Copaxone treatment (200 μg plus an adjuvant to boost the immune response) decreased infarct volumes in rats 7 days post-MCAo [129]. More recent studies, however, found that Copaxone (70 μg , no adjuvant) administered 30 mins prior to stroke did not reduce infarct volumes at 1 day after stroke [130], nor did Copaxone administration (400 μg , no adjuvant) immediately after MCAo reduce

infarct volumes at 3 and 7 days post-stroke [131]. But the latter study also did not find an upregulation of IL-10, which suggests that the lack of protection in these studies may be confounded by the lack of adjuvant, which increases inflammation, and thus Copaxone uptake, at the site of injection. This may also explain the shortened timeframe given for Copaxone to generate a neuroprotective effect, particularly with regard to the 1-day post-stroke study [130]. Therefore future studies should consider Copaxone as a pharmacologic preconditioning agent (similar to LPS, 4.4.1) when determining if it can induce neuroprotection via Breg upregulation prior to stroke onset.

4.3 Post-Stroke Immune Responses Occurring in the Periphery

Stroke induces systemic immune changes, which affect clinical outcomes for weeks and months following stroke onset [132, 133]. As mentioned above, the initial response to tissue injury following stroke is carried out primarily by the innate immune system, as innate cells are recruited to the ischemic parenchyma and contribute to infarct progression [112]. These inflammatory responses are not confined to the brain, however, but are also concurrent with immunological changes in blood, spleen, bone marrow, and secondary lymphoid organs [134, 135]. Gene expression signatures from the peripheral blood after stroke distinguish different causes of stroke, and this implies a characteristic immune response to CNS injury [136]. Stroke induces systemic immunodepression within days of onset, especially in the cases of large infarct volumes [137]. In fact, most patients with larger strokes have higher levels of circulating IL-6 and IL-10, lower lymphocyte counts, and increased susceptibility to infections (e.g., respiratory, urinary tract) that impede long-term recovery [138, 139]. Liesz et al. used different ischemia models to demonstrate the effect of stroke on systemic immunological and microbiological parameters [140]. They found that post-stroke changes associated with immune system, as well as infectious complications, depended on infarct volume. Only huge infarct volumes induced leucopenia (i.e., reduced numbers of lymphocyte counts) in spleen, lymph nodes, and thymus both during acute (after 24 h) and long-term (after 7 days) recovery.

A majority of these systemic immune changes can be attributed to downstream signaling of noradrenaline (NA) receptors, present on almost all immune cells, including 30–50% of B-cells [141]. NA receptors are activated by epinephrine released from post-ganglionic sympathetic nerves within hours of experimental stroke onset [142]. Thus, the sympathetic nervous system dramatically alters post-stroke immune responses to increase plasma cortisol, catecholamines, and monocyte-derived IL-10, which in turn decreases the number and activity of circulating immune cells, including B-cells [134, 142]. Transient decreases in IgG levels were reported in post-stroke patients, which was also associated with increased bacterial infections [19]. Stroke injury is itself a strong stress signal that also activates the hypothalamic–pituitary axis, resulting in the production of glucocorticoids

which further induce immunosuppression [142, 143]. Treatment with propranol, a β -adrenergic receptor antagonist, as well as other steroid antagonists rescued lymphocyte apoptosis and reduced susceptibility to infection after stroke in animal models. Clinical studies confirmed that patients on beta-blockers at the time of stroke onset exhibited reduced stroke injury [144], in-hospital mortality [145], and certain infectious co-morbidities [146], but translational to an effective post-stroke therapeutic and the effect on B-cell populations has yet to be established.

4.4 B-Cell Contribution to Neuronal Plasticity

Functional recovery after stroke is limited, as only ~65 % of stroke survivors exhibit full functional recovery even 1 year post-stroke [147]. Unfortunately, many of the long-term sequelae that both inhibit and promote the neuronal plasticity required for functional recovery in the injured CNS are still being investigated. Under physiological conditions, axonal growth and sprouting is thought to be minimized in the adult brain [148], but after CNS injury, there is re-expression of axonal growth factors (e.g., GAP43, CAP23, and MARCKS) within the glial scar bordering the infarct [149–152]. Unfortunately, axonal growth inhibitors are also upregulated, including chondroitin sulfate proteoglycans (CSPGs), NogoA, and semaphorin IIIa (Sema3a), which often impair the permissive environment for post-stroke plasticity [153–158]. Recovery from stroke requires substantial axonal regeneration into the injured brain to connect novel cortical areas, and strengthen relevant connections in an effort to subsume function lost after stroke onset [159–162].

Unfortunately, there are few studies linking leukocyte-derived factors and neuronal plasticity in stroke, though some evidence has been found in experimental models of both autoimmune disease and spinal cord injury, particularly with regard to T-cell-axon interactions [163–167]. But B-cells harbor the potential for direct neuronal effects as well, particularly as T-cell-activated B-cells isolated from germinal centers in the spleen express genes related to axonal growth, differentiation, and repair [168]. Furthermore, antibodies derived from B-cells in the CSF of MS patients show a direct ability to bind demyelinated axons in the CNS, with several distinct axon-binding patterns, though their physiological or pathological function(s) are unclear [169]. B-cells accumulate in lesions following spinal cord injury and produce autoantibodies that cross-react with neuronal antigens, with improved recovery and decreased spinal pathology in the absence of B-cells, with respect to wild-type mice [170, 171].

Of the aforementioned axonal growth-related proteins, T- and B-cell immune responses to Nogo-66 can be both encephalitogenic, increasing the severity of the disease, as well as suppressive, lowering the disease score in EAE [172]. Also BAFF, secreted by B-cells, potentially binds Nogo receptors expressed in neurons and astrocytes, directly affecting the neuronal outgrowth [173]. Bregs also express Sema3A on their surface; hence interaction of B-cells with neurons can potentially be detrimental to post-stroke axon growth [99]. Finally, B-cells produce significant

amounts of brain-derived neurotrophic factor (BDNF), a growth factor widely known to support post-stroke neuronal growth in both in vitro and in vivo studies [174–176]. While many mechanisms exist to support the direct interaction of B-cells on neurons, future studies need to confirm a role specific to post-stroke plasticity.

4.5 B-Cells Mediate Post-Stroke Cognitive Impairment

Stroke patients are highly susceptible to dementia, particularly with regard to vascular dementia [177–179]. In fact, a single stroke increases the risk of developing dementia two-fold when compared to other risk factors like hypertension, hypercholesterolemia, diabetes, and age [20, 178, 180]. However, the underlying mechanism linking stroke and the development of dementia is largely unexplored. Recent work by Doyle and colleagues found that B-cell-mediated post-stroke inflammation induced delayed cognitive deficits at 7 weeks after experimental stroke, which mirrors the clinical risk of developing dementia over the years following stroke onset [20]. Four different experimental stroke models in two mouse strains colocalized B-cells to the infarct region 4–7 weeks after onset, with B-cells clustering and producing IgA and IgG antibodies. Doyle et al. hypothesized that these antibodies activate Fc receptors and complement pathways to induce neuronal damage and subsequent neurological dysfunction, similar to antibody actions during MS [169], with the potential for detrimental antibodies to diffuse into surrounding healthy tissues to further exacerbate recovery. This post-stroke antibody production impaired hippocampal long-term potentiation (LTP), resulting in short-term memory deficits beginning weeks after stroke. Delayed cognitive defects did not occur in $\mu\text{MT}^{-/-}$ mice or in mice treated with Rituximab. This role of B-cells in long-term cognitive impairment is in contrast to the role of B-cells in beneficial neuroprotection during the acute stages of stroke [181]. But whether B-cells provide acute neuronal protection, support functional plasticity, or even impede recovery may be dependent on the location, timing, and magnitude of the B-cell response within the injured CNS, which will hopefully be elucidated in future studies.

5 B-Cells During Preconditioning and Ischemic Tolerance to Stroke

5.1 Preconditioning and the Immune System

Preconditioning is the use of sublethal stressors to induce endogenous, global tolerance against subsequent injury [182, 183]. Preconditioning was originally developed to study myocardial ischemia, as Murry et al. observed that four, 5-min coronary branch occlusions, each followed by 5 min of reperfusion, protected against a 40-min

ischemic insult [184]. These brief exposures significantly reduced infarct volume in the myocardium without causing cell death. Since this discovery, preconditioning has been applied to ischemic injury in different organs and tissues, including the liver, kidney, small bowel, skeletal muscle, and retina [185–189]. One promising avenue of research is the use of preconditioning to induce ischemic tolerance in the brain. Preconditioning produces robust neuroprotection in rodent models of stroke, including significant reductions in infarct volumes, strengthened BBB integrity, and improved neurocognitive functioning after stroke [190, 191].

A wide range of noxious stimuli induce neuroprotection, which has led to the development of several preconditioning paradigms (see reviews [183, 192, 193]). One of the classic paradigms of preconditioning is ischemic preconditioning, which includes both brief cerebral ischemia and remote limb ischemia. Other well-studied preconditioning models include oxygen (hypoxia or hyperoxia), temperature (hypothermic or hyperthermic exposures), exercise (voluntary exercise or forced exercise), and various forms of pharmacological preconditioning. Despite the heterogeneity of stimuli used in preconditioning studies, there is a remarkable amount of overlap in the underlying mechanisms responsible for protection. Diverse preconditioning methods are dependent on similar cellular, molecular, and physiological mechanisms involving neurons, astrocytes, and vasculature, suggesting that a complex, multifaceted integration of different systems is necessary for ischemic tolerance [183].

There is increasing evidence that alterations to the immune system are a critical component of preconditioning. During preconditioning, the exposure to a stressful stimulus provokes an inflammatory response within the brain, activating many pathways that play a pathogenic role in stroke [194, 195]. TLRs, cytokines, chemokines, cannabinoids, inducible nitric oxide, and ROS have all been implicated as important mediators of both ischemic injury and preconditioning-induced neuroprotection [196, 197]. Despite their negative role in stroke, these inflammatory molecules appear to be necessary for the induction of endogenous neuroprotection. Studies consistently show that blocking these pro-inflammatory signaling cascades diminishes, if not demolishes, the ischemic tolerance produced by preconditioning [198–200]. The inflammatory response after preconditioning, although not severe enough to cause injury, significantly affects the immune system, “reprogramming” it to respond differently to subsequent injury in favor of a less “pro-inflammatory” response [194, 195].

Despite the growing interest in the role of the immune system in stroke [115, 201], there has been limited exploration of the relationship between adaptive immunity, in particular B-cells, and preconditioning. In the following section, we will discuss how three methods of preconditioning, using the systemic interventions of hypoxia and exercise, as well as the pharmacologic intervention of LPS, are associated with B-cell function, with emphasis on the future translational potential of these findings.

5.1.1 Single-Exposure Hypoxic Preconditioning

Hypoxic preconditioning has been shown to be neuroprotective, both in vivo and in vitro in neonates and adults [192]. However, the efficacy of this protection is dependent on the duration, intensity, and frequency of low-oxygen exposures. Many studies use single-exposure hypoxic preconditioning (SHP), in which an animal or tissue is placed inside an oxygen-controlled chamber filled with 8% O₂/92% N₂ gas for several hours. Exposures that are too brief (<3 h) are not protective against subsequent injury, while those that are too long (>5 h) worsen stroke outcome, increasing neuronal death and behavioral deficits [183, 202]. A single moderate exposure to hypoxia (~4 h) is robustly protective, reducing infarct volume, neuronal apoptosis, and neurologic deficits [203]. However, this protection is brief, as a single exposure protects only for 72 h [202, 204].

The ischemic tolerance seen after SHP has been attributed to both neuronal and non-neuronal mechanisms, including increases in antioxidant enzymes, calcium signaling, expression of hypoxia-inducible factors, erythropoietin, and vascular growth factors, activation of opioid receptors, as well as metabolic suppression [202, 204–211]. In addition to these mechanisms, there is increasing evidence that modulation of the immune system by SHP contributes to ischemic tolerance, specifically through activation of pro-inflammatory pathways (Table 1). After SHP, rat cortical neurons increased production of TNF- α and exhibited higher intracellular levels of sphingolipid ceramide, a secondary messenger involved in TNF- α signaling [200]. Inhibiting TNF- α and ceramide attenuated SHP-induced neuroprotection. Additionally, CCL2, a chemokine responsible for the recruitment of pro-inflammatory monocytes, dendritic cells, and memory T-cells, is upregulated in cortical neurons and endothelium after SHP [198]. In the absence of CCL2, the protective effects of SHP are significantly diminished. Additionally, Stowe and colleagues found that independent of CCL2, SHP significantly altered circulating leukocytes prior to injury, decreasing T-cells, monocytes, and granulocytes 12 h after the hypoxic exposure. Interestingly, despite the decline in other immune cell populations, there was a significant increase in the B-cells following SHP, suggesting a potentially unique relationship between SHP-induced neuroprotection and B-cells.

5.1.2 Repetitive Hypoxic Preconditioning

An important limitation of SHP paradigms is the brief time-window of protection [202, 204]. Thus, recent research has begun to investigate the benefits of repeated exposures to hypoxia prior to stroke, or repetitive hypoxic preconditioning (RHP). In a progressive hypoxia model, mice are placed in air-tight jars until gasping occurs, removed for 30 min of recovery, and then returned for a total of four hypoxic exposures, with some evidence that progressive hypoxia can induce neuroprotection [212, 213]. Another RHP method found that 15-h exposures to hypobaric hypoxia

Table 1 Immune system regulation by hypoxic exposures

Paradigm	Stimuli	Animal	Injury model	Duration of protection	Immune-mediated results	Reference
SHP	8 % O ₂ for 4 h	Mouse	tMCAo	48 h	Histamine is necessary for hypoxia-induced protection and is necessary for neuroprotective increases in VEGF protein	[207]
	20 min hypoxic exposure	Rat	OGD	24 h	TNF- α and ceramide are necessary for hypoxia-induced protection in vitro	[273]
	8 % O ₂ for 4 h	Mouse	tMCAo	48 h	CCL2 is necessary for hypoxia-induced protection	[198]
	8 % O ₂ for 4 h	Mouse	tMCAo	24 h	HIF, SphK, S1P, CCL2 are necessary for hypoxia-induced protection	[276]
RHP	Stochastic oxygen exposures	Mouse	tMCAo	8 weeks	RHP reduced infarct volumes	[215]
				4 weeks	RHP reduced leukocyte diapedesis & leukocyte-endothelial adherence	
	Stochastic oxygen exposures	Mouse	tMCAo	2 weeks	RHP increased CXCL13 expression in the cortex and subcortex	[216]
					RHP increased immunosuppressive B-cells while reduced populations of other leukocytes	

tMCAo transient middle cerebral artery occlusion; *OGD* oxygen glucose deprivation; *TNF- α* tumor necrosis factor alpha; *CCL2* chemokine (C-C motif) ligand 2; *HIF* hypoxia-inducible factor; *SphK* sphingosine kinase 2; *S1P* sphingosine-1-phosphate; *CXCL13* chemokine (C-X-C motif) ligand 13

for 4 weeks decreased infarct volumes for up to 1 week following the final exposure [214]. However, the most effective RHP method to date is a preconditioning using stochastic exposures to hypoxia. Over the course of 2 weeks, animals receive hypoxic exposures varying in duration (2 or 4 h) and intensity (8 or 11 % O₂) [215]. This method of RHP produced robust neuroprotection at 2 weeks after the last

exposure, with a 48 % reduction in infarct volume, but included a 32 % reduction in infarct volume still present 8 weeks after RHP, a significant improvement over untreated animals. RHP decreased vascular inflammation, BBB disruption, leukocyte adherence to vessels, as well as overall leukocyte egress into the ischemic hemisphere [190, 215]. The initial paper describing the stochastic RHP methodology did not investigate specific leukocyte populations present in the brain after preconditioning, but subsequent work focused more extensively on the immune mechanisms behind neuroprotection after RHP.

5.1.3 B-Cells in Hypoxic Preconditioning

The potent protection induced by RHP is, in part, due to significant alterations in the post-stroke inflammatory response during ischemic tolerance. RHP reduced the influx of leukocytes into the ischemic brain, including CD4 T-cells, monocytes, and activated macrophages, concurrent with increased B-cell representation compared to untreated controls [216]. RHP maintained B-cell populations at a level equivalent to that of the uninjured, contralateral hemisphere and, similarly, the ratio of B-cells to monocytes was unchanged. This ratio is used to describe pathological immune environments in the CNS, making it a useful prognostic tool in MS [217] and B-cell lymphoma [198, 218]. Overall, RHP modulation of the immune system after stroke was a CNS-specific phenomenon, with minimal changes seen in peripheral immune populations. CNS specificity may be subsequent to an early RHP-mediated increase in CXCL13, a chemokine that recruits B-cells to sites of inflammation [216]. While CXCL13 is widely present throughout the post-stroke brain, it was largely expressed on cortical vessels, and particularly notable on endothelial cells surrounded by astrocytic endfeet at the BBB. Given the established benefits of RHP on BBB integrity [190, 215], it is possible that the recruitment of B-cells into the ischemic hemisphere contributes to improved neurovascular integrity in preconditioned mice.

Prior to stroke onset, RHP exposure altered the immunophenotype of resident splenic B-cells [216]. Microarray analysis found that over 1900 genes were changed 2 weeks following RHP, producing a unique immunosuppressive B-cell phenotype prior to the induction of injury. These genetic alterations suggest that RHP suppresses B–T-cell interactions. In addition to limiting B–T-cell interactions, RHP also inhibited B-cell proliferation, development, and differentiation, contributing to protective immunosuppression. Complementing these findings, *ex vivo* phenotyping using flow cytometry showed a significant reduction in mature and activated B-cells, as well as a decreased response to stimulation with LPS, suggesting RHP leads to functional impairment and decreased proliferative capacity in B-cells. RHP also induced a higher representation of Bregs, previously implicated in improved outcome after stroke [181]. Indeed, RHP-induced Bregs may directly contribute to the neurovascular protection seen after RHP, although further studies are necessary to confirm a causative role [216].

5.1.4 Clinical Relevance of Hypoxic Preconditioning

Most of the efforts to translate findings from animal models of preconditioning to the clinic have focused on remote ischemic preconditioning, particularly given its success in protecting against myocardial injury [193]. However, there is evidence that hypoxic exposures in moderate doses modulate the immune system in ways that may be protective against stroke. While many studies focus on the innate immune system activation during hypoxia [219], suppression of adaptive immunity is also an important component in the effects of hypoxia on the immune system. After 5 days of extreme hypobaric hypoxia (3200–3800 m), T-cell blastogenesis was impaired in individuals for up to 24 days [220]. Similarly, brief repetitive exposures to normobaric hypoxia have been shown to be anti-inflammatory, significantly decreasing TNF- α in the serum up to 1 week after final exposure [221]. Again, this decrease in TNF- α is consistent with our findings that this cytokine is suppressed through activation of IL-4 pathways after RHP [216]. These clinical studies are limited, though, as they do not look at the effects of hypoxia in the context of illness. However, there is epidemiological evidence that altitude has a positive effect on a wide range of illnesses and injuries such as stroke. A study in Switzerland found that individuals who lived at higher altitudes had decreased risk of coronary heart disease and stroke [222]. But overall, these studies suggest that changes in the immune system due to hypoxia at higher altitudes may lead to a sustained anti-inflammatory phenotype, which may in turn affect susceptibility to injury and illness.

5.1.5 Exercise Preconditioning

Another promising form of preconditioning is exercise preconditioning, which, according to a recent meta-analysis of experimental data, reduces infarct volumes by 42.2% and decreases neurological deficits by 54.6% [223]. Two main forms of exercise preconditioning have been used in animal studies: voluntary exercise and forced exercise. During voluntary exercise, rodents are given unrestricted access to running wheels, wherein they usually run in short, rapid spurts [224]. Conversely, forced exercise on a treadmill requires that animals maintain a slower speed for a longer duration [224]. The efficacy of these two methods varies given that preconditioning-induced neuroprotection is contingent on frequency, intensity, and duration of stimuli. Forced exercise produces more potent protection, reducing infarct volumes by 58.5% as opposed to voluntary exercise, which only led to a 10.4% reduction [223]. However, rats that underwent voluntary exercise preconditioning had greater motor recovery and higher levels of hippocampal BDNF levels after stroke [225]. Forced exercise also increased corticosterone [226], which can impair the immune system by reducing the functionality of T-cells and reducing the number of circulating B-cells [227].

The benefits of forced and voluntary exercise can be attributed to both neuronal and non-neuronal mechanisms. Exercise increases neurogenesis [228–230], angiogenesis [231, 232], and neuronal metabolism [226, 233], while simultaneously

decreasing BBB disruption [208, 234, 235] and apoptosis [236, 237]. Exercise preconditioning also improves stroke outcome through immune-mediated mechanisms (Table 2). Similar to many other forms of preconditioning, exercise preconditioning relies on pro-inflammatory molecules to induce its neuroprotection. Intracellular adhesion molecule (ICAM)-1, an adhesion molecule that promotes leukocyte rolling, and TNF- α mRNA expression, was greater in rats with 3 weeks of forced exercise [238]. But after stroke, the animals that underwent this exercise preconditioning exhibited decreased expression of the TNF- α receptors TNFR1 and TNFR2 [239], I-CAM expression [238], and subsequent leukocyte infiltration [238], similar to our results in RHP-treated mice [215]. Furthermore, exercised rats had lower levels of TLR4 expression [199], which could decrease the production of pro-inflammatory cytokines and contribute to neuroprotection, though this remains to be confirmed.

5.1.6 B-Cells in Exercise Preconditioning

During acute exercise, epinephrine activates the β 2-adrenergic receptors on B-cells, leading to a mobilization of these lymphocytes in the blood [240]. The intensity and duration of exercise modulates this B-cell mobilization, with increased lymphocytosis after intense exercise [240]. Although changes in antibody production in serum appear to be negligible in humans, IgM concentration increases following in vitro mitogen-stimulation [240]. Similarly, rats that had unrestricted access to running wheels showed an increase in natural IgM (nIgM) in their serum and peritoneal cavity [241, 242]. nIgM is beneficial in mounting attacks against bacteria and viruses, as well as activating the complement system, which may be beneficial in post-stroke infection, a common and life-threatening complication in stroke patients described above [242]. Other studies have shown that voluntary and forced exercise increases serum IgG in response to antigens in mice [243–245]. Unfortunately, most studies investigated the effect of exercise on B-cell function within the context of infection, rather than CNS injury. Further studies will need to focus more on the role of immune cells, particularly B-cells, in exercise-induced neuroprotection, in order to determine the extent of their participation in ischemic tolerance.

5.1.7 Clinical Relevance of Exercise Preconditioning

In terms of clinical applications, exercise is one of the most promising forms of preconditioning-induced ischemic tolerance for individuals at a high risk for stroke [246]. Exercise is safe and noninvasive, without the risk of systemic inflammation or immunosuppression, as seen with other forms of preconditioning. It improves overall health, exerting a global anti-inflammatory effect [247]. Clinical studies already show that moderate exercise can reduce the risk of stroke [248, 249], improve vascular function, and diminish other risk factors such as obesity and

Table 2 Effects of exercise on the immune response after stroke

Paradigm	Stimuli	Animal	Injury model	Duration of protection	Immune-mediated results	Reference
Exercise	Forced treadmill running	Rat	tMCAo	48 h	Exercise increased TNF- α and ICAM-1 mRNA levels prior to stroke	[238]
					Exercise decreased leukocyte infiltration and decreased ICAM-1 expression after stroke	
	Forced treadmill running	Rat	tMCAo	48 h	Exercise decreased expression of TNFR1 and TNFR2 receptor expression in the brain	[234]
	Forced treadmill running	Rat	tMCAo	48 h	TNF- α and the ERK 1/2 pathway are necessary for decreased leukocyte infiltration after exercise and exercise-induced protection	[199]
	Forced treadmill running	Rat	tMCAo	48 h	TNF- α and the ERK 1/2 pathway are necessary for exercise-induced reduction of infarct volume, reduction of BBB disruption, reduction of MMP-9 protein, and increased collagen IV	[275]
	Forced treadmill running	Rat	tMCAo	48 h	Exercise reduced TLR-4 mRNA and protein expression in the brain	[274]

TNF- α tumor necrosis factor; *ICAM-1* intracellular adhesion molecule 1; *TNFR1* tumor necrosis factor receptor 1; *TNFR2* tumor necrosis factor receptor 2; *ERK 1/2* extracellular signal-regulated kinases 1 and 2; *MMP-9* matrix metalloproteinase 9; *TLR-4* toll-like receptor 4

high blood pressure [246]. In fact, individuals with active lifestyles prior to stroke have milder strokes, fewer motor deficits, and better functional recovery [250]. Determining the optimal intensity, frequency, and duration of exercise, and determining how exercise can beneficially modulate B-cell phenotypes in individuals, particularly with regard to co-morbidities, requires further research.

5.1.8 LPS Preconditioning

Activation of TLRs and TLR-triggered induction of regulatory factors are critical components of preconditioning, as TLR4-deficient mice were not protected against stroke by ischemic preconditioning [251]. In addition to being necessary for preconditioning-induced ischemic tolerance, systemic administration of ligands for TLR2, TLR4, TLR7, and TLR9 induce neuroprotection in rodent models of stroke [252–255]. Of the endotoxins that have been studied, the benefits of LPS preconditioning have been best characterized [256]. In mice, low systemic doses of LPS (< 1 mg/kg) prior to stroke produces an extended window of preconditioning, reducing infarct volumes for up to 1 week in mice, and 4 days in rats [253, 257]. Neuroprotection of neurons and endothelium significantly contributes to ischemic tolerance after LPS pretreatment. LPS administration 24 h prior to in vitro injury protects cortical neurons and cerebellar granular neurons [257, 258], as well as improves cerebrovascular functioning by reducing vascular permeability and increasing endothelial relaxation and microvascular reperfusion [259–261].

There is strong evidence that the efficacy of LPS preconditioning is largely due to immediate inflammatory responses that reprogram the immune system (Table 3). Within 24 h of LPS administration, there are significant genomic alterations associated with TLR signaling pathways [262]. These alterations have been associated with increased gene expression of anti-inflammatory cytokines and chemokines after stroke in preconditioned animals [263]. The initial LPS-induced increase in TNF- α is followed by a significant reduction at 1 day [257, 264]. Similar to exercise preconditioning, there was also a decrease in TNFR1 after stroke in preconditioned animals [257]. Activation of CCL2, a chemokine previously shown to be important in SHP (see Sect. 4.2.1), induces downstream effectors that reduce inflammation [265] and are required for efficacy of LPS preconditioning [38]. The involvement of TNF- α - and CCL2-mediated pathways in multiple paradigms of preconditioning supports the hypothesis that similar mechanisms may be responsible for preconditioning-induced ischemic tolerance. Finally, LPS preconditioning altered post-stroke leukocyte populations, decreasing neutrophils and monocytes in the brain while increasing circulating lymphocytes [254].

5.1.9 B-Cells in LPS Preconditioning

The role of B-cells in LPS preconditioning has unfortunately not been well-characterized. B-cells were decreased in the plasma after LPS preconditioning, though a similar phenomenon was seen in saline-treated mice, suggesting that loss of circulating B-cells may be subsequent to stroke as opposed to LPS preconditioning [254]. Rosenzweig and colleagues also found that LPS preconditioning did not significantly alter B-cell egress into the ischemic brain, as occurs after RHP [216]. However, this does not exclude the possibility of alterations of B-cell

Table 3 Effects of LPS pre-conditioning in the post-stroke response

Paradigm	Stimuli	Animal	Injury model	Duration of protection	Immune-mediated results	Reference
LPS	0.9 mg/kg, i.v.	Rat	pMCAo	1–4 days	TNF- α is necessary for LPS-induced reduction of infarct volumes	[253]
	0.05 mg/kg i.v.	Rat	pMCAo	24 h	LPS increases PMN infiltration after stroke	[259]
	0.2 mg/kg, i.p.	Mouse	tMCAo	48 h	LPS reduced microglia activation and neutrophil infiltration into the CNS	[254]
					LPS reduced monocytes and neutrophils, increased lymphocytes, and reduced monocyte activation in the blood	
	0.2 mg/kg, i.p.	Mouse	tMCAo	1–7 days	LPS increased pre-stroke TNF- α plasma levels but decreased TNF- α levels after stroke and decreased TNFR1 expression	[257]
			OGD	24 h	TNF- α is necessary for LPS-induced protection in vitro	
	0.2 or 0.8 mg/kg, i.p.	Mouse	tMCAo	72 h	LPS increased TGF- β , CCL5, and IFIT1 gene expression in the brain and IL-10 and CCL5 plasma levels	[263]
	0.2 mg/kg, i.p.	Mouse	tMCAo	72 h	LPS alters TLR signaling pathways and cytokine–cytokine receptor interaction pathways	[262]
	0.2 mg/kg, i.p.	Mouse	tMCAo	24 h	LPS increases MCIP1 mRNA and protein levels in the brain and is necessary for LPS-induced reduction of infarct volumes	[38]

PMN polymorphonuclear leukocytes; tMCAo transient middle cerebral artery occlusion; pMCAo permanent middle cerebral artery occlusion; TNF- α tumor necrosis factor; LPS lipopolysaccharide; CCL5 chemokine (C–C motif) ligand 5; CNS central nervous system; OGD oxygen glucose deprivation; TGF- β transforming growth factor β ; TLR toll-like receptor; PMN polymorphonuclear

phenotype, such as the induction of an immunosuppressive B-cell population by LPS or changes in function, including increased IL-10 production. In EAE, TLR stimulation by LPS induces a Breg population capable of reducing EAE severity [266] with sustained IL-10 production [23]. A proposed two-step model in EAE [266] may describe the role of B-cells in LPS preconditioning, as initial exposure to LPS induces a regulatory B-cell population that maintains an immunosuppressive phenotype via CD40 activation after stroke onset [267, 268], though this has yet to be confirmed.

5.1.10 Clinical Relevance of LPS Preconditioning

Currently, several TLR4 agonists are approved for human use to treat bladder carcinoma and HPV-associated cervical cancer [269]. TLR9 agonists have become increasingly attractive as stroke therapeutic, as they protect from stroke in nonhuman primates and show efficacy in clinical trials as adjuvants to vaccines to treat cancer, HIV, and allergies [270, 271]. Despite the promise of these trials, there are still significant concerns about the use of LPS in stroke patients. Clinical LPS administration is often used as a model of endotoxin tolerance because of the resultant acute inflammatory response followed by extended immunosuppression [272], a potential confounder that may not benefit long-term stroke recovery, considering prophylactic treatments with even low doses of LPS may induce endotoxin tolerance in stroke patients, worsening immunosuppression and increasing risk of infection [195].

6 Summary

This chapter reviews the current literature regarding the role of B-cells in stroke recovery, as well during ischemic tolerance to stroke induced by several preconditioning paradigms. B-cells predominantly mediate pathology in several autoimmune diseases, yet early evidence in stroke suggests that B-cells mediate both acute neuroprotection, as well as long-term pathology, in the injured CNS. What is most apparent from this review is that there is a lack of investigative studies linking B-cells directly to autoimmunity, neuronal plasticity, and overall functional recovery following stroke—in either experimental or clinical studies. While it is useful to review data from other CNS autoimmune diseases to predict potential mechanisms for B-cells during pathology, future studies should use the immunologic techniques established in MS, SLE, and RA models to determine how B-cells, particularly during an autoimmune response, shape the microenvironment within the ischemic CNS to support ischemic brain repair.

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Mast Cell as an Early Responder in Ischemic Brain Injury

Perttu J. Lindsberg, Olli S. Mattila, and Daniel Strbian

Abbreviations

BBB	blood–brain barrier
eNOS	endothelial nitric oxide synthase
IL	interleukin
LPS	lipopolysaccharide
MC	mast cell
MMP	matrix metalloproteinase
MCAO	middle cerebral artery occlusion
NVU	neurovascular unit
PAF	platelet-activating factor
PAI-1	plasminogen activator inhibitor-1
PAR	proteinase activated receptor 2
SCF	stem cell factor
TIMP	tissue inhibitor of metalloproteinases
tPA	tissue plasminogen activator
TNF α	tumor necrosis factor- α
WPBs	Weibel–Palade bodies
VWF	von Willebrand factor

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1 Introduction

Mast cells (MCs) are bone marrow derived, tissue-homing mononuclear white blood cells (leukocytes) that undergo differentiation upon entry into target organs. Committed bone marrow mast cell progenitors are released into the circulation from where they subsequently migrate into peripheral tissues, undergo maturation, and become terminally differentiated by the influence of cytokines within the surrounding milieu [1]. The progression of these cells to fully mature MCs is dependent on KIT activation which occurs as a consequence of stem cell factor (SCF)-induced KIT dimerization and autophosphorylation [1]. Central to this process is stem cell factor (SCF), the ligand for the c-kit tyrosine kinase receptor expressed on the surface of mast cells. KIT activation significantly regulates many aspects of MC differentiation, proliferation, activation, and survival [1, 2]. Mutant mice lacking either c-kit or SCF are MC deficient. MCs characteristically express a high-affinity receptor (FcεRI) for the Fc region of IgE, the least-abundant member of the antibodies. This receptor is of high affinity so that binding of IgE molecules is in essence irreversible.

MCs reside within or in the proximity to tissues that form a barrier to diverse noxious environmental challenges, i.e., in epithelium, mucosa, and the perivascular space. They are most commonly regarded as key effectors in the pathogenesis of allergic diseases and anaphylaxis [2]. However, the meticulous study of MC biology has established that they can generate or release various cytokines such as TNFα, potent proteases such as chymase and tryptase, vasoactive substances such as histamine and eicosanoids, and regulators of hemostasis such as heparin and tPA [3]. They also produce and release numerous interstitial matrix-degrading enzymes such as matrix metalloproteases (e.g., MMP-9). This armamentarium of mediators equips MCs to take part in a wide array of biological processes. This chapter describes recent evidence strongly supporting a role for MCs in the pathophysiology of cerebral ischemia.

2 Mast Cells in Health and Disease

MCs have gained a notorious reputation as mediators of anaphylaxis and allergic, atopic disorders. The spectrum of mast cell mediators, however, indicate a key role in diverse pathophysiological processes, such as chronic inflammatory processes, wound healing, angiogenesis, fibrosis, and tumors. Recent research has also demonstrated an important role in cardiac diseases, cardiac failure, cardiomyopathy, coronary syndromes, and atherosclerosis. Terminally differentiated, tissue-resident MCs are long lived [4], which coheres with a role in modulating local immunopathogenetic effects in chronic vascular disease. At least in the human the longevity of MCs is dependent upon the continued presence of SCF [1]. Clearly, MCs perform a critical role in protecting the host from a variety of environmental hazards [4, 5], but in the long-lived human, the cellular functions of MCs become involved in many

disorders with a strong autoimmune component. The role of MCs has been established in the pathophysiology of atopic dermatitis, psoriasis, asthma, and interstitial cystitis, and is highly probable in multiple sclerosis, coronary artery disease, irritable bowel disorder, and arthritis. The sphere of potential diseases where MCs and their mediators may play a role is, however, extensive and we must refer to other comprehensive reviews for more detailed scrutiny [6, 7].

3 Mechanisms of Action as an Inflammatory Cell

MCs are most efficiently activated via the classical IgE-mediated pathway but they can also be activated by a host of substances such as lipopolysaccharide, cytokines, hormones, immunoglobulins, neuropeptides (such as substance P), endothelin I, and activated complement components [8]. Functional activation of MCs leads to rapid degranulation of their mediators, many of which are preformed and stored in secretory granules. Piecemeal degranulation is an alternative form of mediator discharge permitting the selective secretion of certain mediators (and not the whole content) of the secretory granules. The preformed mediators include tryptase, histamine, serotonin (5-hydroxytryptamine), serine proteases, proteoglycans and cytokines such as tumor necrosis factor α (TNF α) and neutrophil chemokines CXCL1 and CXCL2 [9].

MC activation also leads to *de novo* synthesis of chemokines, cytokines, and eicosanoids that requires time for synthesis. For example, arachidonic acid metabolites (prostaglandins and leukotrienes), platelet activating factor (PAF), and several chemokines (e.g., CXCL1 and CXCL2) and cytokines (IL-1 β and IL-6) can be synthesized *de novo*. The balance of engaging inhibitory and activatory cell-surface receptors on MCs determines whether the cell becomes active on encountering a challenge. Once activated, MC's response is further regulated by the balance of both positive and negative intracellular molecular events that extend well afar from the conventional role of kinases and phosphatases [10, 11]. MCs are equipped with particular antigen presenting capacity and phagocytosis of microbes and antigens [2, 12], although their main mode of immunologic function is to act as local effector cells and attract other immune cells to the site of inflammation.

4 Cerebrocranial Mast Cells

The presence and activity of MCs within the cerebrum, already recognized by Nobel-prize winning immunologist Paul Ehrlich in the late nineteenth century, has received little attention up until now. MCs are not easily visible in standard histopathological stainings such as hematoxylin-eosin. They can be clearly detected with a handful of specialized staining procedures, many of which utilize visualization of the abundant heparin sulfate contained in the cytoplasmic MC granules. These can be demonstrated by metachromatic dyes like toluidine blue (Fig. 1a), azures A

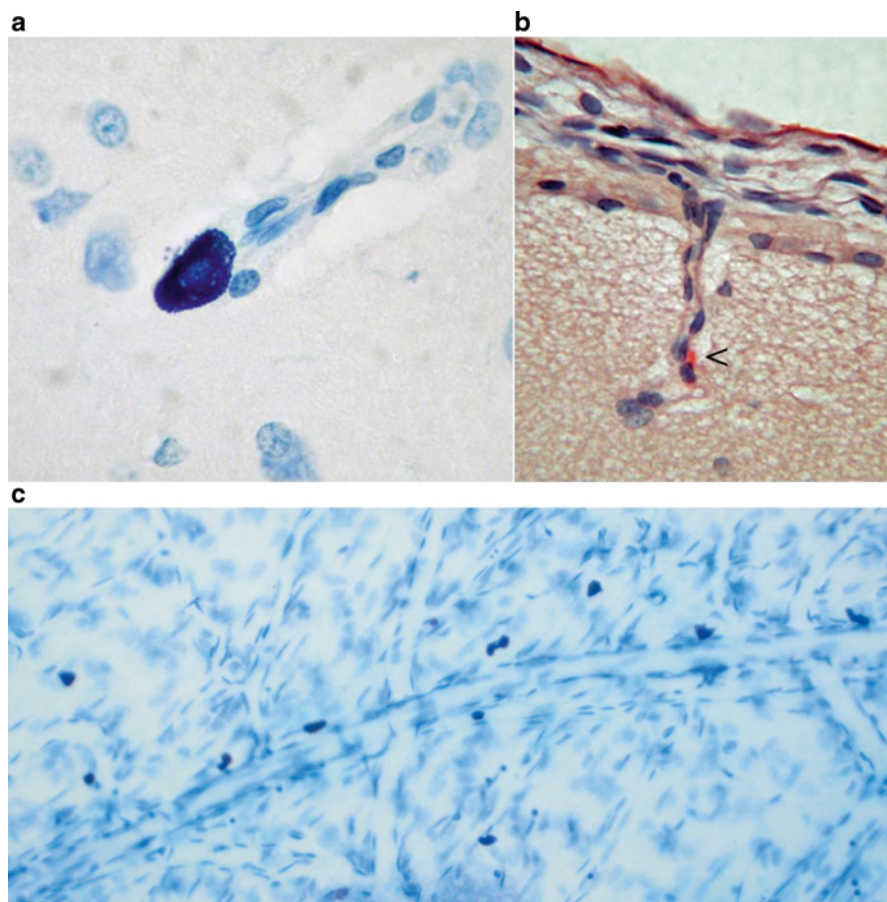


Fig. 1 Mast cells in cerebral and meningeal tissues. **(a)** Toluidine blue staining of a rat brain section demonstrating a densely packed granule-containing MC in association to a small blood vessel. A few solitary granules are also visible surrounding the outer cell membrane. **(b)** Demonstration of a rat brain MC with naphtol AS-D chloroacetate esterase (Leder) staining. A typical position close to a cortical penetrating arteriole can be observed. **(c)** Toluidine blue staining of mouse dura, showing the closely set localisation of MC in meningeal tissues

and B, thionin and methylene blue, which induce purple or violet staining of the granules. Toluidine blue and Giemsa are the most routinely used stainings. Egg white avidin attaches to heparin as well and has been used in a fluorophore-conjugated staining regimen. Tissue MCs can also be detected by enzyme cytochemistry by demonstration of Naphtol AS-D chloroacetate esterase activity (Leder staining) (Fig. 1b), by immunocytochemistry against tryptase, and by visualization of histamine fluorescence after *o*-phthaldialdehyde exposure [13].

Using these techniques, the presence of MCs has long been observed in substantial numbers in various craniocerebral structures. In the mammalian brain, MCs are found abundantly in the meninges, choroid plexus, hypothalamus, thalamus, olfactory bulb, and the midbrain [14–17]. MCs are also observed in the cerebral cortex

[14, 15, 18], characteristically at the branching points of cortical penetrating arterioles (Fig. 1b). A typical position for MCs is the Virchow–Robin perivascular space. They are situated in the brain parenchymal side of basal lamina, nesting between glial processes [19]. MCs are very abundant also in the dural meninges (Fig. 1c). From these locations, MCs can influence not only the blood vessels but interact also with neurons, glial cells, microglia, and extracellular matrix components. Intramuscular injection of compound 48/80 in several species has triggered striking Evans blue extravasation in MC-rich brain areas but not in MC-devoid regions equipped with fenestrated capillaries [20]. A tremendous challenge for research is that the type of MCs is to a large extent species-dependent, and the cell population fluctuates dynamically during development as well as in the course of behavioral and physiological events, and various kinds of stress.

The role of craniocerebral MCs remains elusive, but can be reflected upon an evolutionary perspective. MCs might have transformed from a cell which was directly involved in the compartmentalization and killing of pathogens into a cell type which orchestrates complex reactions leading to segregation and clearing of invading microbes. From a simple effector cell, the MC has become a coordinating cell which operates not only through innate mechanisms but also with the contribution of adaptive immune mechanisms [21]. It is of fundamental importance for the pathogenic mechanisms within the CNS, that of the resident cells in the brain, only MCs are able to acutely respond by releasing massive amounts of preformed mediators (reviewed in [22, 23]). The impact of MC activation is not uniformly proinflammatory, but can also attenuate the response of circulating leukocytes as in the mouse hippocampal brain trauma model [24].

In the brain, MCs have been ascribed a wide spectrum of physiological functions serving not only host response and innate immunity but also endocrine regulation [20]. MCs have been traditionally held as a first line of host defense against pathogens, allergens, toxins, and tissue injury, but with evolutionary accumulation of various membrane-bound receptors and diverse granule-contained mediators, MCs have engaged in quite diverse regulating functions also in many normal physiological and homeostatic functions along with the evolution of CNS.

5 Diversity of MCs

MCs of different organotypic origins show a substantial heterogeneity regarding their granule content and certain outer membrane epitopes. The organotypic difference has been used to classify MCs, which traditionally have been divided into mucosal or connective tissue type serosal (peritoneal) types. One key diversity is that of the main protease content: in humans, MCs can be distinguished based upon whether they contain tryptase (MCT) or tryptase and chymase (MCTC) [25]. Another example of diversity is the lack of c-kit receptor for SCF reported in some cerebral MCs [26]. Interestingly, blood transfusion results in MCs of donor origin settling in tissues of WT mice including the brain, and this relocation is restricted to

regions bearing host MCs, but transfusion in Kit^{W-sh/W-sh} MC-deficient mice strain results in MCs in the pinna of the ear, not in the brain [27]. Another brain tissue-specific MC characteristic is the low expression of the IgE-binding, high-affinity receptor FcεRI. The molecular and anatomical heterogeneity of the brain MCs has recently been reviewed in detail [20].

However, experimental studies have indicated that, in keeping with the perivascular position of MCs in the brain, the released vasoactive mediators furnish them with a capacity to regulate the permeability of BBB [28, 29].

6 MCs in the Priming of Inflammatory Cell Response

MCs are best known for their potent effector functions in allergic and hypersensitivity disorders. These phlogistic responses have long been known to be biphasic, the immediate phase occurring within minutes, and the late phase within several hours. The former is caused by the release of MC granule contents (histamine, TNFα, chymase, tryptase, heparin, cathepsin-G, and assorted chemokines). The second phase is characterized by *de novo* synthesis of MC mediators, such as leukotrienes, cytokines, prostaglandins, and additional chemotactic factors. However, MCs recently have been demonstrated to be involved in a vast array of considerably more complex immune functions that go well beyond allergies and contain the development of autoimmune disorders and peripheral tolerance, as well as the initiation and maintenance of adaptive and innate host responses [30].

In the development of local inflammatory tissue response, MCs have been implicated in recruiting neutrophils that are frequently the first immune cells to enter an inflamed or infected tissue site [31, 32]. In a mouse model of lipopolysaccharide (LPS)-induced peritonitis, MC granules containing preformed chemokines CXCL1 and CXCL2 were shown to be released by MCs within 15 min of *in vivo* stimulation, constituting an ideal mechanism to stimulate local neutrophil entry from the circulation at an early stage following the inflammatory signal [33]. Earlier work has emphasized the role of MC-granule dependent TNFα in mediating the initial phase of neutrophil recruitment following immune-complex mediated peritonitis in mice [34]. In addition, MCs, like macrophages, also were demonstrated to newly synthesize CXCL1 and CXCL2, making detectable amounts within 1 h of LPS treatment. Neutrophil extravasation was diminished in mice devoid of MCs [33].

Another facet of the local inflammatory response is the increase of vasopermeability and tissue edema. A well-established view is that several MC mediators, especially the vasoactive amines histamine, serotonin, and vascular endothelial growth factor A (VEGF-A), increase capillary leakage and can rapidly trigger tissue edema [35], but numerous other MC mediators such as PAF and leukotrienes participate in this response. Histamine receptor antagonists are therapeutically used to treat edema formation and allergic reactions associated with aberrant MC activity e.g., in pulmonary asthma, which can be also treated by the classic MC-stabilizing medicament, sodium cromoglycate. MCs can also trigger vascular permeability by paracrine mechanisms such as heparin-initiated bradykinin formation [36].

Although sparsely scattered in tissues, MCs are strategically positioned in close proximity to blood vessels and thus in a position to directly influence circulating leukocytes [33]. The same principle seems to be active in mounting local inflammation also in the various inflammatory diseases of the brain [6]. It is noteworthy that the described physiological and pathophysiological roles of MCs are likely to be strongly species-specific, because the immunological key characteristic such as the palette of granule constituents, membrane receptors, and activation modes have significant variability across different animal species.

7 Mast Cells as a Member of the Neurovascular Unit

The co-orchestrated roles of endothelial cells, neurons, astrocytes, pericytes, and extracellular matrix in the pathobiology of cerebral blood vessels have been encompassed under the term “neurovascular unit” (NVU). We and others have viewed MCs in the context of the NVU responding to cerebral ischemia [23, 37]. In the micromilieu of the brain parenchyma resident MCs lie in close proximity to other cell types of the NVU (Fig. 2). In addition to their role as initiators of acute inflammatory and vasoactive events, functional, unstressed MCs are certain to interact with these surrounding cell types, with bidirectional effects. Modes of interaction are likely to include direct cellular connections and released mediators, but interestingly, may also include cell signaling via release of exosomes, small membrane vesicles packed with RNA, capable of altering function of surrounding cell types [38]. For now, most research on cerebral MCs has focused on the pathophysiology of different neurological disorders, and more detailed investigation of MC interaction with other NVU cell types has been scarce, mostly relying on *in vitro* cell culture methods. Therefore the participation of MCs in the various physiological functions of the NVU is still to be characterized.

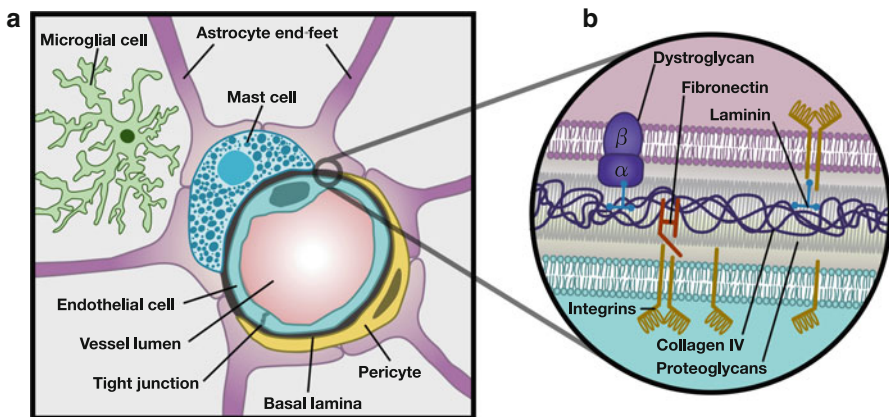


Fig. 2 Illustration of MCs at the NVU (a) and components of the basal lamina (b)

8 Evidence for a Role of MCs in Cerebral Ischemia

8.1 *Mast Cell Activation in Experimental Models of Cerebral Ischemia*

Initial observations of MC activation in experimental focal cerebral ischemia were made 15 years ago in our laboratory [39, 40]. In these experiments, degranulated cerebral MCs were observed in perivascular locations often associated with adjacent perivascular edema, based on which we hypothesized MCs to be involved in the pathophysiology of cerebral ischemia. Later on, MC activation and subsequent mediator release were reported in an *in vitro* model of oxygen-glucose deprivation [41]. In a separate study we further showed a dose-dependent activation of MCs (as measured by histamine release) *in vitro* as a reaction to tissue plasminogen activator (tPA), used for treatment of focal cerebral ischemia [42]. MC activation has also been described in neonatal rat stroke models. Biran et al. showed that ischemia induced histamine accumulation in the infarct core at 6–12 h after ischemia induction, whereas it was located in the penumbra at 24–48 h [43]. This study also showed that histamine accumulated in neuronal cells before they degenerated, which was accompanied by an increase in the MC count at 12 and 48 h. Activation of MCs after experimental hypoxic-ischemia injury in the neonate brain has later been replicated by additional studies [44].

8.2 *The Effects of MCs in Experimental Focal Ischemic Brain Damage*

Investigation on the role of MCs in focal cerebral ischemia began with experiments using a transient middle cerebral artery occlusion (MCAO) model in rats. Applying pharmacological activation and inhibition of MCs, we found MCs to play a central part in the development of space occupying brain edema, BBB disruption, and neutrophil infiltration 3 h after transient ischemia and reperfusion [18]. For example, MC stabilization with intraventricularly applied cromoglycate led to a 39% reduction in hemispheric expansion, 51% reduction in BBB leakage, and a 37% reduction in postischemic neutrophil infiltration. Involvement of MCs was subsequently studied using the same model together with intravenous tPA, simulating thrombolytic treatment [42]. MC stabilization showed reduction not only of edema, BBB disruption, and neutrophil infiltration, but also of perilous hemorrhage formation, neurological deterioration, and mortality. In both of these proof-of-concept studies, experiments were repeated using gene-manipulated MC-devoid rats lacking the *c-kit* ligand necessary for stem cell-dependent MC differentiation, demonstrating an even stronger protective effect with MC deficiency [18, 42].

More recently, our initial experimental findings in rats were replicated by two groups using a mouse MCAO model of transient ischemia. Studying later timepoints at 3 days and 2 weeks, Arac et al. showed significant reductions in brain edema, granulocyte infiltration, and infarct size in two different MC-deficient mouse strains [45]. Importantly, the study demonstrated the central participation of the meningeal MC population mediated via interleukin (IL)-6 and, to a lesser extent, chemokine ligand 7. Another study with pharmacological stabilization of MCs and genetically MC-deficient mice supported the involvement MCs in the pathophysiology of ischemia-mediated edema and inflammation, and suggested a role for endoglin, endothelin-1, and MMP-9, but not for TNF α [46]. Interestingly, this study also reported an increase of MC count by 50 % in the infarcted hemisphere 4 h after transient MCAO. Together, these observations provide solid evidence to support direct involvement of MCs in the pathophysiology of focal cerebral ischemia.

8.3 *MCs and Neuroprotection*

The initial *in vivo* stroke model experiments were not designed to study late-stage neuroprotection, and did not notice any effect regarding the lesion volume in association with MC stabilization or genotypic MC deficiency [18]. However, another study with longer follow-up of 24-h revealed a clear effect on functional recovery both after pharmacological stabilization of mast cells and in rats with mast cell deficiency [42], although no significant differences in infarct sizes were seen. More recent studies have also showed a reduction of infarct volume after MC stabilization in adult wild-type mice [45, 46]. Again, additional supporting evidence of MC involvement in tissue injury comes from neonate models. One set of experiments showed MC involvement in hypoxic-ischemic brain damage in the immature rat [47] and later observations showed that MC stabilization translated into reduced neuronal loss and brain atrophy [44]. The authors suggested the possibility of MC-derived IL-9 to be involved in the detrimental effect, which was supported by others [48]. Immunohistochemical co-localization studies of histamine and microtubule-associated protein 2 revealed accumulation in neuronal cells prior to their degeneration, and increased MC counts in the corresponding regions [43]. Histamine immunoreactivity was detected in MCs at 2, 6, and 12 h after ischemia, but disappeared at 24 h along with a concomitant observation of MC degranulation. Another study showed an effect of cyclosporin A in protecting against mild ischemic injury in neonatal rat brain [49]. The observed effect in reduction of histamine release from MCs is, however, IgE-dependent, and may not be of major importance in this setting. These data support a role for MCs in experimental ischemia-induced neuronal death, at least in neonatal cerebral ischemia.

9 Mast Cells and BBB in Ischemia–Reperfusion Injury

As the neurovascular unit responds to the sudden insult of ischemia and subsequent reperfusion, MCs are ideally located to initiate and aggravate known pathways of BBB disruption [23]. As discussed earlier, the activation of MCs is known to occur in a biphasic manner [3], starting with acute release of potent preformed granule contents that quickly spread to interact with the abluminal side of endothelial cells, the surrounding basal lamina, and other cell types of the neurovascular unit (Fig. 3). The second, later phase of MC activation is characterized by *de novo* production and release of mediators to support and prolong the initiated inflammatory response [3].

Very early on hypoxia, acidosis, formation of reactive oxygen species, and changes in blood flow act to disrupt cellular homeostasis of the neurovascular unit [37]. These reactions, together with intravascular blood coagulation, complement activation, and activation of the sympathetic nervous system, are likely initiators of MC activation next to parenchymal microvessels and within meningeal tissues [23].

The most imminent MC effects are amplified through the adjacent endothelium, which is the initial site of BBB leakage and failure early after reperfusion [50]. Histamine, an abundant and highly soluble MC mediator, acts through endothelial histamine receptors to activate calcium influx and convert the cells into a proinflammatory state [51]. The carefully characterized effects of histamine include increased endothelial permeability [51, 52], activation of endothelial nitric oxide synthase (eNOS) [53], and acute release of Weibel–Palade bodies (WPBs) [54], the main storage site for von Willebrand Factor (VWF) and P-selectin that act to support acute leukocyte infiltration [55, 56]. The MC protease tryptase may further activate endothelium through cleavage of the proteinase activated receptor 2 (PAR-2), with similar

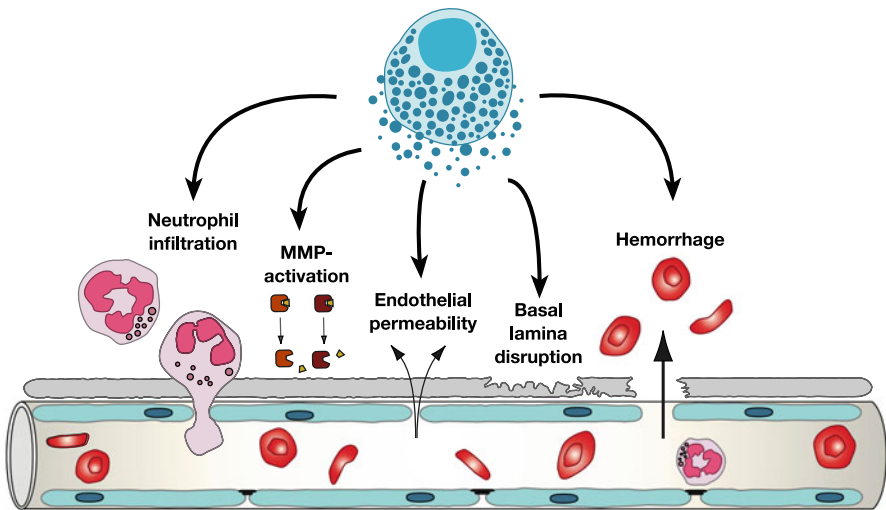


Fig. 3 Main pathways of MC-mediated blood–brain barrier disruption

effects to histamine [57]. A third MC mediator, Cathepsin G, has also been shown to induce endothelial permeability and influx of calcium into endothelial cells [58, 59].

As reperfusion injury advances, a storm of proteolytic enzymes is activated within the structures of the cerebral microvasculature [60, 61]. Ultrastructural evidence suggests that an early disruption of basal lamina [62], endothelial tight junctions [63], and other cellular connections ensues [64], and may locally progress to cause structural failure of the vascular wall, leading to edema and eventually hemorrhage [65]. Experimental evidence indicates that the wide armamentarium of MC mediators may have a central role in enhancing this proteolytic cascade [23].

The family of matrix metalloproteinase (MMP) enzymes, especially the gelatinases MMP-2 and -9, are thought to be a central proteolytic pathway, minutely characterized in experimental stroke models [61, 66–68]. For example, in human stroke patients plasma levels of MMP-9 are correlated with the incidence of significant hemorrhagic transformation [69, 70]. Using a rat MCAO stroke model, we found that both genetic MC deficiency and pharmacological MC stabilization with intracerebroventricular cromoglycate were able to significantly reduce the percentage of microvessels with high gelatinase activity in the ischemic hemisphere as early as 3 h after reperfusion (–64 % and –36 %, respectively) [71]. This finding is likely a sum of several MC-mediated effects on the MMP-cascade.

MCs have been shown to release both MMP-2 and -9 [72], which fits with the gelatinolytic activity we observed in the granules of activated cerebral MCs [71]. The MC protease chymase is capable of activating proMMP-1, proMMP-2, and proMMP-9 [72–74] and degrades tissue inhibitor of metalloproteinases (TIMP) -1, an important endogenous MMP-inhibitor [75]. MC tryptase has also been shown to activate proMMP-2 and proMMP-3 [76, 77]. Further on, *in vitro* studies have shown that histamine induces MMP-2 production in endothelial cells [78] and MMP-9 production in astrocytes [79]. MC proteases can also degrade components of the basal lamina directly: chymase is capable of degrading fibronectin [80] and cathepsin G, found in a subset of MCs, is able to degrade fibronectin and laminin [81, 82].

Progression of postischemic BBB disruption is accompanied by unrestrained granulocyte infiltration, beginning hours after reperfusion [83], which acts to further drive inflammation, increase proteolysis and barrier permeability [84], and disrupt microvascular flow [85]. MCs seem to have a central role in activating leukocyte recruitment, as data from three individual laboratories show that MC inhibition significantly reduces both early and late granulocyte infiltration after transient MCAO (3–6 h and 3 days postreperfusion, respectively) [18, 42, 45, 46].

MCs secrete a wide range of mediators that can augment granulocyte infiltration. In addition to the endothelial dependent effects of histamine and tryptase described above, MCs are capable of releasing preformed TNF α , which further increases endothelial permeability, endothelial adhesion molecule expression, and neutrophil infiltration [86–88]. Moreover, chymase is thought to have direct chemotactic effects on neutrophils [89]. As MC activation endures, *de novo* production of mediators continues to support infiltration of granulocytes. IL-1 is capable of increasing both endothelial barrier permeability and neutrophil infiltration [90, 91]. Further, in a recent report, Arac et al. demonstrated that IL-6 is central for MC-dependent neutrophil infiltration

in a later phase, 3 days postreperfusion. In these experiments reconstitution of MC-deficient mice with wild-type MCs returned typical neutrophil infiltration and brain swelling, while reconstitution with IL-6-deficient MCs did not [45].

To sum up, the effects of MCs in experimental models of ischemic stroke are well in line with the known effects of the wide armamentarium of MC mediators. However, the individual contribution of these mediators is still unknown, and will require further experimental work, preferably by reconstituting MC-deficient mice with MCs deficient for the studied mediator [45]. Although meningeal MCs appear to be central at later timepoints after reperfusion (3 days and 2 weeks) [45], the relative contribution of different MC populations in the ultra-acute and early phases (0–48 h) of ischemic stroke is still unknown.

10 Mast Cells, Blood Coagulation, and Fibrinolysis

In addition to their potent vasoactive, proteolytic, and chemotactic effects, MCs are known as a profibrinolytic, anticoagulant, and antithrombotic cell type, with several effects on thrombotic pathways [23, 92]. As reperfusion injury advances to break down structures of the vascular wall, the unphysiologically strong activation of MCs may partake in initiating dangerous intraparenchymal hemorrhage, one of the most feared complications of acute stroke treatment. Supporting this hypothesis, in experiments using a rat stroke model with 90 min of transient MCAO combined with intravenous tPA, both genetic and pharmacological MC inhibition led to almost total abrogation of intraparenchymal hemorrhage [42]. Again, a spectrum of MC mediators have effects on hemostasis.

Heparin is the central anticoagulant mediator released from MCs, a negatively charged glycosaminoglycan which catalyzes antithrombin III-mediated inactivation of coagulation factors, most importantly activated factor X and thrombin [93]. Heparin can inhibit binding of platelets onto collagen IV [94], revealed upon disruption of the vascular wall. Further, heparin also releases tissue factor pathway inhibitor (TFPI) from the surface of endothelial cells, a mediator which can inhibit arterial thrombosis [95]. Lastly, heparin has recently been shown to activate the plasma contact system, inducing rapid generation of bradykinin without activation of blood coagulation [36], which may act to further increase endothelial permeability and leukocyte infiltration.

MCs also have more direct effects on fibrinogen and fibrin. MC tryptase has been shown to degrade fibrinogen, preventing normal fibrin formation [96, 97]. Tryptase also activates pro-urokinase [98], an important plasminogen activator, initiating plasmin-mediated breakdown of fibrin. Moreover, MCs have been shown to directly secrete tissue plasminogen activator (tPA), another central plasminogen activator, without accompanying secretion of plasminogen activator inhibitors, like plasminogen activator inhibitor-1 (PAI-1) [99]. Importantly, a wide collection of experimental evidence has shown that plasminogen activators, especially tPA, have important effects on proteolysis and inflammation at the BBB, in addition to direct

pro-excitotoxic effects [100]. Of note, a positive feedback loop may exist between fibrinolysis and further MC activation, as certain fibrinolytic breakdown products of fibrinogen have been shown to activate MCs [101].

The endothelial effects of MC mediators contribute an additional pathway for modification of hemostasis. In patients with anaphylaxis, extremely strong MC activation and subsequent release of endothelial WPBs have been shown to induce a rapid increase in circulating levels of both vWF and tPA, and induce systemic plasminogen activation [102]. The significance of this pathway during localized MC activation is still to be uncovered, but may have both fibrinolytic and proaggregatory effects.

In the setting of acute inflammation, the physiological purpose of these described anticoagulant, fibrinolytic, and antithrombotic MC effects may be in regulating thrombosis activated by inflammatory pathways, to ensure adequate blood flow to the inflamed tissue area, and counteract the inhibitory effects of fibrin formation on leukocyte recruitment [103]. More generally, in the resting state, MCs have been suggested to protect the brain microvasculature against thrombotic challenges [104]. In line with these hypotheses, several products of the coagulation cascade have been shown to activate MCs, including bradykinin, thrombin, and activated factor X [105–107].

11 Conclusions

To conclude, recent experimental research suggests that MCs and their nominal responses increasing the permeability of the vascular wall play a significant, deleterious role following acute cerebral ischemia. MCs should be regarded as a potent inflammatory cell type that can interact via a multitude of mediators and signalosomes with its neighboring cells within the NVU, and may also have more distant effects within the CNS. MCs are a unique resident inflammatory cell type, settled in the proximity of the vascular wall already at the outset of ischemia, capable of quickly degranulating, leading to degradation of the basal membrane, BBB damage, brain edema, and hemorrhage.

Future goals of MC research include examination of whether MC mediators are released early after cerebral ischemia in man. In view of the significant species differences in the immunological characteristics and mediator contents of MCs, more evidence is needed on the magnitude of MC-dependent chemokine release, neutrophil targeting, and other secondary effects on postischemic tissue integrity. The potential involvement of the meningeal MC population is an attractive area of future research, as these MCs may be more readily influenced with pharmacological means due to their localization outside the BBB. MCs are candidate cells to become novel pharmacologic targets at the NVU to limit ischemic and hemorrhagic brain damage associated with reperfusion therapies.

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Roles of Neutrophils in Stroke

Glen C. Jickling and Frank R. Sharp

1 Introduction

Neutrophils are among the first immune cells to respond to ischemic brain, and contribute to processes central to the pathogenesis of ischemic stroke including clot formation and atherosclerosis. Neutrophils have been implicated in worsening outcomes in several diseases including ischemic stroke, myocardial infarction, diabetic retinopathy, sickle cell disease, transfusion related acute lung injury, acute respiratory distress syndrome, and renal microvasculopathy [1].

Many of the neutrophil functions designed to kill pathogens also influence a variety of processes that can initiate or promote cerebral infarction. During the process of phagocytosis neutrophils produce Reactive Oxygen Species (ROS) such as superoxide and hypochlorous acid via NADPH oxidase and Myeloperoxidase (MPO), respectively, [2] which can directly injure the Blood Brain Barrier (BBB) and possibly brain cells. During the process of degranulation, normally designed to kill pathogens, neutrophils release proteases including MMPs (-1, -2, -8, -9, -13), elastase, cathepsin G, and Proteinase 3 that can degrade extracellular matrix and damage the BBB and brain cells. Neutrophils can also release or respond to pro-inflammatory cytokines (e.g., IL-1 β , IL6, IL-8, TNF- α) and chemokines (MCP-1/CCL2, MIP-1alpha/CCL3, CCL5/CCR5) that could worsen ischemic brain injury [3]. A variety of neutrophil receptors interact with damaged endothelium and with platelets to promote clotting/thrombosis [4, 5] and promote atherosclerosis [5, 6] (Fig. 1). The current status of research in these areas will be summarized here.

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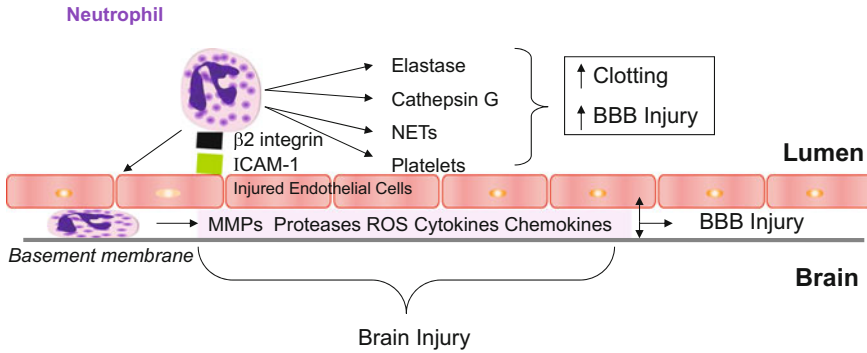


Fig. 1 Neutrophil activation and adhesion in acute ischemic stroke. Following ischemic stroke cytokines and damage associated molecular patterns are released. These promote neutrophil recruitment and activation, including the release of reactive oxygen species, proteases, cytokines. Neutrophils adhere to activated endothelium through adhesion molecules that promote neutrophil–endothelial interactions and neutrophil migration with resulting effects on the blood brain barrier and brain parenchyma (Abbreviations: *CCL*, CC-chemokine ligand; *CXCL*, CXC-chemokine ligand; *CCR*, CC-chemokine receptor; *CXCR*, CXC chemokine receptor; *DAMPs*, damage associated molecular patterns; *IL*, interleukin; *ILR*, interleukin receptor; *FPR*, formyl peptide receptor; *HMGB1*, high mobility group box 1; *HSP72*, heat shock protein 72; *ICAM-1*, intracellular adhesion molecule-1; *MAC-1*, macrophage 1 antigen; *MMP-9*, matrix metalloproteinase 9; *PSGL-1*, P-selectin glycoprotein ligand-1; *TLR*, toll-like receptor; *TNF α* , tumor necrosis factor alpha)

In assessing the role of neutrophils in acute ischemic stroke, this review focuses on the first few days following stroke. This is in part because most human and animal infarct volumes are stable by 24 h and do not change greatly thereafter for many days. In addition, the majority of the animal studies have focused on this early time period presumably for similar reasons. This focus, however, is not meant to say neutrophils do not have an impact on the long-term outcomes after stroke. As discussed shortly, human studies point to neutrophil numbers being extremely important in all types of cardiovascular and cerebrovascular disease in so far as risk of disease, severity of disease, and prognosis/outcomes. In addition, neutrophils have important roles prior to stroke since they promote clotting and atherosclerosis as noted below.

It is notable that several large clinical trials aimed at interfering with leukocyte/neutrophil function have failed in human stroke [7–9]. This is an important reminder that the roles of neutrophils in animal stroke models may or may not recapitulate the role of neutrophils in human stroke. Indeed, 50–70% of leukocytes in humans are neutrophils, whereas only 10–25% of rodent leukocytes are neutrophils [2]. Humans are more susceptible to infection than rodents, and infection clearly worsens stroke outcomes [10, 11]. Thus, suppression of neutrophil function could be more problematic in human than rodent stroke. Thus, identifying neutrophil targets most likely to benefit stroke outcome without affecting anti-microbial functions of neutrophils is likely important and is discussed where information is available.

The failure of anti-neutrophil therapies in humans has also raised the question of whether leukocytes and neutrophils in particular are a pathological variable in

ischemic stroke or simply a bystander reaction to damaged tissue not actually causing damage [12]. Many experimental studies do not demonstrate a causal role for neutrophils in brain injury following ischemic stroke [12]. For instance, demonstrating a change in infiltration of neutrophils or neutrophil markers with a given treatment does not demonstrate the neutrophils contributed to the injury. Even when a causal role is demonstrated, it is often not clear whether this would be relevant for one or all three major causes of stroke in humans: large vessel, cardioembolic, and small vessel/lacunar strokes. In addition, neutrophils appear to play a lesser role in stroke models where there is little or no reperfusion which may complicate translation to humans, and might limit treatments where there is probably little or no reperfusion in ischemic stroke. This chapter will consider some of these issues, and attempt to determine if there are any specific functions or molecules ascribed to neutrophils that might be causative in human stroke, not adversely affect anti-pathogen roles of neutrophils, and represent reasonable treatment targets.

2 Brief Biology of Pathogen Function of Neutrophils

Neutrophils are terminally differentiated cells that develop in the bone marrow [2, 5]). Once released into the circulation neutrophils seek signs of infection and inflammation via local cytokines, chemokines or damage associated molecular patterns (DAMPs) or PAMPs. Infectious or inflammatory activation of neutrophils leads to adhesion to vessel walls and migration inside tissue to the site of infection or inflammation. Notably, blood-derived neutrophils found in ischemic brain often show that they have just proliferated (are BrdU positive) [13].

When an infectious particle is encountered by neutrophils, it is phagocytosed. NADPH oxidase and MPO generate ROS, and proteolytic contents of neutrophil intracellular granules are released into the phagosomes to kill and digest the microorganisms [5]. Neutrophils have been found to form DNA based neutrophil extracellular traps (NETs) (described below) that immobilize pathogens to prevent spread and to also directly kill pathogens [2]. The neutrophils generally die via apoptosis within a day or two, though infection and inflammation can increase neutrophil survival. Once neutrophils die, macrophage removal of dead neutrophils is initiated [5] via a IL-23, IL-17, G-CSF pathway [2]. Besides this traditional role in infection, neutrophils also play a role in noninfectious inflammatory processes including ischemic stroke and heart disease. Evidence for this in humans has come in part from neutrophil responses in peripheral blood of patients with stroke.

3 Neutrophil Response in Ischemic Stroke

In patients with ischemic stroke, the number of circulating neutrophils rises within the first few hours of stroke onset [14] (Figs. 1 and 2). This increase is associated with stroke severity [15], infarct volume [16], and worse functional outcomes [17].

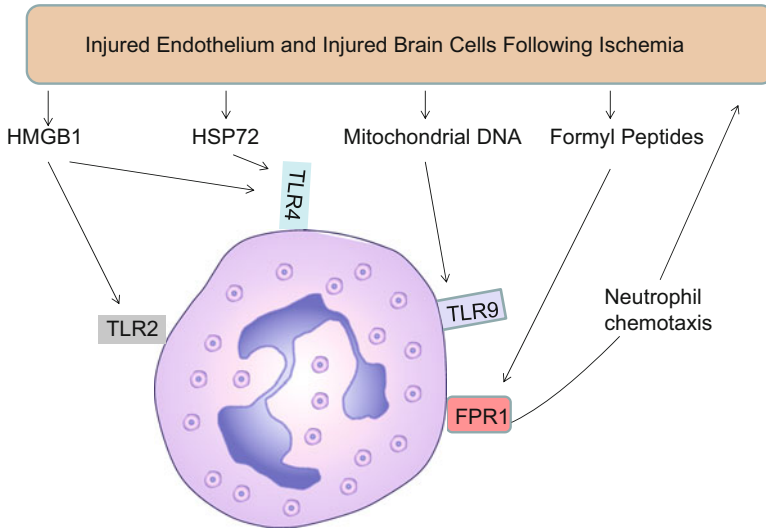


Fig. 2 Ischemic brain injury results in the release in a number of factors that activate circulating neutrophils. These include HMGB1, HSP72, mitochondrial DNA, and formyl peptides. Activated neutrophils are recruited to site of tissue injury

In contrast to neutrophils, lymphocytes decrease following ischemic stroke. Thus, the neutrophil to lymphocytes ratio is increased following stroke, and is associated with mortality and infarct size [18].

The rise in neutrophils following stroke occurs as a result of enhanced production, increased release from the bone marrow and spleen, and possibly from a reduction in neutrophil apoptosis [19]. Neutrophils express several endothelial adhesion molecules (PSGL-1, ESL-1, CD44, LFA-1, MAC-1) within 15 min of ischemia (Fig. 3). By 2 h, neutrophil rolling and adhesion is present in the pial vessels of the brain [20–22]. After 6–8 h, neutrophils have surrounded cerebral vessels and infiltration has begun [23, 24]. By 24–48 h of ischemic stroke, neutrophil infiltration into brain has peaked [22, 25]. The increase in neutrophils following ischemic stroke is associated with increased expression of adhesion molecules, cytokines/chemokines, proteases, and ROS.

A preponderance of data to date suggest neutrophil pro-inflammatory activation post stroke is associated with increased infarct size, increased BBB disruption, hemorrhagic transformation, and worse neurological outcomes. These studies are summarized below. However, often studies of neutrophils in stroke have used markers that are not specific to neutrophils and can be found on other immune cells. For example, MPO is often interpreted as a marker of neutrophils, however is also expressed by monocytes/macrophages and microglia. Thus when evaluating the role of neutrophils in stroke it is important to be cognizant that reported associations of neutrophils may also reflect a potential contribution from other immune cells.

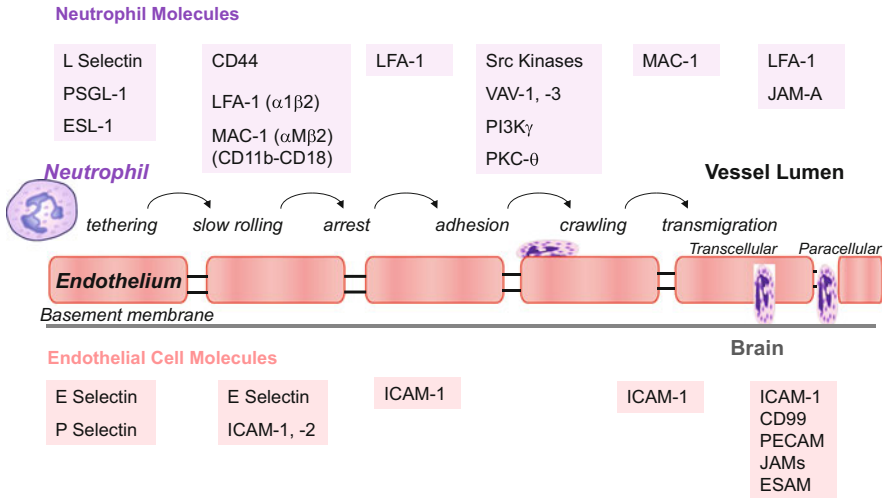


Fig. 3 Neutrophil recruitment, adhesion, and transmigration. Neutrophils express specific adhesion molecules that bind activated endothelium to promote adhesion and migration. In ischemic stroke the adhesion molecules expressed on neutrophils and endothelium involved in neutrophil tethering, rolling, arrest, adhesion, crawling, and transmigration remain poorly defined and may differ from the peripheral vasculature. Further studies are required to delineate the molecules involved in the neutrophil recruitment cascade following ischemic stroke and determine the extent of transmigration. (Abbreviations: “?” indicates unclear role in ischemic stroke; *ICAM-1*, intracellular adhesion molecule-1; *LFA-1*, lymphocyte function-associated antigen 1; *MAC-1*, macrophage 1 antigen; *PSGL-1*, P-selectin glycoprotein ligand-1)

4 Neutrophils and Thrombosis

Neutrophils are important contributors to clot formation (Fig. 4). When neutrophils are depleted, thrombus formation is reduced [26]. A neutrophil promotes thrombosis through several mechanisms including formation of tissue factor, interactions with platelets, release of NETs, and release of proteases that act on coagulation factors [26, 27].

Tissue factor interacts with coagulation factor VIIa to initiate activation of the extrinsic coagulation pathway and promote thrombus formation. Activated neutrophils are an important source of tissue factor [26, 27]. Whether neutrophil-derived tissue factor promotes thrombosis in ischemic stroke requires further study. Tissue factor is increased in rheumatoid arthritis, systemic lupus erythematosus, and Crohn’s disease, each of which is associated with ischemic stroke. Neutrophil-derived tissue factor may contribute to this increased stroke risk.

Neutrophils have a number of interactions with platelets that result in enhanced platelet aggregation and clot formation [28]. Ligands expressed by neutrophils include PSGL-1 that binds platelet P-selectin, MAC-1 (CD11b-CD18, $\alpha M\beta 2$) that binds platelet GPIb α , and binds a fibrin-glycoprotein IIb/IIa (gp IIb/IIIa, $\alpha IIb\beta 3$ integrin)

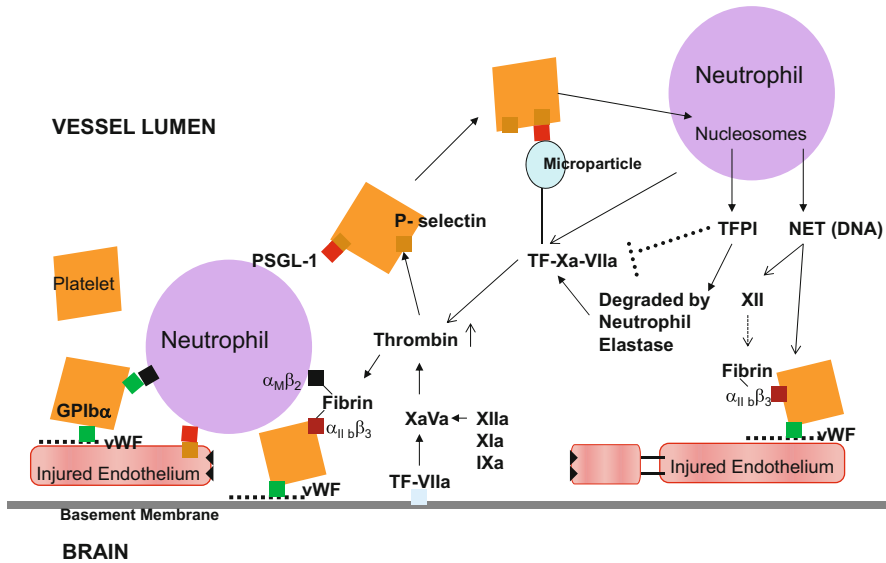


Fig. 4 Role of neutrophils in thrombus formation. Neutrophils promote thrombosis through interactions with platelets, proteolytic cleavage of clotting factors (TFPI, coagulation factor X), and release of prothrombotic molecules (neutrophil extracellular traps, tissue factor) (Abbreviations: *ICAM-1*, intracellular adhesion molecule-1; *MAC-1*, macrophage 1 antigen; *NET*, neutrophil extracellular traps; *PSGL-1*, P-selectin glycoprotein ligand-1; *TF*, tissue factor; *TFPI*, tissue factor pathway inhibitor; *vWF*, von Willebrand factor)

complex (Fig. 4). Platelets also release soluble CD40L that stimulate neutrophil expression of MAC-1. In ischemic stroke, neutrophil–platelet interactions may be important to thrombus formation and vessel occlusion. In patients with recent ischemic stroke, neutrophil–platelet complexes are increased [29]. The stroke prevention therapies dipyridamole and candesartan both inhibit neutrophil expression of adhesion molecules which prevent thrombus formation [30]. Abciximab and eptifibatid act on the gpIIbIIIa receptor, which mediates neutrophil–platelet interactions. Clopidogrel blocks platelet activation and resultant conformation change in gpIIbIIIa, and thus also prevents neutrophil–platelet interactions [31]. In sickle cell disease, ischemic stroke may result from altered neutrophil–platelet interactions and enhanced neutrophil-dependent platelet aggregation [32].

Neutrophil-derived proteases contribute to thrombus formation. Cathepsin G and elastase act on coagulation factor X to promote coagulation [33]. Elastase also degrades tissue factor pathway inhibitor alpha (TFPI α), which increases levels of tissue factor that promote clot formation [33]. Neutrophil cathepsin G plays a role in neutrophil–platelet interactions [34]. Inhibiting cathepsin G reduces bleeding time and has greater antithrombotic effect than aspirin. In rodent stroke, inhibiting cathepsin G improves cerebral blood flow, reduces brain injury, and improves behavioral outcomes [34]. Neutrophils also release the protease ADAMTS13, which cleaves hyperactive ultra-large von Willebrand factor (ULVWF) and affects

thrombosis in ischemic stroke [35]. Thus, targeting neutrophil proteases may have potential as novel antithrombotic therapies to prevent ischemic stroke [34].

NETs may also contribute to thrombus formation in ischemic stroke [36]. NETs are derived from neutrophils and composed primarily of DNA. Though they typically bind pathogens, NETs can trigger platelet activation and promote thrombus formation [37]. Blocking NET formation with DNase reduces clot formation [38]. NETs have been implicated in deep venous thrombosis but as yet not in ischemic stroke [36]. Neutrophils also release DNA–histone complexes that not only trap and eliminate pathogens but also promote thrombus formation. The thrombogenic potential of NETs is supported by the finding that DNase inhibits thrombus formation related to DNA–histone complexes [39].

5 Neutrophils in Atherosclerosis

Atherosclerosis is a major cause of ischemic stroke occurring in both the extracranial and intracranial vasculature that supplies the brain. Neutrophils contribute both to the formation of atherosclerosis and to the rupture of plaque that causes thrombosis and brain ischemia [40] (Fig. 5).

In atherosclerosis, neutrophils are recruited early to sites of endothelial injury through cytokines, chemokines, and adhesion molecules (Fig. 5). In mice deficient in the adhesion molecules P-selectin (platelets) [41], CD18 (neutrophils, monocytes), or ICAM-1 (endothelial cells), formation of atherosclerotic plaque is reduced [42]. P-selectin is important for platelet deposition in plaques. Platelets secrete CCL5 which acts on neutrophil CCR5 receptor to promote neutrophil recruitment to injured endothelium [43]. Neutrophils adhere to endothelium through CD18 and ICAM-1. In turn, recruited neutrophils promote monocyte recruitment via CRAMP (cathelicidin antimicrobial peptide) [5]. In mice, deletion of CRAMP or depletion of neutrophils reduces atherosclerosis [44, 45].

Chemokines and cytokines may be potential targets to reduce atherosclerosis in stroke (Fig. 5). Evasin-3 is a CXC chemokine-binding protein that inhibits neutrophil activation. When administered in rodent models, carotid atherosclerosis is decreased, intraplaque neutrophil content is reduced, and matrix metalloproteinase-9 (MMP-9) activity is diminished [46]. The beneficial effects of increased HDL (high-density lipoprotein) on atherosclerosis may be mediated in part through neutrophils. Increasing HDL decreases neutrophil activation by proinflammatory cytokines (TNF- α , IL-1, and IL-8) [6].

Neutrophils also contribute to atherosclerosis by promoting the formation of oxidized lipids (Fig. 5). Neutrophil NADPH oxidase and myeloperoxidase (MPO) are a major source of ROS that promote formation of oxidized low density lipoproteins (oxLDL). oxLDL are taken up by scavenger macrophages to form foam cells in atherosclerotic plaques (Fig. 5). Neutrophil-derived ROS also contribute to vessel injury and endothelial dysfunction in atherosclerosis, cause vascular smooth muscle proliferation, and activate MMPs which contribute to plaque rupture [6, 47].

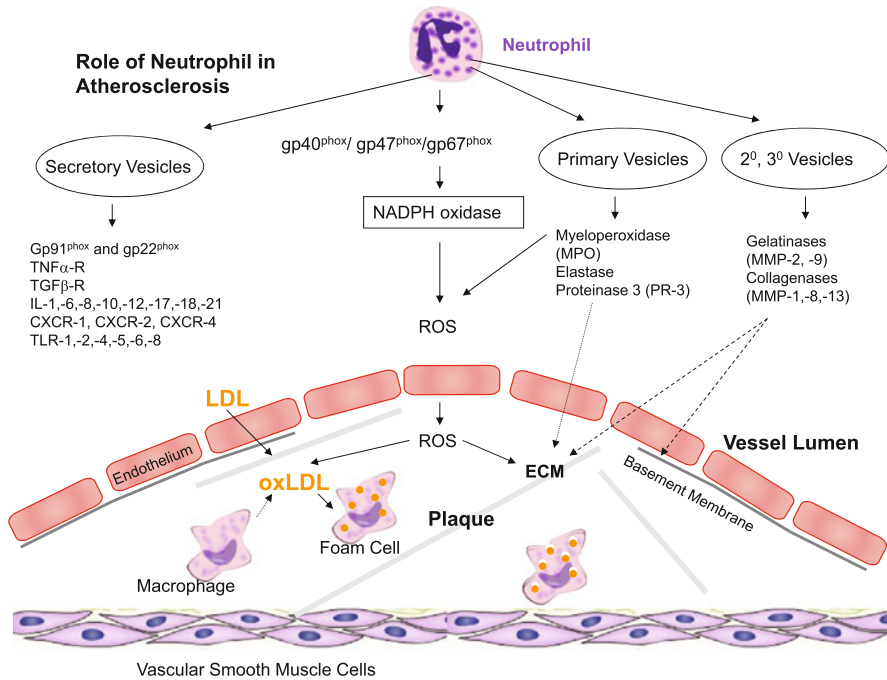


Fig. 5 Role of neutrophils in atherosclerosis. Neutrophils promote the formation of atherosclerosis through interactions with platelets and by enhancing monocyte infiltration into damaged endothelium. Neutrophils promote atherosclerotic plaque progression and rupture via the release of cytokines, reactive oxygen species that activated macrophage foam cells, producing oxidized-LDL, and proteolytic degradation of the fibrous cap (Abbreviations: *BM*, basement membrane; *CCL*, CC-chemokine ligand; *CRAMP*, cathelicidin antimicrobial peptide; *ECM*, extracellular matrix; *FPR*, formyl peptic receptor; *ICAM-1*, intracellular adhesion molecule-1; *IL*, interleukin; *INF- γ* , interferon gamma; *LL37*, Cathelicidin; *MAC-1*, macrophage 1 antigen; *MMP*, matrix metalloproteinase; *ox-LDL*, oxidized low density lipoprotein; *PSGL-1*, P-selectin glycoprotein ligand-1; *TLR*, toll-like receptor; *TNF α* , tumor necrosis factor alpha)

The beneficial effect of statins in atherosclerosis may relate in part to effects on neutrophils. Statins decrease the production of ROS by inhibiting neutrophil NADPH oxidase, as well as reducing neutrophil activation, adhesion molecule expression, and platelet interactions [6].

Proteases released by neutrophils also contribute to atherosclerosis (Fig. 5). Neutrophil-derived extracellular matrix-degrading proteases can promote endothelial dysfunction by degrading vascular basement membrane and type IV collagen. In turn, dysfunctional endothelium promotes additional neutrophil recruitment through increased expression of adhesion molecules (β 2 integrins, P-selectin, E-selectin, ICAM-1) and IL-8. Neutrophils are an important source of MMPs which contribute to atherosclerosis. MMP-1, MMP-8, and MMP-13 have been shown to promote atherosclerosis in mice [48, 49] and to be overexpressed in human atherosclerotic plaque [50]. MMP-8 has also been associated with plaque instability [50], and genetic polymorphism in MMP-8 is related to progression of atherosclerosis

and plasma levels of the adhesion molecule VCAM-1 [49]. Neutrophils are also a primary source of MMP-9. MMP-9 promotes plaque instability and rupture through effects on extracellular matrix. Treatment with statins increases plaque stability in part by reducing neutrophil MMP-9 expression and neutrophil infiltration [51].

In humans, atherosclerotic plaque instability correlates with the presence of intra-plaque neutrophils [52]. Progression of aortic arch atheroma is associated with increased neutrophils in addition to increased risk of stroke and myocardial infarction [53]. In humans, an increase in neutrophil counts is associated with hypoechoic unstable carotid plaques, symptomatic carotid artery stenosis, and cerebral micro-embolization [54].

6 Neutrophil Response in Acute Ischemic Stroke: Humans

Neutrophils respond acutely to ischemic stroke, and remain activated for days to weeks (Fig. 1). CD11b expression increases on neutrophils by 15 min following experimental strokes and remains elevated for over 24 h [55]. Deep hypothermic cardiac arrest in humans immediately increases counts of activated neutrophils (CD11) [56]. After permanent middle cerebral artery occlusion (MCAO) in rodents, there is significant rolling and adherence of leukocytes in pial venules and arterioles by 2 h [20]. In rodent permanent MCAO models, neutrophil infiltration was moderate and typically within and adjacent to blood vessels bordering the infarcted cortex whereas temporary MCAO with reperfusion resulted in neutrophil infiltration throughout the infarcted cortex [57].

Labeled polymorphonuclear leukocytes (neutrophils) can be detected in the ischemic hemisphere by 1 h following cerebral ischemia in the dog [21]. Intravascular neutrophils in the rodent ischemic hemisphere peak around 6 h [22]. Neutrophil “invasion” into ischemic brain begins between 6 and 8 h after reperfusion in rodent stroke models [23, 24] (Fig. 1) and can peak in rodent brain as early as 24 h after MCAO [22, 25]. Cytokine-induced neutrophil chemoattractant (CINC) mRNA is increased by 6 h post MCAO, peaks at 12 h, and decreases but is still elevated at 24 h in a rat stroke model [58]. Animal studies that have tracked myeloperoxidase (MPO) find neutrophil numbers peak between 1 and 5 days following stroke and decrease thereafter whereas monocytes peak later between 3 and 7 days [24, 59–61]. One study disputes these time lines, finding that macrophages, lymphocytes, and dendritic cells enter cerebral infarcts prior to neutrophil entry (based upon flow cytometry), though equal numbers of neutrophils and activated microglia were observed at day 3 [62].

The inflammatory response post stroke can be quite prolonged, taking 12 months for fibrinogen and CRP levels to return to baseline levels [63]. IL-1ra, produced in part by neutrophils [64], takes 6 months to return to baseline following stroke whereas IL-6 levels, produced in part by neutrophils, return to baseline around 8 weeks after ischemic stroke [63]. Direct assessment of neutrophils in human stroke brain over time shows three groups: severe neutrophil accumulation that dramatically increased within 12 h after stroke onset and persisted beyond 30 days; those with moderate neutrophil accumulation that significantly decreased by 30 days; those with mild

neutrophil accumulation that decreased by 6–9 days [65]. In patients with severe neutrophil accumulation, neurological outcome was worse and infarction volume larger than in patients with less marked accumulation—suggesting the magnitude of the neutrophil response correlates with the magnitude of the brain injury [65]. Neutrophil numbers decrease in part after stroke because of apoptosis and engulfment by brain microglia [66] and engulfment by macrophages [60], and because of decreased release from bone marrow.

7 Neutrophil Biomarkers in Blood Following Acute Stroke: Humans

Additional evidence for neutrophil responses to acute stroke in humans comes from measurements of neutrophil biomarkers in serum and plasma. PSGL-1, P-selectin glycoprotein ligand 1 which binds to P-selectin on platelets and endothelial cells, showed increased expression on neutrophils and monocytes from day 1 to 90 after stroke as compared with control subjects ($p < 0.05$) [67]. Neutrophil Mac-1 expression was significantly increased on days 1 and 7 after stroke compared to controls ($p < 0.05$). Neutrophil PSGL-1 expression on day 1 was significantly higher in patients with early neurologic deterioration (END) ($p < 0.01$) [67].

Neutrophil elastase from blood was elevated on the first day of a stroke (216.5 ± 26.8 $\mu\text{g/L}$) and significantly increased by 7 days post stroke (384.7 ± 31.4 $\mu\text{g/L}$) compared to control levels (67.4 ± 5.2 $\mu\text{g/L}$) [68]. Plasma neutrophil gelatinase-associated lipocalin (NGAL) is increased in the first few days following acute ischemic stroke [69, 70] and can remain elevated for over a year following the stroke [70]. Neutrophil proteinase 4 (NP4) is also elevated in blood following ischemic stroke [69].

IL8, neutrophil chemotactic factor, is increased in serum within 24 h of ischemic stroke and is paralleled by an increase in blood neutrophils [71]. Though not specific for neutrophils, TNF- α , PAI-1, and tPA showed a significant predictive value of ischemic stroke, whereas IL-6, VICAM-1, ICAM-1, and neutrophil percentage showed only a slight or no association with stroke diagnosis [72]. Serum TNF- α , produced in part by neutrophils, rises early and remains elevated paralleling the prolonged leukocytosis following stroke [73].

8 Neutrophil Response to Acute Stroke: Animals

8.1 Neutrophils Detect Injury

The mechanisms by which neutrophils detect pathogens and organ injury and infiltrate tissue have been well studied [2, 74]. Though there have been an increasing number of studies on how neutrophils respond following stroke, many of the details are derived from peripheral blood vessels and the response to pathogens rather than

ischemic brain [2]. Thus, the following description is based upon a composite derived from various tissues.

Following a stroke, circulating neutrophils detect injury via changes in molecules on the endothelial surface and molecules released by damaged endothelial cells. Once attracted to an area of injury, neutrophil recruitment involves a series of steps: capture/tethering, slow rolling, arrest/adhesion, crawling, and transmigration which is mediated by a series of molecules (Fig. 3) [2, 74]. Attenuation of CXCR4 and activation of CXCR2 lead to neutrophil release from bone marrow in response to inflammation, accounting in part for increased circulating neutrophils [75]. Neutrophil recruitment to damaged endothelium is initiated by changes on the surface of endothelium that result from stimulation by inflammatory mediators including histamine, cysteinyl-leukotrienes (e.g., leukotrienes B4) and cytokines (e.g., IL-1 β) and chemokines that are released by resident sentinel vascular leukocytes, brain endothelial cells, and brain microglia [2]. For example, chemokines activate neutrophil receptors including CCR1, CCR2, and CCR5 which mediate neutrophil recruitment to ischemic tissue [76]. CXCL1, -2, and -5 released from endothelial cells signal via neutrophil CXCR2 to activate neutrophils and promote adhesion to endothelium [2]. Endothelial cells can also be activated by DAMPS (HMGB1, Hsp72 from brain) that increase adhesion molecules.

Following infection or injury, P-selectin expression on endothelial cells increases within minutes after release from Weibel–Palade bodies, and E-selectin is synthesized in endothelial cells within 90 min of injury [2]. P-selectin and E-selectin on endothelial cell surfaces bind P-selectin glycoprotein ligand-1 on neutrophils to capture or tether them (Fig. 3) [74]. DAMPs released from injured endothelial and brain cells can also activate neutrophils (Fig. 2) which may promote adhesion.

Arrest, adhesion, and crawling are promoted by neutrophil integrins LFA-1 (also called α 1 β 2 or β 2 integrin C11a-CD18) and MAC1 (also called α M β 2 or CD11b-CD18) (Fig. 3). Transmigration requires integrin interactions with CAMs (ICAM-1 or -2) for neutrophils to pass between endothelial cells (paracellular) or through endothelial cells (transcellular) and across the basement membrane [2, 74]. Once across the vessel wall, neutrophils infiltrate infected tissue via a second gradient of molecules [2]. However, following stroke it is not as clear whether neutrophils infiltrate tissue very much if at all. In fact most pathological studies show neutrophils on the luminal and abluminal sides of vessels in infarcted brain suggesting they are mostly perivascular [22]. A recent study challenges the concept that neutrophils actually enter ischemic brain tissue [77]. Using an in vivo transient MCAO model in mice, neutrophils were mainly restricted to luminal surfaces or perivascular spaces of cerebral vessels and virtually no PMNs entered the infarcted CNS parenchyma in a period from 1 h to 2 weeks after the stroke. Vascular PMN accumulation did not correlate with vessel permeability, expression of endothelial cell adhesion molecules, platelet aggregation, or release of NETs. The absence of PMN infiltration within infarcted brain per se was supported by studies of 25 human stroke specimens collected at early time points after infarction [77]. This study challenges the conceptual framework of many previous studies, and might suggest that neutrophil interactions with the BBB (endothelium, pericytes, basement membrane, perhaps astrocytes) might explain protective effects of modulating neutrophil functions.

9 Regulating Neutrophils Response in Acute Ischemic Stroke: Animal Models

Given the above biology, a large number of studies, some of which date back several decades, have suggested that modulating interactions between neutrophils and endothelium [78, 79] was associated with decreased neutrophil responses to ischemic brain [80, 81] and improved outcomes following ischemic stroke. Although many of the approaches were not specific for blocking neutrophils, they have provided proof of principle. A summary of selected studies shows there is still much to be learned and more specific approaches are needed in order to develop possible treatments for acute stroke targeted just at neutrophils.

9.1 Cellular Adhesion Molecules

In early experimental studies, ICAM-1 was shown to be upregulated following ischemic stroke. An antibody to ICAM-1 reduced ischemic cell damage after transient but not permanent MCAO in Wistar rats [82]. Knockout of ICAM-1 and immunodepletion of neutrophils decreased infarct volume and mortality and improved outcome following transient MCAO in mice [83]. Another study showed knockout of ICAM-1 decreased infarct volume in transient and permanent MCAO models without having much effect on neutrophil infiltration into ischemic brain, and depletion of neutrophils further decreased infarct volume following transient MCAO [84]. These and other studies led to a study with an antibody (Enlimomab) targeted against ICAM-1 which failed in human ischemic stroke [8] and is discussed below.

At baseline type 2 diabetic rats have higher neutrophil CD11 and endothelial sICAM, and then following stroke have higher levels of both molecules, have greater neutrophil adhesion and cell aggregates and worse outcomes following stroke [85]. The platelet inhibitor cilostazol appears to act on endothelial cells to inhibit expression of adhesion molecules including ICAM-1 and neutrophil adhesion induced by high glucose through increasing NO production [86]. Anti-VCAM-1 antibodies do not protect against focal cerebral ischemia in rats or mice [87]. A problem with targeting CAMs has generally been that this is nonspecific and would prevent adhesion of not only neutrophils but also other leukocytes.

9.2 CD11/CD18

Macrophage-1 antigen (Mac-1) (CD11b/CD18), also called complement receptor 3 (CR3), is a leukocyte beta2 integrin and facilitates leukocyte adhesion, transendothelial migration, phagocytosis, and respiratory burst, all of which may mediate reperfusion-induced injury to ischemic brain tissue. CD11b/CD18 is found not only on neutrophils, but also on monocytes/macrophages and Natural Killer cells and binds to endothelial cell ICAM-1. Pre-ischemic administration of an anti-CD18 monoclonal

antibody increased reflow in microvessels in a primate stroke model [88]. Postischemic treatment at 2–4 h [89, 90] and at 20 min after transient MCAO [91] with anti-CD11b/CD18b monoclonal antibodies decreased infarct volumes and neutrophil migration into ischemic brain. Mice with knockouts of CD18 have smaller infarcts, decreased mortality, and decrease of neutrophils in ischemic brain following transient MCAO [92, 93], though there were no differences following permanent MCAO [93].

9.3 *Neutrophil Inhibitory Factor*

Neutrophil inhibitory factor (NIF) blocks neutrophil adhesion by binding the CD11a and CD11b beta 2 integrins. Administration of recombinant neutrophil inhibitory factor (rNIF) beginning at 2–4 h after temporary MCAO in adult rats decreases infarct volumes at 7 days, but had no effect if treatment were continued for 6 h rather than 48 h, and had no effect in permanent MCAO [94, 95]. In another study, administration of NIF at 4 and 6 h post MCAO decreased infarct volume and brain swelling and improved functional outcomes [96]. A human trial of NIF failed to show improvement in acute ischemic stroke [9]. Though the reasons for the failure are discussed below, it is important to note that NIF would affect not only neutrophils but any beneficial roles of monocytes and Natural Killer cells.

9.4 *CXCR1/2 and CCR2*

CXCR1/2 receptors (IL8 receptors) are involved in recruitment of neutrophils into infected or injured tissue. IL8 is released by endothelial cells and monocytes (and neutrophils themselves) to act on CXCR1/2 receptors to recruit neutrophils. Blockade of CXCR1/2 chemokine receptors with the drug reparixin protected against brain damage in ischemic stroke in mice, decreased MPO staining in the stroke, decreased IL1beta, and improved motor outcomes in two studies [97, 98]. However, two other studies were negative. The CXC chemokine-binding protein Evasin-3 potently inhibits chemokine bioactivity and related neutrophilic inflammation in two mouse models of carotid atherosclerosis [46]. In carotid atherosclerosis, treatment with Evasin-3 was associated with reduction in intraplaque neutrophil and MMP-9 content and weak increase in collagen as compared with Vehicle. However, in acute stroke, Evasin-3 decreased neutrophil infiltration and oxidative stress but without improving stroke outcomes [46]. In a similar study, administration of a CXCR2 antagonist (SB225002) reduced expression of CXC chemokine subfamily of genes and neutrophil-related infiltration following SB225002 administration but did not improve outcome after cerebral ischemia-reperfusion [99]. Of note the CXCR2 ligand, CXCL5, is increased in CSF but not blood of patients within 24 h of ischemic stroke [100]. CXCL1, another neutrophil attractant, is increased in CSF but not serum of patients with ischemic stroke [101].

CCR2 is another neutrophil receptor that mediates recruitment. Though CCR2 knockout mice have decreased infarct volumes, decreased edema, and decreased

infiltration of both neutrophils and monocytes into ischemic brain [102], this has yet to be confirmed in independent studies. In addition, blocking CCR2 would not selectively block neutrophils.

9.5 *Cannabinoid Receptors/CXCL2*

Cannabinoid 2 receptor (CB(2)) agonists decrease ischemic brain injury in an MCAO model [103]. They do this by acting on neutrophil CB(2) receptors and activating p38 in neutrophils via a chemokine CXCL2 dependent mechanism which inhibits neutrophil entry into ischemic brain [103]. In addition cannabinoids may decrease cerebral infarct size in part by decreasing HGMB1 in plasma and decreasing TLR4 activation of neutrophils [104].

9.6 *P- and E-Selectin*

P- and E-selectin are constitutively expressed in endothelial cells and facilitate entry of leukocytes into brain. When P-selectin is stimulated by thrombin, it translocates from cytoplasmic granules to the plasma membrane. Pro-inflammatory mediators like TNFalpha and lipopolysaccharide (LPS) induce P-selectin mRNA. Administration of a humanized antihuman E- and P-selectin monoclonal antibody decreases stroke volume and improved functional outcomes in a primate stroke model without evidence of immunosuppression [105]. This antibody would bind endothelial E- and P-selectins and would tend to decrease infiltration of all leukocytes including neutrophils into ischemic brain [105]. P-selectin knockout mice have decreased damage to the BBB and marked decrease of neutrophil infiltration following MCAO compared to wild type mice [106] and P-selectin KO mice have smaller infarcts [107]. In one study post stroke anti-P-selectin antibody decreased infarct volumes [107], though in another pretreatment but not post treatment with a P-selectin antibody decreased infarct volume following transient MCAO [108]. An E-selectin antibody decreases infarct volume and neutrophil accumulation following transient MCAO in mice [109]. An L-selectin antibody did not protect against focal cerebral ischemia [110], even though it plays a role in adhesion of neutrophils and other leukocytes to endothelial cells [111].

9.7 *CD47*

CD47 is a cell surface glycoprotein that helps mediate neutrophil transmigration across blood vessels. Knockout of CD47 decreased infarct volumes at 24 and 72 h after a 90 min (temporary) MCAO [112]. CD47ko also decreased edema at 72 h, ameliorated decreases of claudin-5, markedly decreased MMP-9, and decreased neutrophil numbers in brain infarcts [112].

9.8 *Slit1/Robo1*

In the normal brain, the presence of Slit1 on PMNs, and Robo1 on cerebral endothelial cells, generated a repulsive force to prevent the infiltration of PMNs into the brain [113]. During stroke recovery, a transient reduction in Robo1 expression on the cerebral endothelial cells allowed the uncontrolled infiltration of Slit1-expressing PMNs into the brain causing inflammatory reactions [113]. Similarly, lack of the immune receptor CCR5 on brain cells was associated with larger infarcts and worse motor deficits compared to wild type mice, and this was associated with greater influx of neutrophils into the infarct [114].

9.9 *DAMPs/Toll-Like Receptors*

Ischemic brain injury results in the release of a number of DAMPs, including HMGB1, Hsp72, S100A9, peroxiredoxin, mitochondrial peptides, and extracellular nucleic acids (DNA, RNA) [115–117] (Figs. 1 and 2). DAMPs act on neutrophils through specific receptors (toll-like receptors (TLR), co-receptors, FPR1) [115], and result in a pro-inflammatory response involving the production of cytokines, proteases, and ROS [115]. Targeting DAMPs and TLRs has shown promise as a treatment in ischemic stroke. When HMGB1 levels in plasma are reduced with cannabinoids, there is a reduction in infarct size and activated neutrophils [104]. Neutralization of peroxiredoxin with antibodies reduces inflammatory response and infarct volume growth [117]. In patients with ischemic stroke, increased neutrophil expression of TLR4 on day 3 and 7 is associated with worse stroke outcome and infarct volume [118]. When TLR4 is knocked out in mice, ischemic brain injury is reduced [119]. Despite the absence of TLR4, MPO+ cells and Iba1+ microglial cells were increased in brain, suggesting an increase of brain MPO+ cells (which could be neutrophils or macrophage/microglia) does not necessarily equate to worsening of brain injury in stroke.

10 **Neutrophil Mediated BBB Disruption in Ischemic Stroke**

The BBB is disrupted following ischemic stroke. Neutrophils contribute to this disruption through the release of proteases (MMPs, elastase, cathepsin G, proteinase 3), ROS and during the process of migrating across cerebral endothelium [2, 77, 120] (Fig. 1). BBB disruption plays an important role in poststroke cerebral edema and hemorrhagic transformation (HT). The importance of neutrophils in BBB disruption and HT is highlighted by the fact that neutrophil proteins (MMP-9) and genes (LTF, NGAL, CEACAM8, CRISP3) are predictive of HT in patients with stroke [121]. Furthermore, inhibiting or depleting neutrophils reduces BBB disruption and the rate of HT [120, 122]. In contrast, promoting neutrophil activation with LPS enhances BBB disruption in rodent stroke [123]. Thus, targeting neutrophil mediated BBB disruption may have utility to reduce cerebral edema and the rate of HT post stroke.

10.1 Neutrophil MMP-9

Neutrophils are an important source of MMP-9 within the first 24 h of ischemic stroke. MMP-9 is increased in plasma within the first 2–6 h of stroke in patients, primates, and rodents [124, 125]. We found increased leukocyte MMP-9 mRNA within 3 h of stroke [126, 127]. Increased plasma MMP-9 levels correlate with BBB disruption and predict tPA related HT [128]. In humans with ischemic stroke, MMP-9-positive neutrophil infiltration is associated with BBB breakdown, basal lamina type IV collagen degradation, and hemorrhagic transformation (HT) [129].

Targeting neutrophil MMP-9 has been evaluated as a treatment in ischemic stroke. Mice lacking leukocyte MMP-9 have decreased BBB disruption and infarct size [130] which appears to be mediated by leukocyte MMP-9 and not brain MMP-9 [130, 131]. In neutrophils, MMP-9 is regulated by carcinoembryonic antigen-related cell adhesion molecule 1 (CEACAM1). Knocking down CEACAM1 promotes neutrophil MMP-9 expression and enhances BBB breakdown in ischemic stroke [132]. Pharmacological inhibition of MMP-9 post stroke also reduces BBB disruption and HT [133]. For example, the rate of HT in rats can be reduced with BB-94, a broad spectrum MMP inhibitor, and minocycline, an inhibitor of MMP-9 and microglia [134]. Ongoing studies are evaluating the role of minocycline to reduce HT in stroke patients. Though early inhibition of MMP-9 reduces HT in animals, delayed MMP-9 inhibition enhances brain injury and worsens stroke in part by impairing vascular remodeling [135].

Granulocyte colony-stimulating factor (G-CSF) has activity to induce neutrophil mobilization. In a rat MCAO stroke model, G-CSF increases peripheral neutrophil numbers at 24 h, and this is associated with an increase in MMP-9 and tPA related hemorrhagic transformation [136]. G-CSF was found to potentiate neutrophil release of MMP-9 in the presence of tPA. In humans, G-CSF administered following ischemic stroke has also been found to increase peripheral leukocyte count. This increase was not associated with any improvement in neurological outcome at 90 days, nor an increase risk of HT. Though the rationale as to why G-CSF did not show clinical benefit is unclear, it may have enhanced early mobilization of inflammatory neutrophils which offset any potential beneficial effect of G-CSF.

10.2 Neutrophil Elastase

Neutrophil elastase degrades basal lamina and extracellular matrix. In experimental stroke pharmacological inhibition of neutrophil elastase reduces BBB permeability, decreases cerebral edema, and improves neurological outcomes [137]. In mice lacking elastase, ischemia-induced BBB disruption is reduced, as is infarct volume, cerebral edema, and leukocyte-endothelial adhesion [138]. Furthermore, pharmacological inhibition of elastase in MMP-9-null mice further decreased infarct volume and BBB disruption, indicating effects independent of MMP-9 [138]. These results are important since they suggest that combined inhibition of MMP-9 and elastase may be more effective than either alone.

10.3 Neutrophil ROS

Neutrophils are an important source of ROS following stroke and reperfusion of ischemic brain [140, 148]. ROS disrupt the neurovascular unit through damage to endothelial cells, pericytes, smooth muscle cells, and astrocytes. This results in increased BBB permeability, cerebral edema, and hemorrhagic transformation.

Superoxide radicals are mediators of reperfusion-induced BBB disruption [139]. Neutrophils generate superoxide by the enzyme nicotinamide adenine dinucleotide phosphate oxidase (NOX). Inhibition of superoxide results in reduced BBB disruption in rodents. When NOX is inhibited with apocynin or by genetic ablation, the severity of BBB disruption is reduced, as are infarct volumes and the degree of neutrophil infiltration [140, 141].

Several studies have evaluated ROS as a target to reduce BBB disruption and HT in ischemic stroke. Though animal studies of the spin trap agent NXY-059 showed promise [142], it failed to improve stroke outcome or reduce hemorrhagic transformation (HT) in stroke patients [143]. Another spin trap agent, N-t-Butyl-Phenylnitron (PBN), also reduces tPA related HT in rodent stroke [144] but worsens HT in rabbit stroke [145]. The free radical scavenger edaravone decreases BBB disruption and HT in rodent stroke [146] but increased HT in stroke patients [147]. Benidipine, a dihydropyridine calcium channel blocker, also inhibits neutrophil ROS production and improves stroke outcomes in rats [148]. Given these variable results, evaluation of candidate compounds in multiple animal models of HT may be beneficial [149].

11 Stroke, Infection, and Neutrophils

Infections, including pneumonia and urinary tract infections, worsen outcomes in patients with ischemic stroke [150–152]. The mechanisms for this worsening appear to be related in part to neutrophils. In one animal study, systemic inflammation was induced in mice by peripheral interleukin-1beta (IL-1beta) challenge and a transient MCAO performed [153]. This resulted in conversion of a transient to a sustained disruption of the tight junction protein, claudin-5, and also markedly exacerbated disruption to the cerebrovascular basal lamina protein, collagen-IV. These alterations were associated with a systemic inflammation-induced increase in neurovascular gelatinolytic activity that was mediated by a fivefold increase in neutrophil-derived MMP-9 in ischemic brain. MMP-9 inhibition attenuated the deleterious impact of systemic inflammation on brain damage, edema, neurological deficit, and incidence of hemorrhagic transformation [153]. Influenza infection has also been found to worsen rodent stroke in part by increasing release of cytokines and neutrophil-derived MMP-9 that are mediated by $\alpha 7$ nicotinic acetylcholine receptors [154].

Another study has implicated neutrophils in worsening outcome from systemic inflammation [123]. Systemic inflammation was produced by intraperitoneal LPS (a gram-negative bacteria coat protein) followed by a transient MCAO. Brain damage and neurological deficit were worsened by LPS. Administration of an IL-1 receptor antagonist abolished the effect of LPS on brain damage. Systemic IL-1 increased ischemic damage to a similar extent as LPS. Neutropenia abolished the damaging effects of systemic IL-1. The data show that LPS induction of neutrophil-induced IL-1, perhaps via neutrophil mobilization via CXC chemokine induction, worsens stroke outcomes following systemic inflammation. The results may help to explain the poorer outcome in stroke patients presenting with infection [123].

12 Protective Role of Neutrophils in Stroke

Though the cerebral ischemia field generally lumps all neutrophils into “bad guys,” it would be important to know if neutrophils serve any neuroprotective functions. If they do have protective functions, it would be useful to avoid inhibiting these functions if anti-neutrophil treatments are to be designed to treat stroke. As described above, patients with stroke are at risk for infections, and infection clearly worsens in stroke in patients and in animal models of stroke [123, 154–156]. Therefore, significantly inhibiting the function of neutrophils in fighting infections might worsen outcomes—as discussed in the human trials below—and any neutrophil therapy for stroke should take this into consideration.

In addition, it has been suggested that neutrophils might adopt a pro-inflammatory N1 phenotype or an anti-inflammatory N2 phenotype in the CNS depending on environmental cues [157]. The N2, anti-inflammatory phenotype, may be adopted when neutrophil numbers exceed a critical threshold.

Clearly protective roles of neutrophils have been described including neutrophils performing wound debridement, and simply removing neutrophils from an inflamed tissue can result in more tissue pathology even in the setting of sterile injury [158]. MMP-9, derived either from brain or neutrophils, activates VEGF to promote revascularization at sites of injury [159]. Neutrophils facilitate recruitment of monocytes to inflamed tissue, and by promoting their own removal from tissue contribute to resolution of inflammation and promote repair [2]. During termination of inflammation, neutrophils release several anti-inflammatory molecules. These include annexin-1, lipoxin A4, resolvins, and protectins that decrease neutrophil migration and recruitment [2]. These molecules or their agonists might also be worth testing in acute ischemic stroke.

A recent dose escalation study of G-CSF in acute stroke did not detect any deleterious effects in spite of expected increases of both neutrophils and monocytes in a 12 h time window post stroke [160]. In animal studies, G-CSF knockout animals have larger infarcts than controls, and administration of G-CSF decreases infarct size whereas it increases peripheral but not brain neutrophil counts [161].

13 Human Stroke Trials Using Immunotherapy

Animal studies in the 1990s demonstrated excellent outcomes in animal stroke models following a variety of therapies that modulated the immune system, including antibodies against ICAM and modulating NIF/CD18 [42, 80, 82–84, 91, 93]. These studies led to two clinical trials both of which were negative and led to loss of enthusiasm for this approach. A third trial (LeukArrest, anti-CD11/CD18) was also performed and stopped during the trial because it was unlikely to show benefit [162], though the final results were never reported.

The results of the Enlimomab Acute Stroke Trial were published in 2001 [8]. Patients with ischemic stroke ($n=625$) were randomized to anti-ICAM antibody (Enlimomab) or placebo within 6 h of stroke onset and treatment continued for 5 days. At 90 days more patients died with Enlimomab and fewer were symptom free compared to placebo ($p=0.004$). There were more infections and fever with Enlimomab compared to placebo, and patients with fever had worse outcomes [8]. After the trial it was realized that the Enlimomab antibody may not have been sufficiently humanized and may have stimulated an immune response on its own [162]. A follow-up animal study to the human Enlimomab trial provided some insight into why this particular trial may have failed. It was found that administration to rats of a murine antibody preparation against ICAM-1, 1A29, elicits the production of host antibodies against the protein, activation of circulating neutrophils, complement activation, and sustained microvascular activation [163]. 1A29 did not improve stroke outcome in the rat or mouse, but pre-administration in the rats worsened stroke outcomes. Thus Enlimomab may have inadvertently stimulated an immune response that worsened stroke outcomes.

An additional reason the Enlimomab trial may not have shown benefit was the nonspecific leukocyte target ICAM-1. Blocking ICAM-1 would be expected to interfere with adhesion of almost all types of leukocytes and potentially decrease response to infection. Six enlimomab-treated stroke patients developed meningitis (none for placebo), suggesting that the anti-ICAM treatment impaired host responses to infection and this may have contributed to worsening of outcomes. Future immune therapy trials should carefully consider and test whether a potential therapy would significantly impair host responses to infections. Since rodents tend to be resistant to many infections, additional animal stroke models with vulnerability to infections similar to that of humans should be tested.

Another human trial of immunotherapy in stroke also failed to show benefit in ischemic stroke. In animals NIF had been shown to prevent neutrophil adhesion to endothelium by inhibition of CD11a and CD11b beta(2) integrins [164] and CD18 on PMNs [165]. In addition, UK-279,276 (NIF) reduced infarct volume in a rat MCAO reperfusion model [165]. Based on these and other animal data ASTIN (Acute Stroke Therapy by Inhibition of Neutrophils) was an adaptive phase 2 dose-response-finding, proof-of-concept study to establish whether UK-279,276 improved recovery in acute ischemic stroke [9]. 966 patients were entered into the trial and treated within 6 h of symptom onset. UK-279,276 did not improve recovery

in acute ischemic stroke patients [9], but it did not produce serious side effects and did not increase infections or fever. Reasons for the failure of the trial could have included: (1) blocking neutrophil adhesion does not improve stroke outcomes in humans but does so in rodents (2) the dose or timing of administration did not produce the desired biological effect (3) the effects of UK-279,276 on neutrophils are different in rodents compared to humans. This trial did suggest that failure was not related to an adverse effect on host immune response to pathogens.

14 Conclusions

The future of modulating neutrophils in ischemic stroke is thus uncertain. It seems clear that specific inhibition of this cell type might be desirable compared to a global suppression of the immune response. Moreover, approaches that maintain some response to pathogens would appear to be preferred. It is interesting that low dose G-CSF increased leukocyte counts post stroke in humans, including neutrophil counts, without worsening clinical status [166]. With a recent study suggesting neutrophils do not infiltrate ischemic brain but rather remain intravascular or perivascular [77], the role of targeting neutrophils to reduce ischemic brain injury seems unclear. Instead, targeting neutrophil effects on the BBB and neurovascular unit may be more fruitful. Though neutrophil MMPs and other proteases (elastase, cathepsin G) disrupt the BBB in stroke, additional studies are required to determine whether their inhibition can improve human ischemic stroke. Targeting neutrophils to prevent future stroke may also be beneficial. As our understanding advances regarding the roles of neutrophils thrombosis and atherosclerotic plaque rupture, modulating neutrophils may serve as a novel method to prevent stroke.

15 Author Contribution Statement

All authors contributed to the presented work and made critical revision of the manuscript for important intellectual content.

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The Function of Cytokines in Ischemic Stroke

Christopher C. Leonardo and Keith R. Pennypacker

1 The Complex Sequelae Resulting from Ischemic Stroke

Ischemic stroke produces a complex injury profile that is characterized by multiple, successive waves of tissue injury. The acute phase primarily involves energy failure and excitotoxic injury, which are hallmarks of most neurodegenerative diseases and set the stage for delayed cell death. During the acute phase, neural tissue most directly irrigated by the occluded vessel experiences rapid energy failure resulting from oxygen and glucose deprivation. The resulting failure of cellular respiration necessitates the switch to anaerobic energy production, leading to acidosis. This metabolic switch, coupled with excessive release of the neurotransmitter glutamate from injured neurons and neighboring astrocytes, promotes ionic instability that compromises cell membranes and facilitates neuronal cell death. In addition to free radical injury, which is achieved in large part through the production of superoxide and peroxynitrite radicals, the release of sequestered intracellular calcium stores also promotes apoptotic signaling and cellular oedema [1].

These mechanisms of ischemic stroke pathology are well documented and have provided the basis for several decades of stroke research, whereby the primary goal was to limit excitotoxicity as means of limiting these injurious responses.

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Unfortunately, this approach was met with very limited success [2, 3], as relatively few stroke patients are eligible for the thrombolytic therapies (i.e., tissue plasminogen activator) that must be administered within 3–6 h of stroke onset. With few viable treatment options, the majority of patients are limited to supportive care during the period of infarct expansion. During this period, the viable tissue surrounding the core infarct becomes compromised through perpetuation of the above mechanisms coupled with detachment-induced injury from loss of synapses and neuroglial support. Without recanalization and restoration of blood flow, neural injury will progress based upon the severity of the initial insult. This will continue until endogenous remodeling mechanisms, such as glial scarring, are successful in protecting healthy surrounding tissue from the inflammatory microenvironment present within the infarction.

The rapid progression of neuronal injury, as well as the inability to diagnose ischemic stroke patients in time to receive thrombolytic therapy, has prompted investigations into other therapeutic targets. Among these was a broad category of neuroinflammatory responses that encompasses a wide range of deleterious phenomena. For example, depending upon the specific context, neuroinflammation may include blood–brain barrier (BBB) permeability, glial scar formation, extravasation of peripheral immune cells, and/or activation of resident immune cells [4]. Additionally, and somewhat paradoxically, immune cell chemotaxis is accompanied by the upregulation and release of cytokines, chemokines, prostaglandins, proteases, and other proinflammatory molecules that perpetuate cellular disruption and tissue injury [5]. To complicate matters, these signals collectively demonstrate dual and often overlapping roles that cloud interpretation of their individual actions within the grand scheme of stroke pathology. For these reasons, the inflammation that ensues following ischemic stroke produces a complicated injury profile with multiple mechanisms that need to be addressed in order for any therapeutic approach to be successful.

A critical component of any therapeutic is the ability to dampen proinflammatory signaling that contributes to delayed cell death. This delayed wave of neural injury is facilitated by BBB disruption in response to the initial ischemic event. Within hours to days of stroke onset, matrix metalloproteinases (MMPs) cleave tight junction proteins involved with sealing the barrier. The result is a biphasic opening of the BBB that is initiated by MMP-2 and enhanced through the actions of MMP-9 [6]. Immune cell signaling exacerbates the inflammatory milieu within the core infarct. Microglia respond to resident death signals, while peripheral leukocytes are activated by CNS antigens that have leaked into the systemic circulation. The result is a feed-forward proinflammatory microenvironment that is modulated by resident glia and peripheral leukocytes through the upregulation and release of cytokines. Thus, cytokine signaling has profound effects on the ultimate fate of the injured brain and is regulated by activated immune cells both within and outside of the brain parenchyma.

2 Cytokines as Therapeutic Targets for Ischemic Stroke

Cytokines are involved in a wide variety of biological processes, including embryonic development, stem cell differentiation, innate and adaptive immunity, cognition, progressive neural injury and neurodegenerative disease pathology. The complexity of cytokine actions is reflected in the distinct families of cytokines, which include interleukins, chemokines, mesenchymal growth factors, adipokines, and the tumor necrosis factor family [7]. Depending upon the cytokine, these signals can influence cell signaling by cell surface receptor binding, antibody binding, and direct DNA binding. Thus, this interesting class of molecules has attracted much attention as researchers have searched for new therapeutic targets.

Ischemic stroke produces a biphasic injury profile that can be characterized in terms of an “acute” excitotoxic response followed by a “delayed” neuroinflammatory response. Indeed, cytokines have emerged as important signaling molecules that play key roles in both injury phases and are intimately connected with immune cell activation. To date, there have been a multitude of studies aimed at uncovering the precise mechanisms by which cytokines influence cellular responses to cerebral ischemia. Results from clinical studies and animal models have identified several key cytokines that can be targeted for the treatment of ischemic stroke.

The body of work regarding cellular immune responses to ischemic stroke is extensive and is discussed in great detail throughout this book. Although it was not historically the case, neuroinflammation is now widely regarded as a principle cause of delayed infarct expansion that is driven, in large part, by cytokine signaling [8, 9]. Both the acute and progressive phases of stroke injury have been linked to the upregulation and release of proinflammatory cytokines [4, 5]. Animal models have offered the means to conduct controlled studies designed to elucidate the mechanisms by which cytokines influence injury and remodeling of the ischemic brain. This section reviews some of the key findings related to cytokine production and signaling after ischemic stroke. Additionally, promising therapeutic approaches predicated on modulating cytokine signaling are discussed within the context of neuroinflammation and delayed infarct expansion.

2.1 Tumor Necrosis Factor Alpha Signaling and Expression

Tumor necrosis factor alpha (TNF α) has long been identified as a key proinflammatory effector of ischemic tissue. Although it is principally produced by activated macrophages, virtually all lymphocytes have the capability of producing TNF α as part of the systemic inflammatory response. This cytokine binds two distinct receptors, TNF receptor type 1 (TNFR1) and TNF receptor type 2 (TNFR2), and TNFR signal transduction can have various consequences including induction of NF- κ B and AP1 transcription factors, as well as initiation of apoptotic

signaling through caspase activation. To complicate matters, TNFR activation can also suppress inflammatory signaling and apoptosis.

The existence of distinct TNFRs and various intracellular signaling pathways coupled to TNFRs explains some of the inconsistencies from ischemic stroke studies. For example, activation of TNFRs can result in two antagonistic responses. Virtually all cells typically present within the brain after stroke express TNFR1. The intracellular death domain contained within TNF α R1 is linked to Fas-associated protein with a death domain (FADD) signaling, where activation can result in the induction of caspases and/or cytochrome c release. Alternatively, TNF α binding to TNFR2 can activate TNF receptor-associated factor 2 (TRAF2), which induces expression of cell survival-associated genes that are regulated by mitogen-activated protein kinase (MAPK), NF- κ B and AP-1 transcription factors [10, 11]. Additionally, TNFR2 lacks an intracellular death domain and is expressed on the surfaces of cells present within the cerebral infarct, including resident glia (oligodendrocytes and astrocytes), cerebrovascular endothelial cells, T-cells, and monocytes. Thus, the dualistic nature of TNF α signaling can produce caveats concerning the measurement of cytokine levels in the brain. In fact, a recent study demonstrated that TNF α production increased in both microglia and monocytes between 24 and 72 h after transient middle cerebral artery occlusion (tMCAO). Additionally, this study showed a positive correlation between TNF α production and the phagocytic capacity of microglia [12], which might suggest a beneficial role in response to ischemia.

Since TNFR binding can result in a multitude of effects depending upon the receptor activated and the local environment (i.e., the cell types present and the extent of inflammatory signals), the timing of cytokine production and release has been a subject of great interest as researchers work to understand the functional consequences of TNF α signaling. Animal models of ischemia, particularly those investigating immunoinflammatory mechanisms, have been instrumental in characterizing protein and mRNA levels of various cytokines, including TNF α . Studies exploring ischemia/reperfusion injury in liver found that Kupffer cells and neutrophils increased expression of TNF α along with other inflammatory mediators including reactive oxygen species (ROS) and nitric oxide (NO) [13]. Importantly, elevated TNF α also correlates with damage to various other body tissues following ischemia [14].

Consistent with ischemic peripheral tissues, TNF α transcript and protein levels are consistently upregulated in the injured and ischemic brain. This cytokine is expressed at early time points following stroke and is a critically important facilitator of immune cell recruitment to the cerebral infarct. TNF α gene transcript was elevated as early as 6 h following tMCAO in mice and remained elevated at extended time points up to 96 h after stroke [8, 15, 16]. Furthermore, TNF- α was elevated in the systemic circulation 24 h after tMCAO in the same model system, but returned to normal levels by 96 h post-stroke [16].

A similar expression pattern was detected in the spleen and provides clues as to its role in progressive inflammation. The spleen is a reservoir of immune cells and plays a substantial role in systemic immune responses. There is now a body of work

implicating splenic signaling in progressive neural injury following various insults including ischemic stroke, hemorrhagic stroke, and traumatic brain injury (TBI). Because the BBB is breached as a result of these and many other insults, activated peripheral immune cells can exert effects both systemically and at the level of the brain. Mice subjected to tMCAO showed a splenic expression pattern where the expression of various cytokines, including TNF α , was increased between 1 and 2 days following stroke [15, 17, 18].

2.2 *TNF α as a Therapeutic Target for Ischemic Stroke*

Various experimental approaches have been used to determine whether TNF α is a viable target for the treatment of stroke. Some of these include exogenous administration and/or blockade of TNF α , the utilization of knockout mice lacking the TNF α gene (TNF α ^{-/-}) or key immune cell populations, and animals subjected to splenectomy prior to insult. These methodologies have provided valuable information as to the role of TNF α in proinflammatory signaling. Furthermore, data from these models provides insights into acute signals originating at the level of the brain as well as those that are regulated by the peripheral immune response.

Popular approaches to investigating cytokine signaling involve boosting levels of the protein, and conversely, deletion or blockade of the protein. Rodent stroke models, including tMCAO and permanent MCAO (pMCAO), have been useful in conducting such studies due to their cost-effectiveness and reproducibility. Studies show that the administration of TNF α increases infarct volume in these models [19], demonstrating a deleterious role for TNF α that is dose-dependent. Similarly, reducing TNF α by blocking the biosynthetic enzyme, TNF α converting enzyme (TACE), was effective in reducing infarct volume and functional deficits in rats subjected to pMCAO [20]. Consistent with these data, other data showed efficacy in reducing tMCAO injury following administration of TNF α neutralizing antibodies [21].

Indeed, this cytokine has traditionally been regarded as a proinflammatory mediator that exacerbates cellular injury due to its elevation in neurodegenerative disease and after brain injury. However, several lines of evidence indicate that TNF α may be at least moderately misunderstood since it also appears to play a beneficial role under certain conditions. Data showed that TNF α upregulates the antioxidant superoxide dismutase 2 (SOD2), which is important in converting superoxide into molecular oxygen or the less damaging radical hydrogen peroxide. Interestingly, SOD2 has been implicated in stroke preconditioning [11], which would suggest a protective role for TNF α . Consistent with a protective role for TNF α , mice lacking the TNF α gene actually demonstrated worse outcomes in terms of total infarct volume relative to control mice [22]. To further complicate the issue, deletion of the gene coding for TNFR1, but not TNFR2, protected mice against pMCAO [22]. These data are consistent with the pro-apoptotic and cell survival-associated roles of TNFR1 and TNFR2, respectively, and thus highlight the dual nature of TNF α .

From a strictly therapeutic perspective, there is evidence that selective targeting of TNF α itself or TNFRs may prove beneficial for the treatment of ischemic stroke. The promise of this approach comes from TNFR effects and altered TNF expression patterns in response to known neuroprotective treatments and strategies. For example, activation of TNFR2 triggered oligodendrocyte (OL) progenitor cell recruitment in a cuprizone-induced white matter injury model [23]. Since white matter injury is directly related to motor deficits in ischemic stroke pathology, this mechanism may prove beneficial to improve these outcomes. Other data showed that administration of cocaine- and amphetamine-regulated transcript (CART) decreased infarct volume at early and delayed time points following tMCAO. Furthermore, the observed neuroprotection was accompanied by reduced TNF α levels in the blood and a dampened proinflammatory T-cell response [16].

Interestingly, studies investigating immune cell function also point to a crucial role for TNFR signaling. Peripheral immune activation is now an emerging area of research due to a growing body of work demonstrating a clear link between exacerbation of brain injury and the activation of T-cells, monocytes, and peripheral macrophages. Removal of the spleen, a lymphoid organ that acts as a reservoir for peripheral immune cells, has been shown to elicit protection in models of stroke, TBI, and ischemia-reperfusion injury to peripheral body tissues [24–27]. Since splenectomized animals are protected against ischemia, both in peripheral tissues and at the level of the brain, researchers have also focused efforts on cytokine signaling in spleen-derived immune cells. One technique involves harvesting splenocytes from animals subjected to focal ischemia. These cells are then activated and assayed to determine the inflammatory profile, including cytokine production and release.

From splenectomy experiments and assays with cultured splenocytes from ischemic tissues, spleen-derived TNF α has emerged as a candidate target due to its response profile. For example, splenectomy caused reductions in hepatic injury and leukocyte infiltration that were accompanied by decreased TNF α release [25]. In addition, neuroprotection and improved spatial memory performance following splenectomy were associated with decreased TNF α in a model of TBI [27]. Splenocytes harvested from CART-protected animals showed a reduced capacity to produce proinflammatory cytokines, including TNF α [16]. In a model of tMCAO, splenocytes harvested from stroke mice showed upregulated TNF α expression after activation with CD3/CD28 antibodies [8]. These experiments were later replicated to show similar effects in splenocytes harvested up to 24 h following tMCAO, demonstrating a prolonged effect of TNF α production following ischemic stroke in which the cytokine was elevated as late as 96 h after the insult [16]. Alterations in splenic TNF α have also been observed in response to stem cell therapies. Treatment with hematopoietic stem cells (HSCs) blocked splenic TNF α gene transcript [17], while human umbilical cord blood (HUCB) cell treatment reduced the extent of concavalin A-induced TNF- α production and splenocyte proliferation [28].

To date, the various approaches to blocking TNF α signaling have been met with some success in animal models and as well as the clinical setting. Administration of MMP/TACE inhibitors has been effective in rodent models, with BB-94

demonstrating protection against transient global ischemia and KB-R7785 reducing neural injury following MCAO. In addition to inhibiting TACE, which is a key regulator of TNF α activation, another promising approach is blocking the soluble form of TNF α (sTNF α) and its receptors (sTNFRs). This method would act to blunt the effects of sTNF α binding, thus offering another level of regulation. Recently, the TNF α blocking compound etanercept has shown preclinical efficacy along with extended efficacy in reducing motoric and gait disturbances in a clinical trial [29, 30]. Although there is some promise in the clinic and various reports of preclinical efficacy, the utility of targeting TNF α remains debatable when considering its complex, dualistic role.

2.3 Interleukin-1 Beta (IL-1 β) Signaling and Expression

Interleukin-1 beta (IL-1 β) is another putative proinflammatory cytokine associated with exacerbated injury following ischemic stroke. A member of the IL-1 cytokine family, IL-1 β is predominantly produced by macrophages as a proprotein that is proteolytically cleaved into its active form by caspase 1. In general, IL-1 β functions in multiple capacities to increase cell proliferation and differentiation, facilitate immune cell chemotaxis through the activation of chemokines, and regulate apoptotic signaling pathways.

In human stroke patients, IL-1 β is increased in plasma and cerebrospinal fluid during the acute phase [30, 31]. Together with TNF α and MMPs, IL-1 β plays a key role in the BBB disruption that renders the brain susceptible to activated peripheral macrophages and monocytes. Perivascular astrocytes are a major source of neural IL-1 β at this early time point [32]. In fact, a recent study demonstrated elevations in IL-1 β as early as 1 h after tMCAO that appeared to be influenced by the activity of MMP-2 [33]. However, microglia begin to upregulate IL-1 β as they transition from the early M2 phenotype, which is predominantly thought to be beneficial, to the putative M1 proinflammatory phenotype [34].

IL-1 β has most frequently been examined by measuring mRNA content at selected time points following experimental stroke. Previous studies using the tMCAO model showed that IL-1 β was elevated in ischemic brain at 6 h post-stroke and remained increased up to 96 h after insult [16].

2.4 IL-1 β as a Therapeutic Target for Ischemic Stroke

Consistent with a proinflammatory role in disease pathology, animal studies showed that therapeutic doses of HSCs inhibited splenic chemokine and cytokine production, including IL-1 β , when administered 24 h after tMCAO. Additionally, treatment reduced apoptotic neurons, activated macrophages/microglia, and T-cell infiltration [17]. Similarly, the deleterious effects of IL-1 β signaling were demonstrated in a

model of neurodegenerative disease (ME7 prion disease) in which intrahippocampal injection of IL-1 β induced astrocytic chemokine production and enhanced infiltration of T-cells, neutrophils, and monocytes into the brain parenchyma [35]. The recruitment of peripheral immune cells observed in this study is consistent with the IL-1 β profile previously detected in the spleen following focal ischemia. For example, IL-1 β gene transcript was elevated by 22 h post-reperfusion in a rodent model of tMCAO [15, 18] and remained elevated at 48 h after stroke [17]. Interestingly, the involvement of spleen-derived IL-1 β in brain injury was also documented following TBI in rats. Removal of the spleen immediately following TBI reduced IL-1 β production, and this effect occurred concomitantly with reductions in mortality and vasogenic oedema. Additionally, the same animals showed improved performance on the Morris Water Maze, demonstrating a spatial memory benefit that was directly correlated with reduced levels of IL-1 β [27].

The preponderance of data demonstrating a role for IL-1 β in perpetuating a pro-inflammatory microenvironment makes this cytokine an attractive target for stroke therapy. As such, many laboratories are investigating mechanisms of inhibiting IL-1 β signaling. A recent study showed that mice deficient in lactadherin, an endogenous glycoprotein with IL-1 β inhibitory actions, showed increased IL-1 β production and greater infarct size following permanent distal MCAO relative to wildtype controls. Furthermore, the increase in cerebral infarction was negated in mice that were administered an IL-1R antagonist [36].

Despite these encouraging data, there are caveats associated with inhibiting IL-1 β . This cytokine is a key modulator of immune cell function, and its complex regulation at the transcriptional level may pose problems in terms of selective targeting of the proinflammatory isoform. To the latter point, a recent study showed that a specific variant of the IL-1 β gene produces a protein that was associated with higher recanalization rates in stroke patients following tPA administration [37]. In light of these data, it will be critical to understand the biological activity of all SNP variants to ensure that the appropriate IL-1 β variant is being targeted.

2.5 Interleukin-6 Signaling and Expression

Interleukin-6 (IL-6) is another interesting cytokine in that experimental evidence suggests a dual nature, including proinflammatory and prosurvival effects. The cytokine has been implicated in the pathogenesis of many neurological diseases. Similar to the other cytokines discussed within this chapter, the purported involvement in ischemic injury (and therefore potential as a therapeutic target) has arisen largely from observed expression profiles in affected patients and experimental subjects.

One noteworthy distinction between IL-6 and many of the other cytokines investigated is that IL-6 serum levels have shown a positive correlation with negative outcomes and brain injury in human stroke patients [38]. In fact, of the clinical variables used to predict mortality, IL-6 remains among the most reliable [39].

Recent data from the Northern Manhattan Study (NOMAS) has expanded upon the predictive validity of IL-6 as a biomarker. The NOMAS study is an ongoing investigation of risk factors conducted using a multiethnic cohort living in the same community. In this study, researchers concluded that IL-6 was an accurate predictor of ischemic stroke when levels were higher than high-sensitivity C-reactive protein, another biomarker that has been useful as a stroke predictor [40].

Consistent with these clinical data, IL-6 gene transcript is also induced in mice after tMCAO [8], and the signaling pathways activated by IL-6 suggest that this cytokine may prove useful as a therapeutic target. IL-6 signaling is complex and signal transduction can result in very different effects on cells. The receptor complex that is activated by IL-6 is composed of IL-6R subunits alpha (IL-6R α) and beta (gp130). Interestingly, the gp130 intracellular domain can initiate MAPK signaling or activate the Janus kinase/transducers and activators of transcription (Jak/STAT) pathway. Ultimately, these signal transduction pathways result in transcriptional regulation through binding to the promoters of genes driven by AP-1, NF- κ B, and other regulatory elements.

The complexity of IL-6 signaling is mirrored by its diverse functions. This cytokine has been shown to play a role in normal physiological functions, including lipid and iron metabolism, osteoclast stimulation during bone remodeling, angiogenesis, and immune system regulation. In terms of the latter, IL-6 is involved in promoting neutrophil production and chemotaxis, enhancing proinflammatory Th17 cell production and inhibiting the putative proinflammatory cytokines TNF α and IL-1 β [41]. As such, IL-6 has been implicated in various disease states including autoimmune disorders, diabetes, cancer, arthritis, Alzheimer's disease, and even some forms of psychopathology.

2.6 IL-6 as a Therapeutic Target for Ischemic Stroke

Data from experimental models has warranted a close look at IL-6 as researchers search for novel therapeutic targets. However, contradictory results have complicated the discussion as to whether the actions and effects of this cytokine can be modulated effectively. For example, one study using IL-6 knockout mice found that IL-6 gene deletion had no effect on cerebral infarction in the tMCAO model [42]. A separate study using IL-6 knockout mice in the same tMCAO model showed worsened long-term infarcts and functional outcomes relative to wildtype mice. This recent study attributed these effects to reduced capacity for angiogenesis and diminished STAT3 activation. Additionally, cultured neurons, endothelial cells, and mixed glia increased IL-6 gene transcript in response to IL-6 application, suggesting that resident cells of the brain are capable of upregulating IL-6 following exogenous administration [43].

Although these conflicting data highlight the difficulty in formulating conclusions from animal studies, other lines of evidence are consistent with the latter study in demonstrating benefits of IL-6 signaling. Intracerebroventricular (i.c.v.)

administration of recombinant IL-6 prior to pMCAO significantly decreased infarct volume in rats, suggesting that IL-6 is directly neuroprotective at the level of the brain [44]. These data are supported by another group that utilized a mouse model of tMCAO to demonstrate reduced infarct volume following i.c.v. administration of IL-6. Here, the afforded protection was linked to STAT3-mediated SOD2 production and this study included another group of IL-6R deficient mice that showed exacerbated neural injury relative to wildtype and IL-6-treated mice [45].

Successful cell therapies have also provided insights into mechanisms of IL-6 protection. For example, cultured NSCs that were preconditioned with IL-6 prior to transplantation effectively increased angiogenesis, reduced infarct volume, and improved functional outcomes in mice subjected to ischemic stroke. Once again, the protective effects in this study were associated with SOD2 induction [46]. In a separate study investigating the splenic contribution to ischemic stroke, NSCs blocked stroke-induced elevations in IL-6 at 24 h and were found to be in direct contact with CD11b⁺ splenic macrophages at 3 days [18]. It is noteworthy that IL-6 is secreted by macrophages in response to pathogen-associated molecular patterns (PAMPs) and other inflammatory signals. Since IL-6 was upregulated prior to maximal infarct expansion and the blunted IL-6 production was linked to macrophages, it is conceivable that selective inhibition of macrophage-derived IL-6 could be a promising strategy. Interestingly, splenocytes harvested from mice 22 h after tMCAO, but not at other time points, increased IL-6 upon activation *in vitro* [8]. These data suggest that splenic immune cells might be preferentially responsive at a time point consistent with delayed BBB opening and infarct expansion. In this case, it may be possible to develop a more selective and/or time-sensitive approach to targeting IL-6 signaling.

2.7 Interleukin-10 Signaling and Expression

Interleukin-10 (IL-10) is widely considered anti-inflammatory and exerts its actions through multiple mechanisms including the suppression of proinflammatory cytokines and the direct activation of cell survival pathways [47]. This cytokine binds to IL-10R, which is composed of two IL-10R1 domains and two IL-10R2 domains. The IL-10 receptor is coupled to the Jak/STAT signal transduction pathway, which is activated when IL-10 binds to IL-10R1 and initiates the signal transduction pathway through the IL-10R2 subunits [48].

Interleukin-10 blocks proinflammatory signaling by macrophages/microglia, inhibits signaling pathways activated by death receptors, and is associated with the Th2 phenotype [48]. Importantly, Th2 cells have been identified and implicated as contributors to the protective effects of anti-inflammatory compounds targeting peripheral immune cell activation following ischemia. In agreement with its role as a protective cytokine, data from human patients showed that those with worsening scores (based upon the Canadian Stroke Scale) within 48 h of ischemic stroke also showed significantly lower levels of systemic IL-10 [49]. However, the clinical data

available is inconsistent. For example, a study examining cytokine levels before and after thrombolytic therapy found a reduction in IL-10 plasma levels in the acute phase of stroke but no significant change after tPA administration [31]. These data were similar to a previous study that documented reduced IL-10 serum levels after stroke, yet no significant relationship between IL-10 levels and clinical outcomes [50].

Although circumstantial, the clinical reduction in IL-10 suggests that this cytokine may be involved in stroke pathology. Furthermore, experimental data does mirror the clinical expression profile in some ways. Unfortunately, data from these models is inconsistent and therefore difficult to interpret. In the pMCAO model of ischemic stroke, IL-10 gene transcript was reduced in rats 12 h following occlusion [51]. Conversely, a separate study demonstrated elevations in IL-10 mRNA 6 h post-MCAO in rodents.

2.8 IL-10 as a Therapeutic Target for Ischemic Stroke

The protective role of IL-10 has been documented in several experimental ischemia models. For example, IL-10 knockout mice showed larger infarct volumes after permanent focal ischemia. In the same study, primary neuronal cultures from knockout animals showed heightened susceptibility to oxygen glucose deprivation (OGD), while application of recombinant IL-10 protected these neurons from OGD [52]. A later study replicated these deleterious outcomes in mice lacking the IL-10 gene [21] while a recent study showed that the beneficial effects of ischemic preconditioning were negated in mice treated with IL-10R blocking antibodies prior to myocardial infarction [53].

In contrast to the previously mentioned knockout study, a separate study showed only a modest effect on infarct volume in IL-10 knockout mice, with slight increases in infarct volume and neurological deficits. Interestingly, several proinflammatory cytokines were elevated 4 days after stroke but other putative anti-inflammatory proteins were increased between 4 and 7 days after ischemia [54]. Taken together, these data once again demonstrate the caveats associated with targeting cytokine signaling. In this case, it would appear that both compensatory responses and temporal effects are critical in cytokine regulation of the ischemic microenvironment.

Other recent studies have shown that regulatory immune cells, particularly T-regulatory (T-reg) and B-reg cells, may suppress proinflammatory pathways that exacerbate neural injury following stroke [15, 21, 55, 56]. B-cell-deficient mice showed increased neural injury and behavioral deficits relative to controls [57]. A follow-up study employed adoptive transfer of IL-10-enriched, splenic B-cells stimulated with LPS *in vitro*. These cells were administered intravenously to mice 1 day prior to tMCAO. Data showed reduced infarct volumes that were accompanied by reduced extravasation of T-cells and monocytes into the brain parenchyma [58].

These data are consistent with previous reports documenting the success of therapies boosting IL-10 signaling. For example, IL-10 overexpression in neuroglia and

brain endothelial cells reduced infarct volume in mice after pMCAO [59], and viral-mediated i.c.v. gene transfer of IL-10 reduced infarct volume after pMCAO and dampened hippocampal injury after global ischemia in rats [60]. Importantly, the selective targeting of IL-10 signaling may lie in the targeting of specific IL-10-expressing immune cell types, rather than the pan activation of IL-10 signaling pathways. Gene transcript for several cytokines discussed in this chapter, including IL-10, was downregulated within the ischemic brains of SCID mice that are deficient in functional lymphocytes [15]. A recent study using IL-10 knockout mice and Rag2-deficient mice, which lack mature lymphocytes, suggested that IL-10 is required for CD4+ T-cells to facilitate protection in a rat model of facial nerve axotomy [61]. The authors linked protective IL-10 signaling to Th2 cell activation based upon these data and a previous report demonstrating the neuroprotective effects of Th2-type CD4+ cells [62].

Although plausible, questions remain regarding the selective targeting of IL-10 and the immune cell phenotype(s) that should be targeted following cerebral ischemia. Since CD4+ T-cells include distinct T-reg phenotypes and also represent both Th1- and Th2-type lymphocytes, it is difficult to ascertain a precise cellular target from these studies alone. Instead, it may be more beneficial to target a specific time point following ischemia. The preponderance of evidence concerning IL-10, and in fact all cytokines that influence the fate of ischemic tissues, indicates that the timing after insult may hold the key to successful therapeutics.

2.9 Interferon-Gamma Signaling and Expression

Interferon-gamma (IFN- γ) is an important cytokine involved in innate immune responses and has also been identified as a potential therapeutic target for ischemic stroke treatment. Due to its potent proinflammatory effects, IFN- γ may be involved in several aspects of ischemic injury. The IFN- γ receptor exhibits ubiquitous expression and is found on virtually all cell types. Similar to IL-10, the receptor complex consists of two IFN γ R1 domains that confer IFN- γ binding and two IFN γ R2 domains that mediate signal transduction through Jak/STAT phosphorylation. Lymphocytes are major producers of IFN- γ , which primarily targets macrophages to protect against pathogens through chemokine production and secretion [63].

In terms of regulation, IFN- γ production is increased through positive feedback by virtue of its own expression, as well as through the actions of IL-12. The net result of IFN- γ signaling can include a multipronged inflammatory milieu created through the induction of MCP chemokines, B-cell activation, and ROS production. These events, in turn, facilitate a switch in microglia/macrophages toward the M1 phenotype and priming of naïve T-cells toward the Th1 phenotype [60]. Since IFN- γ elicits such profound effects after insult, the expression levels of this cytokine have been investigated both clinically and in preclinical stroke paradigms.

Data from multiple studies has shown increased IFN- γ production at various time points following ischemic stroke. For example, IFN- γ mRNA was increased

2 days following tMCAO and remained elevated 6 days after stroke [51]. Another study found elevated plasma levels of IFN- γ 24 h after tMCAO that returned to control levels by the 4-day time point [16]. Consistent with a role in cerebral infarct expansion, IFN- γ increased 72 h after stroke in the brains of rats subjected to pMCAO [64] and mice subjected to tMCAO [65]. Thus, a role for IFN- γ in post-stroke signaling is evident according to data from studies employing multiple models and species.

A hallmark of stroke, both clinically and in animal models, includes infiltration of peripheral leukocytes into ischemic brain tissue. The extravasation of peripheral immune cells results from several processes, including proteolytic cleavage of BBB proteins by MMPs, induction of endothelial adhesion molecules, recruitment of polymorphonuclear cells, and extravasation of other immune cell types through a compromised BBB [33]. With regard to immune cell populations, the spleen is a known mediator of systemic inflammation and has been shown to contribute to ischemic injury in multiple tissues, including brain. Indeed, splenic IFN- γ gene transcript was increased 22 h after focal ischemia in rodents [15].

One major source of IFN- γ is natural killer (NK) cells, which were aptly named due to their ability to destroy cells infected by pathogens. These immune cells were detected in the brains of ischemic stroke patients and peaked within 2–5 days after onset. In the same study, a mouse model of pMCAO was utilized to further examine NK cell infiltration and cytokine expression. These data showed that IFN- γ -expressing NK cells peaked in the ischemic mouse brain 12 h post-stroke, while the interferon-inducible chemokine IP-10 was detected at both acute and delayed time points after stroke [66]. IP-10 expression is induced by IFN- γ and facilitates the Th1 inflammatory response through dual actions, including the direct activation of the CXCR3 receptor on Th1 cells and antagonism of the CCR3 receptor present on Th2 cells. Thus, IP-10 induction through IFN- γ signaling can amplify the Th1 response and dampen the Th2 response simultaneously, resulting in perpetuated tissue inflammation [67]. Importantly, T-cells and NK cells upregulate IFN- γ in response to signals from activated macrophages. In ischemic tissues, these macrophages are activated through toll-like receptors (TLRs) seated on the plasma membrane. In addition to PAMPs, which are the typical TLR ligands in cases of infection, these receptors also bind proteolytic fragments and epitopes of intracellular proteins that are released following cellular necrosis. These signals trigger macrophage secretion of cytokines that ultimately drive the production of IFN- γ in NK cells and T-cells. Thus, there are multiple ways in which IFN- γ production is induced and perpetuated following brain ischemia.

2.10 IFN- γ as a Therapeutic Target for Ischemic Stroke

By virtue of the mechanisms through which IFN- γ is activated, the microenvironment resulting from cerebral ischemia is ripe for IFN- γ production and signaling. Likewise, the fact that IFN- γ is a major orchestrator of the proinflammatory Th1

response suggests that tissue injury resulting from peripheral immune cell actions may be due, in large part, to IFN- γ signaling. Interestingly, there is some clinical data that indirectly supports this mechanism. Patients that developed post-stroke infections exhibited worse outcomes than their non-infected counterparts regardless of stroke severity. These outcomes were also linked to Th1 responsiveness to the brain antigens myelin basic protein (MBP) and glial fibrillary acidic protein (GFAP), which come into contact with T-cells as a consequence of BBB disruption, and patients with elevated Th1 responses 3 months post-stroke showed worsened outcomes [68]. Although these data are far from a smoking gun, they do support the notion that IFN- γ exerts actions involving immune cell modulation of ischemic tissue injury.

In terms of therapeutic potential, multiple lines of evidence support the selective targeting of IFN- γ as means of mitigating ischemic stroke injury. In one study, administration of simvastatin reduced infarct volume and IFN- γ levels in mice subjected to tMCAO. Interestingly, protection was also afforded by splenectomy, including reduced levels of IFN- γ in brain, and these effects were negated in splenectomized mice that received adoptive transfer of splenocytes [65]. These data linked neuroprotection with peripheral splenic immune cells and IFN- γ signaling, as did reports from other laboratories. For example, i.c.v. administration of IFN- γ blocking antibodies reduced infarct volume following tMCAO in mice. In this study, removal of protective T-reg cell populations augmented the activation of resident microglia and infiltration of T-cells, which produced IFN- γ [21].

Consistent with a peripheral source of IFN- γ , levels of this cytokine were elevated in the spleen 24 h following pMCAO while removal of the spleen reduced neural IFN- γ expression by 72 h [64]. Additionally, splenocytes harvested from mice subjected to focal ischemia demonstrated an increased capacity for IFN- γ production. In similar experiments involving harvested splenocytes but assessing the efficacy of HUCB cell therapy, data showed that protective doses of HUCB cells also reduced splenocyte proliferation and production of IFN- γ in vitro upon stimulation with concavalin A [28]. The ability of protective therapies to blunt IFN- γ signaling was also documented following administration of CART in mice subjected to tMCAO. Data showed that CART administration at the time of reperfusion reduced infarct volume and IFN- γ plasma levels 24 h after reperfusion. In addition, harvested splenocytes failed to produce IFN- γ following artificial stimulation in vitro [16].

Neuroprotection studies and in vitro assays have expanded our understanding of cytokine signaling and the potential for selective targeting in brain disease, including ischemic stroke. Knockout studies are another valuable tool for identifying mechanisms of action and have been utilized to explore the role of IFN- γ in cerebral ischemia. Mice lacking IFN- γ have decreased infarcts compared to WT mice, consistent with a role in exacerbating neural injury. Interestingly, these knockout mice exhibit infarcts that are similar in magnitude to Rag1 knockout mice lacking T-cells and B-cells [69], suggesting a prominent role for peripheral immune cell-derived IFN- γ . In separate studies utilizing a variety of knockouts for IFN- γ and various immune cell populations, the protection afforded by deletion of RAG1 was reversed

by adoptive transfer of splenocytes from wildtype mice, but only partially restored upon transfer from IFN- γ ^{-/-} mice. Although none of the mice showed improvements in neurological scores [69], these experiments linked splenic IFN- γ to brain infarction following ischemic stroke. Another study utilizing SCID mice, a transgenic line similar to Rag1^{-/-} in T- and B-cell deficiency, found reduced cortical infarction in SCID mice 22 h after tMCAO that was accompanied by elevations in splenic IFN- γ mRNA [15].

Taken together, these data provide strong support for selective targeting of IFN- γ and/or immune cell targets that are primarily responsible for IFN- γ -mediated infarct expansion. Although experimental models have improved our understanding of the key players regulating the proinflammatory microenvironment, the complexity of this environment mandates a cautious, deliberate approach. Future studies will determine whether IFN- γ can be modulated effectively to expand the therapeutic window for ischemic stroke treatment.

2.11 Leukemia Inhibitory Factor Signaling and Expression

Leukemia inhibitory factor (LIF), although far less studied compared to the other cytokines discussed in this chapter, may hold important therapeutic potential for the treatment of ischemic stroke and other neurological disorders. Under normal conditions, LIF is instrumental in facilitating cellular functions related to proliferation, differentiation, and cell survival. Under pathological conditions, however, LIF appears to be beneficial through its ability to induce antioxidant- and prosurvival-related gene expression [70]. This anti-inflammatory cytokine is a member of the IL-6 cytokine family [71] and binds to a heteromeric receptor dimer consisting of a LIF-binding chain (LIFR) and a converter subunit (gp130). Receptor activation by LIF can transduce signaling of multiple pathways, particularly those regulated by MAPK [72], phosphatidylinositol-3-kinase (PI3K)/Akt [73], and JAK/STAT [74, 75].

To date, few studies have conducted detailed investigations of LIF expression following brain ischemia. Nevertheless, LIF expression has been documented both clinically and experimentally to some degree. Postmortem tissues from human stroke patients revealed increased LIF expression in peri-infarct zones that localized to neurons and endothelial cells, while LIF plasma levels were found to be reduced within the first 6 h following ischemia. Alterations in LIF expression were also documented in rats after MCAO, with greatest levels detected 90 min post-stroke [76]. In response to experimental cortical lesion, rats showed a 30-fold increase in astrocytic LIF mRNA relative to controls. Specifically, LIF gene transcript was elevated as early as 6 h post-insult and reached its peak at the 24h time point. Closer examination revealed an expression pattern localized predominantly to astrocytes, along with a limited population of microglial cells [77].

The known protective role of astroglia following neural injury, coupled with the induction of LIF gene transcript, suggests that this cytokine may be instrumental in

the endogenous response to injury. The ability to upregulate antioxidant gene expression would be highly beneficial following brain ischemia, as free oxygen and nitrogen radicals are major contributors to the acute phase of injury. Similarly, the ability to bolster expression of prosurvival genes would be beneficial at both early and delayed time points following stroke. With this in mind, the following section summarizes key findings related to the therapeutic potential of targeting LIF for ischemic stroke treatment.

2.12 LIF as a Therapeutic Target for Ischemic Stroke

The prosurvival pathways elicited by LIF, particularly the PI3K/Akt pathway, are now regarded as promising targets for the treatment of any injury involving oxidative injury, tissue necrosis, and apoptotic signaling. The promise of LIF has been demonstrated in several experimental injury models, including those that produce white matter injury. Data showed that exogenous LIF administration reduced demyelinating injury in experimental autoimmune encephalomyelitis (EAE), a model that recapitulates white matter damage reminiscent of multiple sclerosis, by preserving oligodendrocytes (OLs) [78]. Additionally, LIF effectively improved outcomes following spinal cord transection in mice [79]. The protection afforded in the latter case was attributed to activation of JAK/STAT and Akt signaling and was also associated with the induction of anti-apoptotic proteins. Similarly, LIF reduced motor nerve degeneration in the SOD1 G93A murine model of familial amyotrophic lateral sclerosis (ALS) [80].

Importantly, the benefits of LIF treatment observed in these earlier studies have now been extended to models of cerebral ischemia. In these recent studies, similar pathways appear to mediate the protective effects of LIF. Data showed that LIF effectively decreased infarct volume and white matter injury when administered to rats following pMCAO. In addition to mitigating neural injury, LIF treatment also improved functional outcomes. Further investigation revealed a mechanism operating through Akt-mediated induction of the antioxidant protein peroxiredoxin 4 (Prdx4), as LIF efficacy in protecting OL primary cultures from OGD was negated by co-incubation with Akt inhibitors and Prdx4 neutralizing antibodies [81]. Although neuronal cultures were not used in these studies, neurons and OLs are both susceptible to oxidative injury and therefore benefit from the induction of antioxidant proteins in the wake of ischemia. Likewise, stroke injury includes OL cell death and neuronal injury results, in part, from anoikis (detachment-induced cell death) following the disruption of astroglial junctions.

The therapeutic potential of LIF, although promising, lies in the ability to target LIF signaling at time points prior to maximal infarction. Since the infarct expands from hours to days following stroke onset, it is likely that the therapeutic window can be extended providing that the LIF safety profile is acceptable. To date, LIF has been used clinically for separate indications and has demonstrated a good safety profile. Nevertheless, a better option would be to stimulate the endogenous pathways

activated by LIF (i.e., Akt signaling) as means of circumventing potential issues with safety and dosing. Since LIF can be released by several cell types following injury [77, 82, 83], it will be crucial to characterize the sources of endogenous LIF signaling. Once the temporal and cell-specific responses are uncovered with regard to LIF signaling, there will be more therapeutic options to mimic the endogenous protection afforded by this cytokine.

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Part IV
White Matter Injury and Repair in Stroke

Ischemic Injury to White Matter: An Age-Dependent Process

Sylvain Brunet, Chinthasagar Bastian, and Selva Baltan

Abbreviations

AMPA/KA	AMPA/kainate
CKA	7-Chlorokynurenic acid
CNS	Central nervous system
GS	Glutamate synthetase
KB-R	2-[2-[4(4-Nitrobenzyloxy)phenyl]ethyl]isothiourea mesylate
MON	Mouse optic nerve
NCX	Na ⁺ -Ca ²⁺ exchanger
NMDAR	NMDA-type receptors
OGD	Oxygen glucose deprivation
RNS	Reactive nitrogen species
ROS	Reactive oxygen species
WM	White matter

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1 Introduction

In the United States, someone experiences a stroke every 40 s [1]. Axonal injury and dysfunction are responsible for most of the disability observed after a stroke [2] and aging is one of the most significant risk factors for stroke. The human brain comprises equal proportions of gray matter and white matter (WM) and WM is injured in most strokes [2]. However, most research efforts have traditionally been dedicated to protecting the gray matter. While effective in rodents, this approach has failed to translate to humans. Many reasons may underlie this failure, but major differences between humans and rodents include the greater proportion of WM in the human brain and the lack of significant WM involvement following middle cerebral artery occlusion in rodents, which is one of the most widely used animal models of stroke. Thus, over the past several years we and others have focused on examining how WM responds to ischemic injury, with an emphasis on the impact of aging [3–5].

2 WM Is Sensitive to Ischemic Injury

WM axons are dependent on a constant supply of oxygen and glucose to transmit signals. Central nervous system (CNS) WM electrical function is remarkably tolerant to anoxia [6], while there is regional heterogeneity in the ability to function and survive anoxia [7]. On the other hand, young adult WM is susceptible to ischemia induced by combined oxygen and glucose deprivation (OGD, Fig. 1). Mechanisms underlying ischemic WM injury proved to be unpredictably complex (Fig. 2) [8–16]. WM is composed of axons, oligodendrocytes, microglia, and astrocytes [2, 17]. Axons are myelinated by oligodendrocytes and exhibit patterns/gaps called nodes of Ranvier. Astrocytes support axons metabolically and restore the extracellular ionic environment following axonal activity. Microglia is partly responsible for immune surveillance. Thus, WM is composed of a complex cellular environment in which glial cell–cell interactions intricately maintain axon function. During ischemia, WM cellular elements are individually under attack but remain interactive with each other in intricate mechanisms that are currently under investigation.

3 Mechanisms of WM Ischemic Injury

It is now well-established that ischemia in WM sequentially activates three different injury pathways: the ionic, the excitotoxic, and the oxidative stress injury pathways. The ionic pathway attacks axons by collapsing their ionic homeostasis, which is initiated by the failure of the $\text{Na}^+\text{-K}^+$ pump, cell membrane depolarization, Na^+ channel activation, reversal of the $\text{Na}^+/\text{Ca}^{2+}$ exchanger, and Ca^{2+} channel activation, resulting in the accumulation of intracellular Na^+ and Ca^{2+} (Fig. 2, yellow) [18–23]. This increased intracellular Na^+ leads to the reversal of the Na^+ -glutamate transporter and the release of glutamate from astrocytes [16].

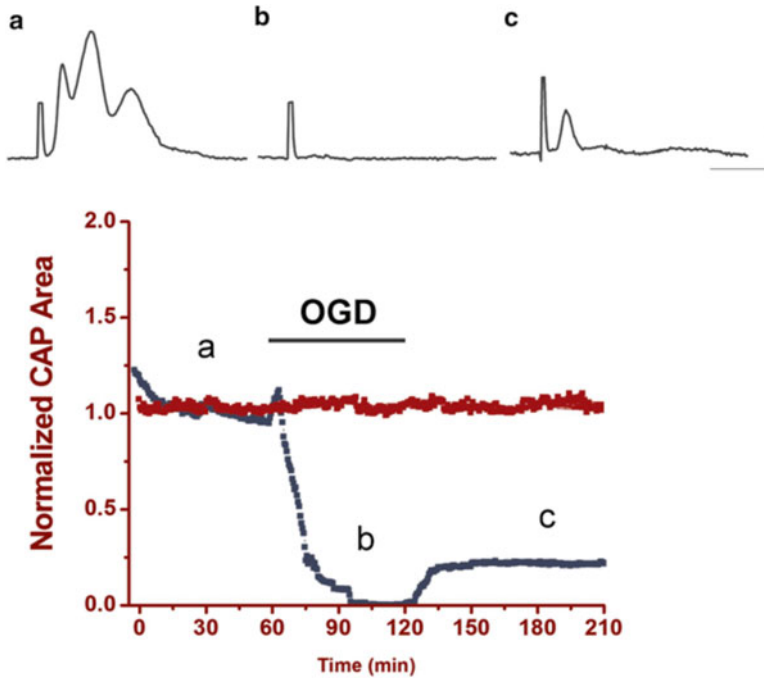


Fig. 1 WM is susceptible to ischemic injury. Axon function is quantified as the area under the compound action potential (CAP), normalized to control area, and plotted against time. Under normal conditions, CAP area is stable for long periods of time (*brown*). A 60 min period of OGD gradually depresses CAP area until conduction along axons is completely lost (*gray*). Restoring oxygen and glucose leads to ~25% axon function recovery. Sample traces from control (*a*), OGD (*b*), and recovery (*c*) periods are shown above the graph (Reproduced in part from Baltan (2014) [18])

Subsequently, the excitotoxic pathway is initiated by an increase in extracellular glutamate (Fig. 2, green). The excitotoxic pathway mainly targets oligodendrocytes by overactivating AMPA/KA receptors [11, 14, 16, 24–27] (redox) [28], which leads to increases in intracellular Na^+ and Ca^{2+} and the activation of downstream toxic intracellular pathways to mediate WM injury.

In parallel with the excitotoxic pathway, increased extracellular glutamate also leads to the activation of the oxidative stress pathway (Fig. 2, blue). The oxidative pathway damages WM components due to the formation of reactive oxygen species (ROS), which arises because the increased extracellular glutamate competes with cysteine at the glutamate-cysteine pump [29], depleting intracellular cysteine to reduce glutathione levels [30] and to cause mitochondrial dysfunction. In addition, the increase in intracellular Ca^{2+} activates NOS to produce nitric oxide [31, 32], which readily reacts with ROS to produce reactive nitrogen species (RNS). ROS and RNS can then attack multiple cellular elements (phospholipids, proteins, DNA, RNA) to mediate injury.

The ionic pathway triggers the injury process, which subsequently reverses the glutamate transporter; however, it is the accumulation of glutamate that dictates the threshold for irreversible injury. Therefore, if the injury is short and only involves

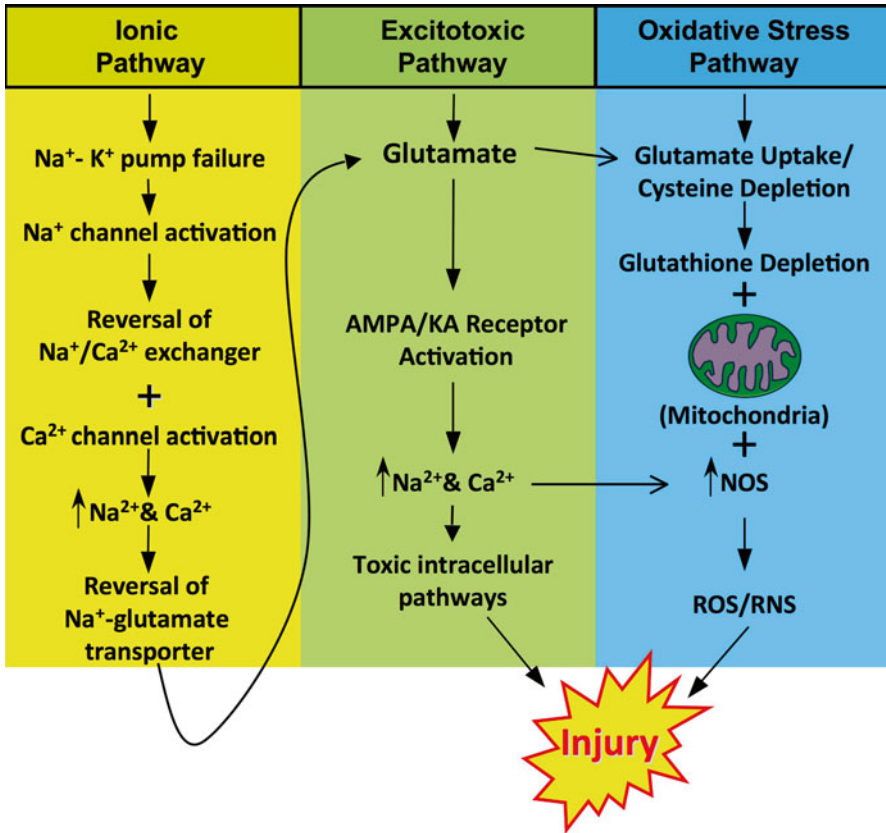


Fig. 2 Ischemia activates three pathways to mediate WM injury. Ischemia leads to the activation of the ionic pathway, which then leads to the sequential activation of the excitotoxic and oxidative stress pathways, which converge to cause irreversible injury to WM during ischemia. Note that glutamate release due to reversed Na⁺-dependent transport dictates the irreversible nature of the injury. *ROS* reactive oxygen species (Reproduced in part from Baltan (2009) [2])

the ionic pathway, then the ischemic injury is completely reversible. Unlike gray matter, this sequential order of events is necessary for the injury to develop, such that bypassing the ionic pathway and applying exogenous glutamate (or glutamate analogues) fails to cause WM injury [16].

4 Mouse Optic Nerve: An Ideal Model to Investigate WM

The mouse optic nerve (MON) is ideal for ischemic studies of WM. The optic nerve, the second cranial nerve, is a purely myelinated central nervous system WM tract and is sensitive to ischemia and to the aging process [33, 34]. In addition, tissue

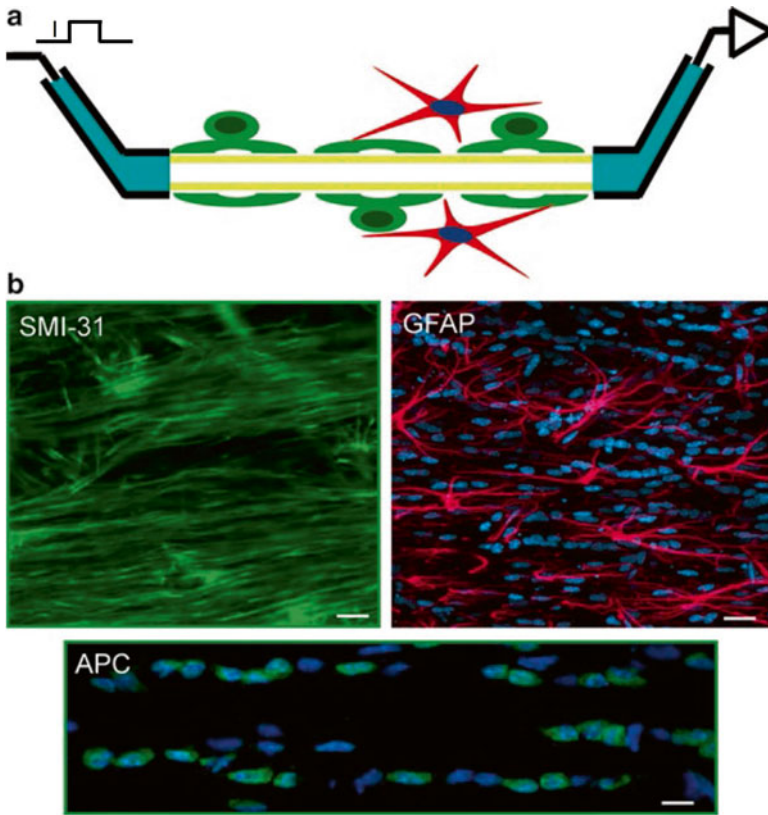


Fig. 3 The MON model is ideal for monitoring WM electrical function and cellular architecture. (a) Use of suction electrodes allows all axons to be stimulated and a CAP to be recorded. Cartoon of mouse optic nerve between two suction electrodes, where the left suction electrode stimulates and the right suction electrode records the CAP. Axons are represented in yellow, oligodendrocytes are in green, and astrocytes are in red. *I* current. (b) Using cell-specific antibodies, WM axons are labeled with SMI-31 for neurofilament (green), GFAP for astrocytes (magenta), and APC for mature oligodendrocyte cell bodies (green). Sytox (+) glial nuclei are in blue. Scale bar= 50 μm for SMI-31 and GFAP, 10 μm for APC (Reproduced in part from Baltan (2014) [18])

isolation does not require extensive surgical interventions; therefore there is negligible surgical injury, its small diameter allows sufficient glucose diffusion [6, 35], there are no neurons or synapses to contribute indirectly to the ischemic injury, and electrical function can be monitored by recording evoked compound action potentials (CAPs). MONs are stable both structurally and electrically for long durations (18 h) and glial cells and axons retain their native relationships to one another within a three-dimensional spatial organization (Fig. 3a). Furthermore, the cellular components can reliably be identified using immunohistochemistry (Fig. 3b), glutamate release can be measured by HPLC [36], proteins of interest can be quantified by Western blot analysis [3], and intravitreal injections provide a path for axonal

delivery [37]. Finally, the current and future availability of genetically engineered mice to experimentally test the role(s) of specific injury pathway components make the MON a model ideally suited to study WM injury mechanisms.

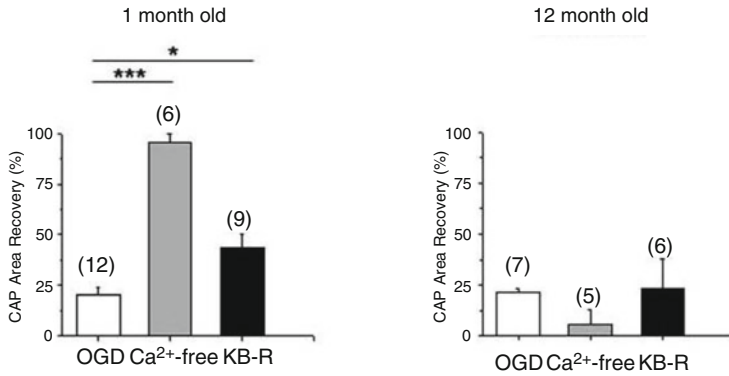
5 Why Is Aging WM More Susceptible to Ischemic Injury?

5.1 A Ca^{2+} -Independent Excitotoxicity Pathway in Aging WM

In aging WM, there is an enhanced contribution of a Ca^{2+} -independent excitotoxicity pathway to ischemic injury. In young WM, intracellular Ca^{2+} accumulation is important in the development of WM ischemic injury, and removal of extracellular Ca^{2+} , or blockade of Ca^{2+} entry via reversal of the Na^+ - Ca^{2+} exchanger (NCX) with 2-[2-[4(4-nitrobenzyloxy)phenyl]ethyl]isothioureia mesylate (KB-R), improves WM electrical function (Fig. 4A, left). In contrast, in older WM these interventions fail to improve injury. In fact, OGD applied in Ca^{2+} -free conditions harms recovery, suggesting that extracellular Ca^{2+} entry does not significantly contribute to ischemic injury in aging WM (Fig. 4A, right). In addition, the lack of protection afforded by blockade of Ca^{2+} entry secondary from the reversal of the NCX confirms the Ca^{2+} -independent nature of ischemic injury in aging WM. On the other hand, in aging MONs, it is possible that Ca^{2+} release from intracellular Ca^{2+} stores becomes accentuated during ischemia. However, the exact relationship between transmembrane Ca^{2+} influx and intracellular Ca^{2+} concentration, and the role of intracellular Ca^{2+} stores in aging WM, all remain to be established. Though these results suggest an attenuated role for the ionic pathway with aging, whether WM injury can be triggered directly at the excitotoxic pathway needs to be tested.

Overactivation of AMPA/kainate (AMPA/KA) receptors mediates excitotoxic injury in both young and aging WM. Thus, blockade of these receptors protects axon function against OGD in both young and aging WM. Interestingly, unless combined with a brief period of OGD [16], activating AMPA/KA receptors with glutamate (or glutamate agonists) does not cause WM injury. This is presumably because of efficient uptake of glutamate from the extracellular compartment by Na^+ -dependent transporters, such as GLT-1, as long as energy supply is maintained (Figs. 2 and 5). These findings suggest that OGD needs to activate ionic dysfunction to prime WM to the toxic effects of glutamate [16]. In aging WM, the Ca^{2+} -independent nature of excitotoxicity raises the question as to how activation of AMPA/KA receptors, which plays a prevalent role during ischemia in older WM, mediates injury. Either OGD-induced injury in older WM is specific to Ca^{2+} entry through AMPA/KA receptors or Na^+ entry through AMPA/KA receptors. The Ca^{2+} -mediated neurotoxicity requires distinct signaling pathways, such that some pathways are more efficiently triggered when Ca^{2+} ions enter at specific entry points such as the Ca^{2+} -permeable glutamate receptors [38]. In addition, overload of neurons by intracellular Na^+ exhibits irreversible neurotoxic swelling in the absence of extracellular Ca^{2+} [39]. Moreover, an increase in intracellular Na^+ leads to the reversal of the Na^+ -dependent glutamate transporter, leading to an increase in extracellular

A Ionic Pathway



B Excitotoxic Pathway

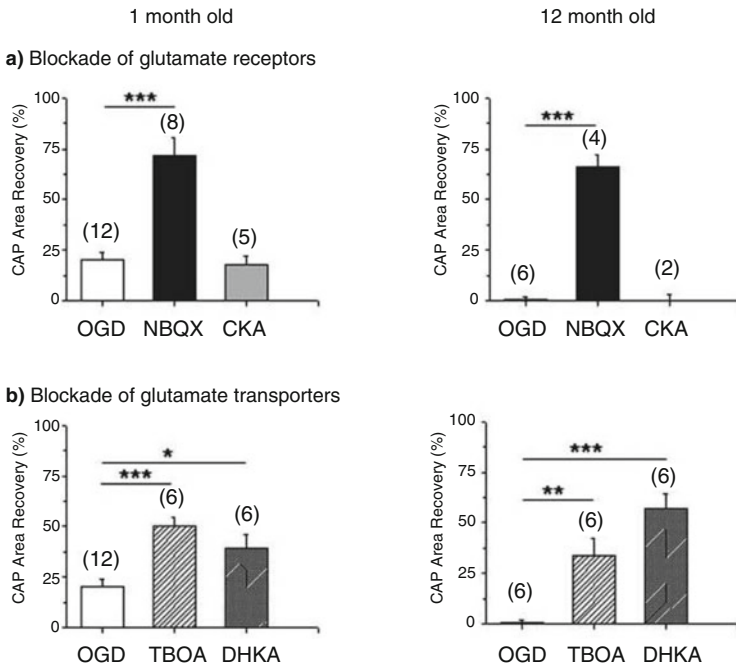


Fig. 4 Ca²⁺-independent excitotoxicity mediates ischemic injury in older WM. **(A)** The role of the ionic pathway in ischemic injury diminishes in older WM. Removal of extracellular Ca²⁺ or blockade of Ca²⁺ entry via reversing the Na⁺-Ca²⁺ exchanger (NCX) with KB-R (2-[2-[4(4-nitrobenzyloxy) phenyl]ethyl]isothiourea mesylate) is protective of axon function in 1-month-old WM after 60 min of OGD (*left*). KB-R failed to improve axon function in 12-month-old WM, even after 45 min of OGD (*right*). Note the reduced recovery in Ca²⁺-free conditions of 12-month-old WM (*right*). **(B, a)** Blockade of AMPA/KA receptors with NBQX (2,3-Dioxo-6-nitro-1,2,3,4-tetrahydrobenzof[*l*] quinoxaline-7-sulfonamide) provides identical protection to axon function in 1- and 12-month-old WM after 60 min of OGD. Blockade of NMDA-type receptors (NMDARs) with chlorokynurenic acid (CKA) does not improve axon function recovery in 1- or 12-month-old WM. **(b)** Blockade of Na⁺-dependent glutamate transporters improves axon function recovery in 1- and 12-month-old WM. Note that dihydrokainic acid (DHKA) afforded greater protection in aging axons. CAP compound action potential; TBOA (DL-threo- β -benzyloxyaspartate). **p*<0.05, ***p*<0.01, ****p*<0.001, two-way ANOVA. Data replotted from Tekkok et al. (2007) [16] and Baltan et al. (2008) [3] (Reproduced from Baltan (2009) [2])

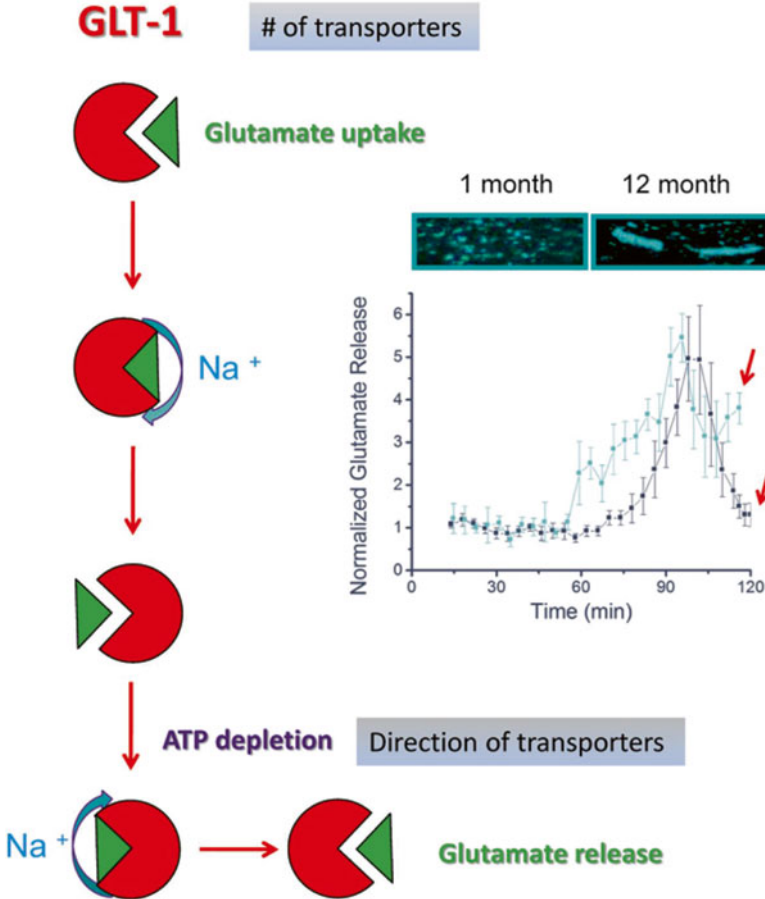


Fig. 5 In aging WM, enhanced excitotoxicity due to impaired mitochondrial function leads to early and robust glutamate release. The principal Na^+ -dependent glutamate transporter, GLUT-1, usually takes up glutamate with co-transport of Na^+ . During ATP depletion, due to increased intracellular Na^+ levels, the transporter reverses and releases glutamate. Therefore, the number of transporters determines the capacity of the system, but it is the ATP levels that determine the direction of the transport (to either remove or release glutamate). Mitochondria in aging axons become longer and thicker compared to young axons, which may hinder ATP production and drive GLUT-1 in reverse mode. Consistent with this, there is an early and robust glutamate release in aging WM (blue) compared to young MONs (gray). Note that the glutamate levels return to baseline in young WM but remain elevated in aging WM (red arrows) (Reproduced from in part from Baltan (2008) [3] and Baltan (2014) [18])

glutamate [40]. Together with the upregulation in GLUT-1 expression in aged WM [3], increased intracellular Na^+ may cause increased and early release of glutamate, overactivating AMPA/KA receptors and creating a vicious cycle that underlies the vulnerability of aging WM to ischemia.

The NMDA-type receptors (NMDAR) are activated during ischemia by glutamate [41], and the resultant increases in intracellular Ca^{2+} lead to myelin injury [42] or detachment of oligodendrocyte processes [43]. Consequently, axon function recovery would be predicted to improve following OGD if NMDARs are blocked. However, blockade of NMDAR in 1-month-old MONs (Fig. 4B, left) [16] or corpus callosum [14] does not improve axon function recovery following OGD. Likewise, blockade of NMDARs with 7-chlorokynurenic acid (CKA) does not protect axon function in older animals (Fig. 4B, right). Moreover, even after shorter durations of OGD (30–45 min), blockade of NMDARs worsens axon function recovery in older WM (data not shown). These results suggest that NMDAR activation during OGD does not contribute to failure of axon function and raises caution for the therapeutic use of NMDAR antagonists during ischemia, particularly in aging WM. These results do not negate activation of NMDARs under ischemic conditions, but suggest that their activation does not specifically contribute to axonal injury.

6 Reorganization of Glutamate Homeostasis in Aging WM

WM glutamate homeostasis and related regulatory proteins also go through age-related remodeling (Fig. 6). Expression of GLT-1, the dominant glutamate transporter, is upregulated in aging WM [3]. GLT-1 plays a key role in the removal of glutamate from the extracellular space to maintain glutamate below neurotoxic levels [44, 45]. Even though GLT-1 is predominantly expressed on astrocytes in young WM, it extends to additional structures with aging, implying that additional WM elements may contribute to the toxic glutamate accumulation in aging WM [3].

In addition to GLT-1, other essential elements for maintaining glutamate homeostasis include GLAST, glutamate, and glutamate synthetase (GS). WM glutamate content increases significantly with aging, which correlates with increased GS levels (Fig. 6). Together with a two-fold increase in GLT-1 levels in older WM [3], these adjustments may support an age-dependent adaptive mechanism in WM to remove glutamate from the extracellular space and to convert excessive glutamate to glutamine to maintain glutamate homeostasis. As a result, glutamate levels [3] and axon conduction across aging axons are stable under normal conditions.

The number of GLT-1 transporters determines the capacity of WM to move glutamate between the intracellular and extracellular space (Fig. 5). However, the direction of the GLT-1 transporter acts to either protect or injure WM. During ischemia, GLT-1 transporter reversal acts to injure the WM due to an accelerated Na^+ overload as a result of decreased tolerance to energy deprivation in aging WM (Fig. 7). Therefore, in old WM, more GLT-1 transporters are reversed, leading to earlier and more robust release of glutamate and enhanced excitotoxicity (Fig. 7). Moreover, in young WM, glutamate levels return to baseline after the end of OGD, which suggests efficient uptake of glutamate by astrocytes. However, in aging WM,

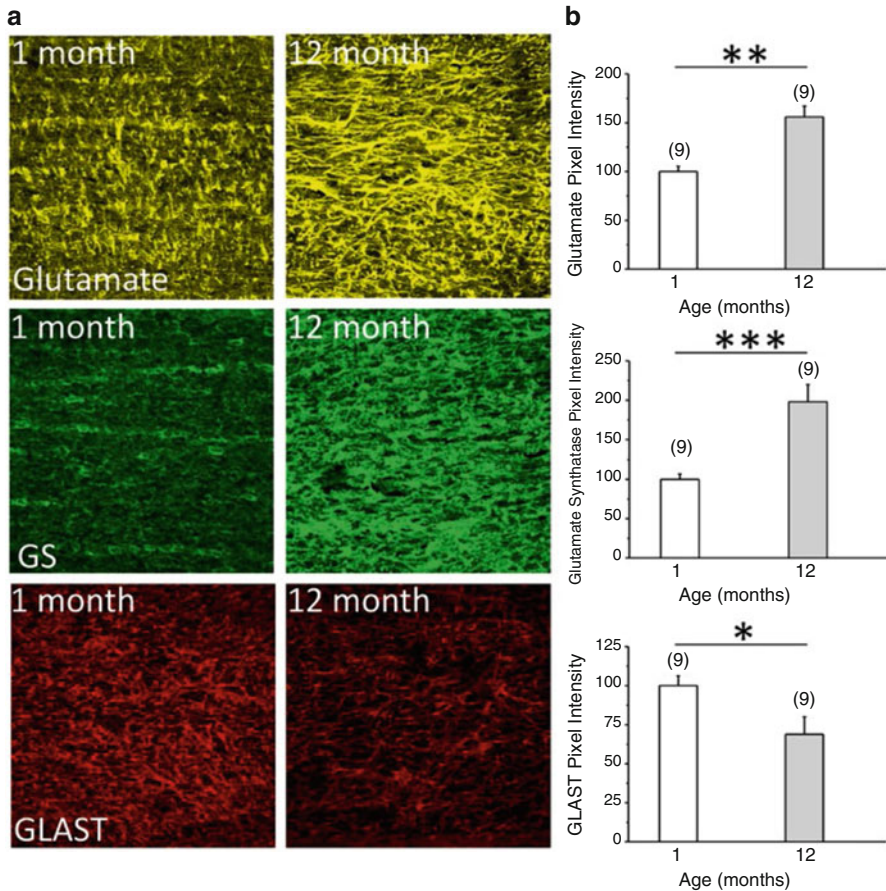
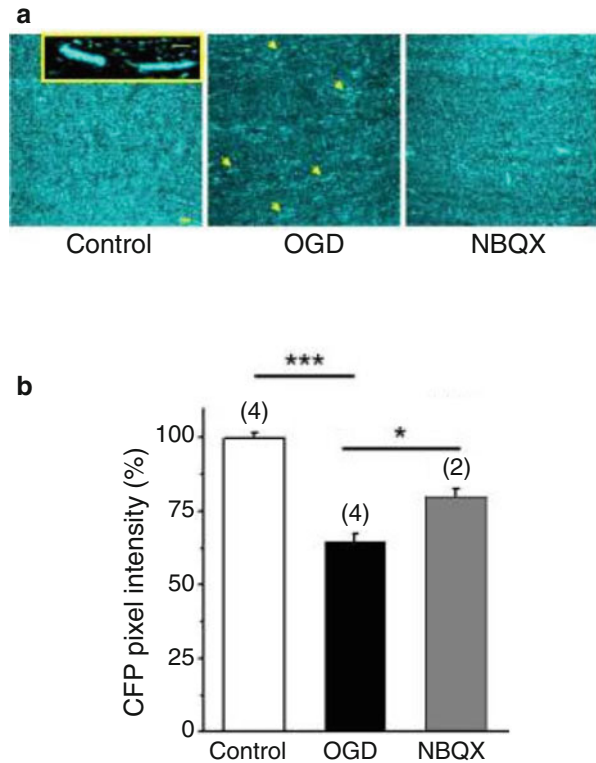


Fig. 6 Glutamate and glutamate synthetase (GS) expression are increased in 12-month-old MONs. **(a)** Immunolabeling and **(b)** quantification of glutamate, GS, and GLAST immunolabeling revealed that glutamate and GS labeling intensity is increased by $156 \pm 11\%$ and $198 \pm 22\%$, respectively, in 12-month-old MONs. Note that GLAST-labeling intensity decreased to $69 \pm 11\%$ in 12-month-old MONs. $*p=0.0278$, $**p=0.004$, $***p=0.0006$, two-tailed Student's *t*-test (Reproduced from Baltan (2014) [18])

glutamate levels remain elevated, suggesting that aging astrocytes cannot take up the excess glutamate and thereby extending the duration of excitotoxicity into the recovery period [3]. Despite the possibility that glutamate may be released from multiple sources in aging WM, astrocytes are expected to remove and store glutamate efficiently. Therefore, these results suggest a prominent change in aging astrocyte capacity to remove glutamate. On the other hand, GLAST expression in WM decreases with aging, raising the possibility that glutamate transporters can functionally substitute for one another with aging (Fig. 6).

Fig. 7 In aged MONs, blockade of excitotoxicity preserves CFP (+) axonal mitochondria. (a) CFP (+) mitochondria were longer and thicker in MONs from 12-month-old compared to 1-month-old Thy-1 CFP mice (inset; scale bar = 2 μ m). OGD reduced CFP pixel intensity in 12-month-old MONs despite longer and brighter CFP (+) mitochondria (yellow arrows). (b) Blockade of AMPA/KAR receptors with NBQX (30 μ M) preserved CFP pixel intensity during OGD. * $p < 0.05$, and ** $p < 0.0001$, one-way ANOVA. Calibration bar = 10 μ m (Reproduced in part from Baltan (2012) [4])



7 Mitochondria-Enhanced Excitotoxicity Underlies the Increased Vulnerability of Aging WM to Ischemic Injury

Mitochondria contribute to the enhanced excitotoxicity of aging WM and underlie the increased vulnerability of aging WM to ischemia. Mitochondria bioenergetics in neurons (gray matter) and their role in glutamate excitotoxicity are well-described [46]. Mitochondrial dysfunction and excitotoxicity share common features and are believed to act synergistically by potentiating one another [47–49]. Mitochondria are dynamic organelles that travel using axonal transport, in both the anterograde and retrograde directions, to reach peripheral locations and provide local energy supply [50, 51]. They constantly undergo fission and fusion events [52] and the relative rates of mitochondrial fusion and fission are implicated in the regulation of their size, number, and shape [53–55]. The balanced delivery of mitochondria to cell bodies, dendrites, axons, and axon terminals helps them serve multiple functions, including energy generation, regulation of Ca^{2+} homeostasis, cell death, and synaptic transmission and plasticity [56]. Expectedly, associations between many

neurological diseases, including aging, and defects in mitochondrial dynamics are emerging [55, 57].

In gray matter, aging neurons become more susceptible to glutamate excitotoxicity because of collapsed mitochondrial membrane potentials and increased generation of reactive oxygen and nitrogen species (ROS/RNS), leading to further reductions in mitochondrial function and energy production [58]. Mitochondrial function appears to decline in older animals, presumably causing reduced ATP production. This has been shown in the heart [59], liver [60], and brain [61]. Ion homeostasis accounts for $\approx 50\%$ of all ATP consumption, for which the Na^+/K^+ ATPase, the key enzyme to maintain ion homeostasis, is responsible for the majority of this consumption [62]. Therefore, the combined loss of ATP reserves and the high energy requirements of the Na^+/K^+ ATPase, diminishing the activity of this enzyme with aging, may heighten the sensitivity of aging WM to injury [63]. Consistent with this possibility, axon function in aging MONs, when transiently challenged with OGD, was slower to restore normal ion gradients, permitting pathological processes related to disruption of ion homeostasis to operate for a longer time (earlier reversal of the Na^+ -dependent glutamate transporter, Fig. 7), thus producing more injury in aging MONs [3]. In addition, aging MONs showed greater recovery of WM function in older animals when OGD was applied at a lower temperature [3], supporting the hypothesis that ATP reserves are compromised in aged WM. Finally, axons with higher ATP requirements have many more mitochondria per unit length of process [64]; therefore, these axons would be preferentially targeted by low ATP reserve conditions.

In neurons, excitotoxicity and elevated Ca^{2+} induce marked changes in mitochondrial morphology, stopping their movement [56, 65, 66] and generating ROS [46]. In young WM, activation of either AMPA or kainate receptors [3] loads mitochondria with Ca^{2+} and fission is enhanced, associated with loss of fluorescence of mitochondria genetically tagged with CFP (Fig. 8) [67]. Ca^{2+} overload activates NOS to produce nitric oxide and ROS/RNS, which could act either directly on oligodendrocytes [68–70] to cause injury or as diffusible second messengers linking oligodendrocyte excitotoxicity to axonal injury [31, 32]. Axon function directly correlates with WM energy reserves, since Na^+/K^+ ATPase activity is dependent on ATP levels. As a result, OGD causes a reduction in ATP levels and CFP (+) mitochondria, which is prevented by AMPA/KA receptor blockade in young and old WM (Figs. 8 and 9). Despite the structural and functional changes in aging mitochondria that enhance oxidative stress and amplify glutamate-mediated excitotoxicity during OGD, blockade of AMPA/KA receptors proved to be a successful strategy in improving WM function after stroke.

In summary, we and others have identified WM as an important therapeutic target for stroke [3, 5, 13, 15, 16, 41]. We have proposed the optic nerve as an ideal model for the study of ischemic WM injury and have described, step-by-step, ischemic injury pathways and the glial cells that are impacted. We have identified that aging modifies these ischemic injury pathways to render WM more susceptible to injury, such that in aging WM, the removal of extracellular Ca^{2+} is injurious, while

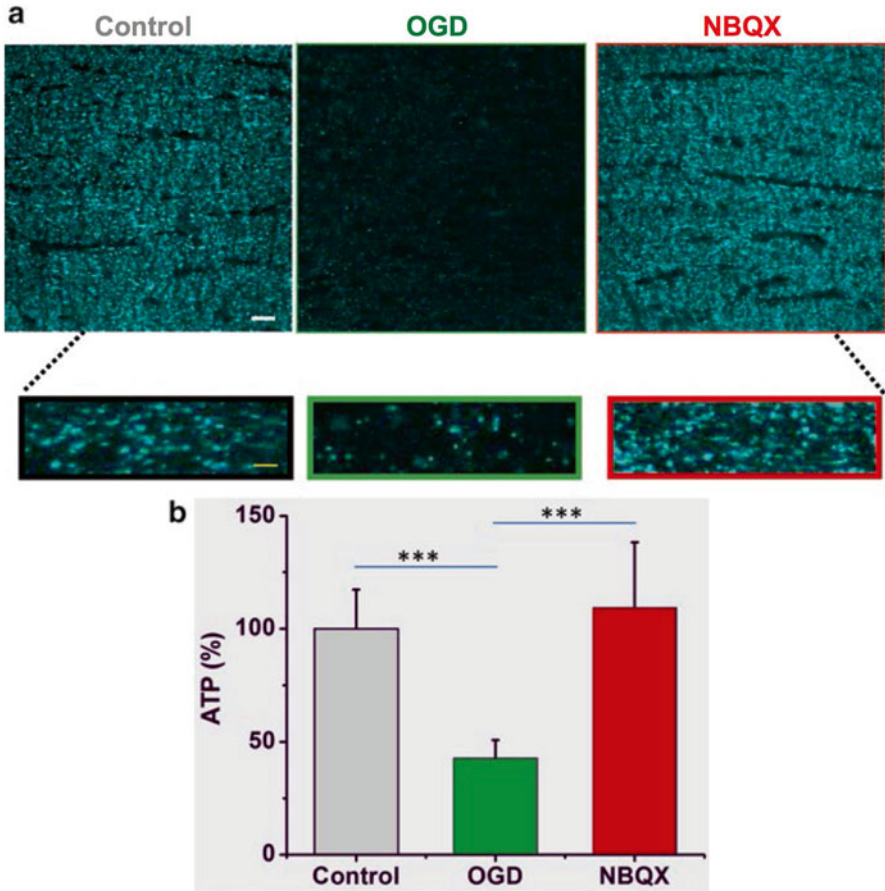


Fig. 8 In young MONs, Blockade of excitotoxicity preserves CFP (+) axonal mitochondria and ATP levels in response to OGD. **(a)** OGD severely reduced CFP fluorescence in MONs from mito CFP (+) mice and pretreatment with NBQX (30 μ M) protected against this loss. Note the change in mitochondrial morphology from small and tubular under control conditions to tiny and punctate following OGD. Calibration bar = 10 μ m (insets = 2 μ m) **(b)** NBQX pretreatment conserved ATP levels in MONs. *** p < 0.0001, one-way ANOVA (Reproduced in part from Baltan et al. (2011) [36])

it was protective in young WM (Fig. 9). In addition, aging alters WM glutamate homeostasis and mitochondrial dynamics, which lead to an enhanced glutamate excitotoxicity period with ischemia, which starts earlier and extends into the recovery period. Furthermore, we have identified that AMPA/KA receptor blockade protects both young and old WM, whereas NMDA receptor blockade is not protective in neither young nor old WM. Unexpectedly, NMDA receptor blockade worsened OGD recovery in old WM. Our results suggest that NMDAR activation during OGD

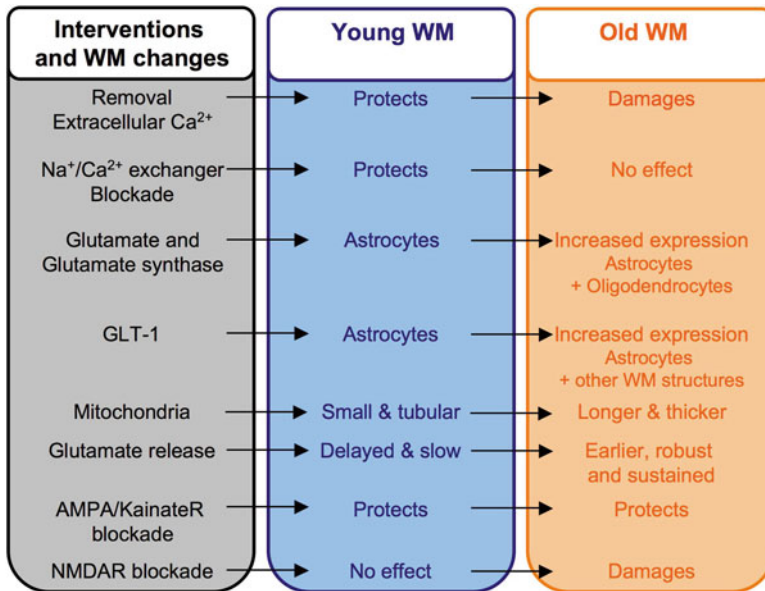


Fig. 9 Working molecular and cellular mechanisms responsible for ischemic WM injury. Age-dependent cellular remodeling of WM elements modifies the injury mechanisms and functional outcome to increase the sensitivity of aging WM to ischemic injury (Modified in part from Baltan 2014 [18])

does not contribute to failure of axon function and raises caution for the therapeutic use of NMDAR antagonists during ischemia, particularly in aging WM. Overall, our research suggests that for the development successful of future stroke therapies, we must tailor our approach to protect both gray matter and white matter as a function of age.

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Part V
Emerging Therapies to Target
Non-neuronal Mechanisms After Stroke

Neurovascular Repair After Stroke

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Abbreviations

[Ca ²⁺] _i	Intracellular calcium concentration
BBB	Blood–brain barrier
CBF	Cerebral blood flow

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CNS	Central nervous system
DCI	Delayed cerebral ischemia
EBI	Early brain injury
eNOS	Endothelial nitric oxide synthase
ET-1	Endothelin 1
ICAM-1	Intercellular adhesion molecule 1
ICH	Intracerebral hemorrhage
ICP	Intracranial pressure
IL	Interleukin
iNOS	Inducible nitric oxide synthase
mGluR	Metabotropic glutamate receptors
MMP	Matrix metalloproteinase
nNOS	Neuronal nitric oxide synthase
NO	Nitric oxide
NSC	Neural stem cell
PDGF	Platelet-derived growth factor
PPAR- γ	Peroxisome proliferator-activated receptor gamma
ROS	Reactive oxygen species
SAH	Subarachnoid hemorrhage
SGZ	Subgranular zone
SVZ	Subventricular zone
TGF- β	Transforming growth factor beta
TLR	Toll-like receptor
TNF- α	Tumor necrosis factor alpha
tPA	Tissue plasminogen activator
VCAM-1	Vascular cell adhesion molecule 1
VEGF	Vascular endothelial growth factor
VNN	Vascular neural network
VSMC	Vascular smooth muscle cell
ZO-1	Zona occludens 1

1 Introduction: Hemorrhagic Stroke and the Vascular Neural Network

Hemorrhagic strokes (subarachnoid hemorrhage (SAH), intracerebral hemorrhage (ICH), germinal matrix hemorrhage (GMH)) result from weakened vessels that rupture and bleed into the surrounding brain, leading to the accumulation of blood, an increase in intracranial pressure (ICP), and compression of the surrounding tissue. Hemorrhagic strokes also result in an immediate or postponed reduction in cerebral blood flow (CBF) [1]. These phenomena have important effects on the vasculature, and successful recovery from stroke requires a functional cerebral circulation; thus, vascular cells play an important role in the sequence of events in hemorrhagic

stroke. A comprehensive model that can be used to evaluate the pathophysiology and repair of hemorrhagic stroke is the vascular neural network (VNN), as proposed by Zhang and colleagues [2]. The VNN includes all the structures that are required to maintain CBF under physiological and pathological conditions, and encompasses vascular smooth muscle cells (VSMCs), arterial endothelial cells, and perivascular nerves connected to arteries and arterioles upstream of the cerebral microcirculation [2, 3]. This network is an extension of the neurovascular unit, which is comprised of neurons, glial cells (astrocytes, microglia, perivascular macrophages), pericytes, endothelial cells, and extracellular matrix, whose primary function is maintaining the homeostasis of the brain's microenvironment by mediating communication between the central nervous system (CNS) and the vascular network [4]. While the neurovascular unit model focuses on the areas immediately surrounding the capillaries, the VNN includes both venous downstream vasculature and upstream arteries and arterioles, and is therefore more inclusive for the mechanisms of vascular injury and impaired reperfusion induced by stroke [2].

The blood–brain barrier (BBB), which is between the CNS and the vasculature, is a part of the neurovascular network and is a highly organized multicellular complex that serves to control many metabolic and signaling actions occurring between the CNS and the rest of the body. It was originally thought that the barrier was principally mediated by tight junctions in endothelial cells [5]. However, it is now known that astrocytes, glial cells, and pericytes play important roles in BBB function [6, 7]. Consequently, any disruption of signaling between these cells and the cerebral endothelium can interfere with the integrity of the BBB and the brain microvasculature [8]. Many current therapies for hemorrhagic stroke are targeted at BBB repair, as loss of BBB integrity is one of the hallmarks of vascular disruption.

An understanding of the mechanisms regulating interactions among cells in the VNN is thus essential for the development of effective therapies against hemorrhagic stroke. In fact, both the beneficial and adverse effects of tissue plasminogen activator (tPA), the only Food and Drug Administration-approved treatment for ischemic stroke, have been linked to its interaction with components of this network [9]. Treatment with tPA was shown to attenuate early brain injury (EBI) after SAH by improving CBF and cortical perfusion, which are reduced due to increased ICP [10]. However, tPA is known to increase the levels of matrix metalloproteinases (MMP)-9 and 3, via the lipoprotein receptor (LRP)-1 in endothelial cells and astrocytes, which in turn increases the occurrence of hemorrhagic transformation in ischemic stroke [11–13]. Niogo et al. reported a concentration-dependent increase in BBB permeability with tPA administration *in vitro*; tPA also induced changes in the morphology of brain endothelial cells and astrocytes, and its modulation of the astrocytic skeleton via plasmin formation led to a decrease in BBB integrity [14]. Furthermore, the interaction of tPA with LRP has been shown to play a role in the permeability of the neurovascular unit by promoting the detachment of astrocyte endfoot processes [15, 16], and MMPs themselves are known to degrade components of the neurovascular matrix [17]. (Figure 1 outlines the pathophysiology of hemorrhagic stroke, illustrating the interactions of the various cell components).

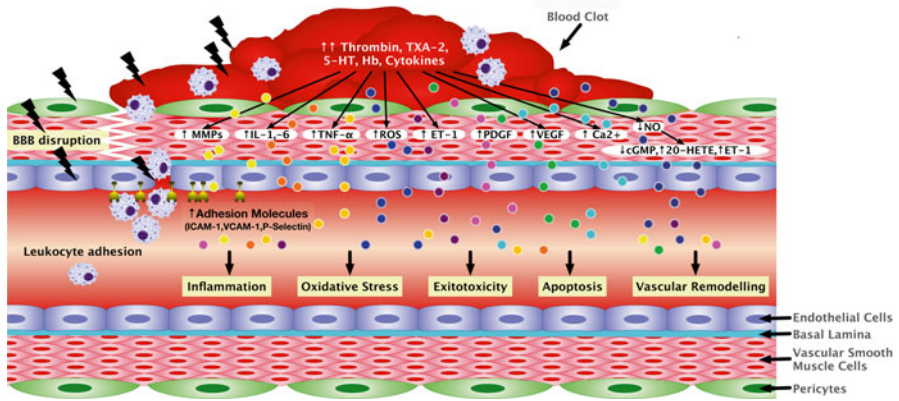


Fig. 1 Pathophysiology of hemorrhagic stroke. Vascular inflammation, oxidative stress, excitotoxicity, apoptosis, and vascular remodeling contribute to neurovascular injury. After stroke, thrombin, thromboxane-2 (TXA-2), serotonin (5-HT), hemoglobin, and cytokines are released. In response, signaling pathways are activated, leading to increased matrix metalloproteinases (MMPs), inflammatory cytokines (interleukin (IL)-1, IL-6, tumor necrosis factor alpha (TNF- α), reactive oxygen species (ROS), endothelin-1 (ET-1), platelet-derived growth factor (PDGF), vascular endothelial growth factor (VEGF), intracellular calcium (Ca^{2+}), and decreased nitric oxide (NO), cyclic guanosine monophosphate (cGMP), and the eicosanoid hydroxyicosatetraenoic acid (HETE). These are the main factors inducing vascular injury. Additionally, the inflammatory response leads to increased expression of intraluminal adhesion molecules (intercellular adhesion molecule 1 (ICAM-1), vascular cell adhesion molecule 1 (VCAM-1)), and P-Selectin, resulting in leukocyte adhesion, and eventually, blood–brain-barrier (BBB) disruption

2 Hemorrhagic Stroke-Induced Neurovascular Disruption: How VNN Interactions Are Altered During Stroke

2.1 Heme Breakdown Leads to Neurotoxicity

Due to the buildup of potentially neurotoxic iron-rich blood, and the products of heme breakdown, neuronal injury and severe cognitive impairment occur following a hemorrhagic stroke [18–24]. Additionally, thrombin, erythrocyte lysis, toll-like receptor (TLR)-4 activation, and other blood components induce activation of microglia and other inflammatory agents and lead to the recruitment of leukocytes into a normally immune privileged area [25–28]. Thus, limiting the effects of this toxicity by hematoma clearance, presents potential for protecting the neurovascular unit after hemorrhagic stroke [29, 30]. In fact, Zhao et al. reported that peroxisome proliferator-activated receptor (PPAR)- γ activation resulted in resolution of the hematoma following murine ICH, which further led to decreased neuronal damage and improved functional outcomes [31].

Heme is degraded into biliverdin, iron, and carbon monoxide by heme oxygenases (HO). HO-1 is the ubiquitously expressed isoform of the enzyme that possesses cytoprotective abilities. Schallner, et al. found that substituting for the absence of

microglial HO-1 by inhalation of carbon monoxide after SAH reduced injury by erythrophagocytosis, and that HO-1 is necessary to attenuate neuronal cell death, vasospasm, impaired cognitive function, and clearance of blood in the subarachnoid space [32]. HO-1 clears the hematoma and thus reduces the oxidation burden and neuronal injury via microglial phagocytosis, similar to the action of PPAR- γ .

2.2 *Excitotoxicity and the Role of Astrocytes*

Glutamate excitotoxicity is another mechanism that induces neuronal death after hemorrhagic stroke. As such, a recent study found that inhibition of glutamate α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptors was neuroprotective in EBI after SAH [33]. Cerebral glutamate levels increase within minutes after SAH, and excitotoxicity through the glutamate receptors is thought to play an important role in SAH pathophysiology [34]. EBI involves vasogenic edema and apoptotic cell death, and is compounded by inflammation, oxidative stress, and ionic disturbances, which leads to BBB disruption, edema, and neuronal apoptosis [35]. These events, along with delayed cerebral ischemia (DCI), cortical spreading depression, and development of microthrombi, impact glutamate receptor physiology following SAH. Microthrombi accumulate in the parenchymal vessels after SAH, and platelet aggregation is critical to their formation [36]. Platelets are released into the brain following the breakdown of vessel walls in SAH, which they help to initiate, and are thought to release glutamate as part of their cell–cell signaling [37, 38]. Indeed, Bell et al. recently found that when platelets were activated with thrombin, glutamate release was significantly increased, which led to an increase in neuronal cell death in vitro due to the proximity of neurons to microthrombi. Interestingly, they also found that platelet-rich plasma downregulated glutamate receptor 2 surface expression (which was likely downstream of excessive receptor stimulation) [37]. This is significant as glutamate receptors play an integral role in memory consolidation and executive functioning.

Many cell types in the CNS express glutamate transporters: astrocytes, neurons, oligodendrocytes, microglia, and the endothelium. However, the role of astrocytes in the transporter-mediated uptake of the neurotransmitter is the most important for maintaining normal glutamate levels, and reducing excitotoxicity, particularly following an insult to the brain [39]. Neuronal vulnerability to glutamate excitotoxicity is 100 times greater in cultures low in astrocytes than in astrocyte-rich cultures [40]. Additionally, while neurons do play a role in synaptic glutamate uptake, the role of astrocytes in this process seems significantly more important. Glutamate can also move in the opposite direction through the transporters. Since membrane electrical potentials and transmembrane ion gradients are maintained by sodium/potassium adenosine triphosphatases, changes in adenosine triphosphate (ATP) concentrations influence glutamate transport. Thus, since ATP levels are severely depleted in the case of cerebral ischemia, which occurs secondary to SAH, glutamate uptake is then reversed, and astrocytes release the neurotransmitter. Neurons can also undergo glutamate uptake reversal, thus increasing extracellular glutamate concentrations, resulting in greater neurotoxicity.

2.3 *Vascular Disruption and Constriction*

The entire cerebral circulation includes large arteries and arterioles, capillaries, and downstream venules and veins. The increased heme and heme breakdown products from the rupturing of vessels and the expanding hematoma during a hemorrhagic stroke were found in a recent study to induce vasoconstriction of cerebral vessels [41]. After SAH, following the usual reduction in CBF, there is a DCI, beginning between day 2 and 4 post-SAH, up to 14 days in humans [42, 43]. Posthemorrhagic vasospasm, induced by heme and its breakdown products, is what leads to DCI, which is also associated with spreading depression and microthrombosis [42]. Several reports have described acute vasoconstriction within hours after SAH, particularly small arteries and arterioles. In guinea pigs, topical blood application to the exposed brain cortex resulted in vasoconstriction of pial vessels [44]. The occurrence of superficial pial vessel constriction was observed predominantly in arterioles, though the time of occurrence varied between the perforation model and the blood-injection model. Nevertheless, smaller arterioles were more constricted than larger vessels in both models [45, 46]. Decreased blood flow was observed in venules as well [47], other reports have shown that venule diameter was unchanged after SAH [48].

The venous compartment of cerebral circulation contains about 70–80% of the circulating volume in the cranial cavity and is surrounded by adrenergic nerve fibers, while postcapillary venules are covered with pericytes [49, 50]. There are several differences between the venous and arterial systems, which affect the roles they play in hemorrhagic injuries. Small veins and venules lack smooth muscle cells; thus, they do not contract as extensively as arteries [51]. As a result, they do not function as much in the regulation of CBF, and only undergo small changes in diameter according to ICP under normal physiological conditions [52, 53]. However, in pathological conditions, the veins may be easily compressed by high ICP. Secondly, veins have thin walls and do not have valves to prevent the backflow of venous blood [54]. The severe brain edema that occurs due to hemorrhagic stroke leads to an increase in ICP that may compress or collapse the thin walls of the veins [55]. Swollen astrocyte endfeet and adherent leukocytes also induce compression of the venule walls. Deep cerebral vein vasospasm has been reported in a rabbit model of SAH [56]. This may be due to the presence of pericytes on the vessels, as contraction of pericytes is essential for active vein constriction [57]. Additionally, in a blood-injection SAH model, leukocyte-platelet aggregates were observed in the cerebral veins at 2 h after SAH [58]. The resultant oxidative stress and inflammation may induce damage to venous endothelial cells and trigger clot formation in the cerebral venous system [56]. This suggests that alterations to the venous system may play a significant role in hemorrhagic stroke-induced secondary brain injury (Fig. 2).

Beyond constriction, alteration in the reactivity and ultrastructural changes in the vessel walls also occur after SAH. For example, platelet activation releases vasoconstrictors, such as thromboxane-A₂, endothelin (ET)-1, serotonin (5-HT), transforming growth factor (TGF)- β , vascular endothelial growth factor (VEGF), and platelet-derived growth factor (PDGF) [59–61]. Furthermore, PDGF, TGF- β , and

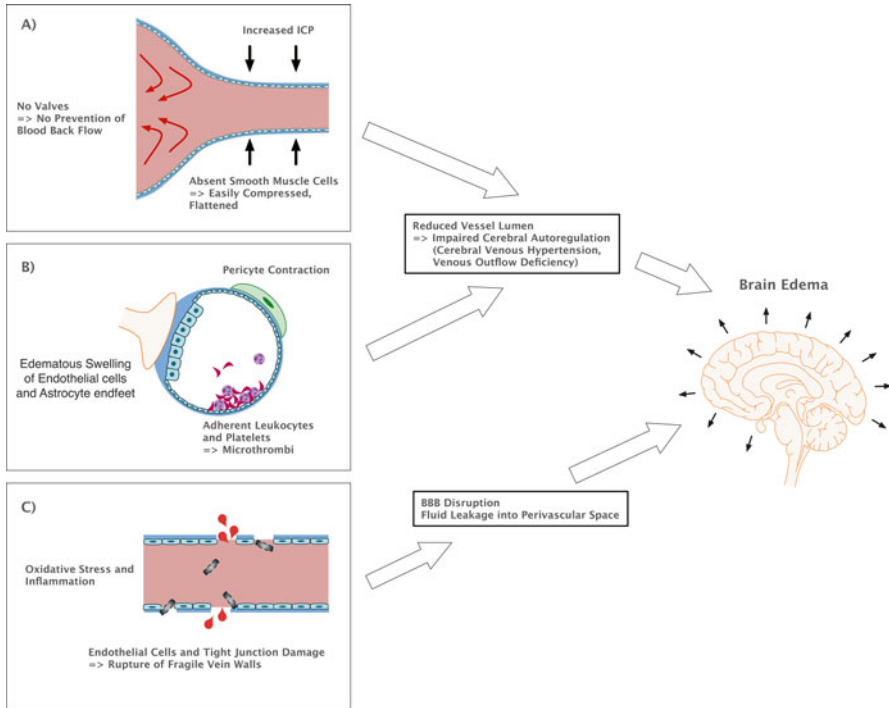


Fig. 2 Alterations in veins, venules, and capillaries after hemorrhagic stroke. (a) Increased ICP flattens and compresses veins easily compared to arteries due to the absence of smooth muscle cells. This leads to reduced vessel diameter and increased back flow of blood. (b) Edematous, swollen endothelial cells and astrocyte endfeet, active pericyte contraction, and adherent leukocytes and platelets also result in reduced vessel lumen. (c) Oxidative stress and inflammation induce endothelial cell and tight junction damage, leading to BBB disruption and brain edema

ET-1 promote proliferation and migration of smooth muscle cells located in the medial and adventitial fibroblasts. These changes may result in the synthesis of additional collagen, increasing the thickness and structural stiffness of the vessel wall [62]. ET-1, which is released from damaged endothelial cells, has also been shown to stimulate phospholipase C-induced extracellular calcium influx through calcium-permeable nonselective cation channels and store-operated calcium channels in vascular smooth-muscle cells, leading to contraction of arteries following SAH in a rabbit model [63]. Further, ET-1-induced vasoconstriction in SAH cells was blocked by inhibition of transient receptor potential canonical 1 and 4, which mediate calcium entry through nonselective cationic pathways, suggesting that ET-1 works by increasing calcium influx [64]. Moreover, ET-1 contributes to hemoglobin-mediated vascular injury via RhoA/Rho kinase and protein kinase C (PKC), which inhibits myosin phosphatase [65]. It was reported that inhibiting the ET-1-induced vasospasm after SAH led to reduced expression of VEGF, the most potent angiogenic factor; thus, the ischemic effect of ET-1-induced vasoconstriction enhances angiogenesis [66]. Figure 3 highlights some vasoconstrictors and vasodilators known to impact vessel diameter following hemorrhagic stroke.

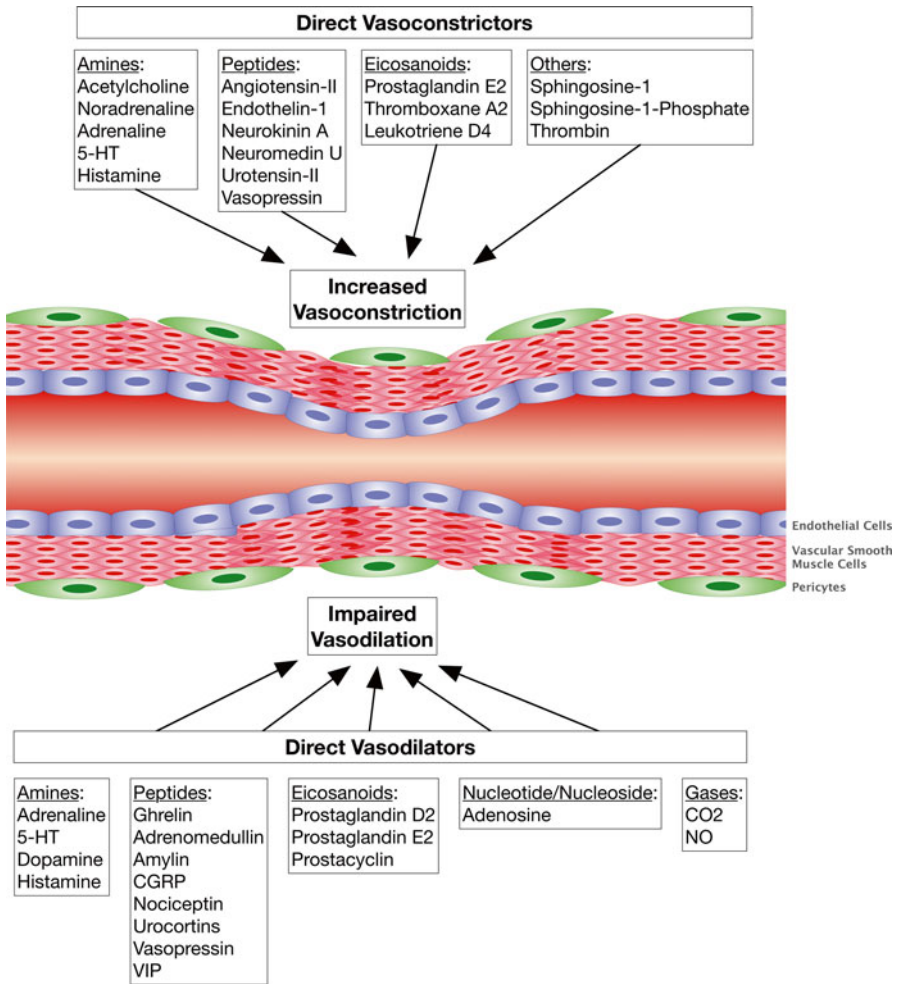


Fig. 3 Vasoconstriction and impaired vasodilation. Hemorrhagic stroke alters cerebral autoregulation, impairing vascular responses. After injury, there is an immediate or delayed reduction in cerebral blood flow due to constriction of cerebral vessels. Vasoconstricting mediators lead to significant contraction of the vessel lumen, decreasing blood flow, while the vasodilators, trying to restore perfusion to the injured area, can induce only restricted expansion of the cerebral vessels. 5-HT serotonin, CGRP calcitonin gene-related peptide, VIP vasoactive intestinal peptide, CO₂ carbon dioxide, NO nitric oxide

2.4 Vascular Injury and Blood Flow

Intertwined with the vascular disruption after hemorrhagic stroke is an increase in the permeability of the BBB. This is further enhanced by the hemoglobin-induced increase in oxidative stress by reactive oxygen species (ROS) and reactive nitrogen

species (RNS), including superoxide radical and nitric oxide [18], which, along with their metabolites, such as peroxynitrite, activate MMP-9, and disrupt the vascular matrix [67–69]. MMPs are upregulated in neurons and reactive astrocytes in hemorrhagic stroke, and play a role in mediating BBB leakage and edema by degrading the neurovascular matrix. Brain edema is the clinical observation of BBB disruption, and leads to a greater increase in ICP [70]. The increased ICP interferes with the flow of blood after injury by inducing constriction of the blood vessels.

With the induced vasoconstriction and vascular disruption after hemorrhagic stroke, the metabolic demands of the brain cannot be sufficiently met, and an increase in neuronal death may result. This is because the brain utilizes approximately 20% of cardiac output, and is extremely dependent on a constant supply of nutrients and oxygen via the blood, as evidenced by its extensive distribution of blood vessels. Furthermore, it is highly specialized to fulfill this function, such that, when neuronal activity increases in a specific brain region, blood flow increases to this area to meet the enhanced metabolic demand, a phenomenon referred to as functional hyperemia [71].

2.5 *Astrocytes in the Cerebrovascular Response*

Coupling of the neurovascular components to meet neuronal activity is regulated by various neurotransmitters [72], as well as astrocytes, which are known to have important effects on vessel diameter, as their endfeet are layered around over 99% of cerebral capillaries [53]. Furthermore, receptors for the vasoactive mediators released by neurons are found on astrocytes and microvascular endothelial and/or smooth muscle cells, and can either dilate or constrict cortical microvessels upon activation [73]. Thus, though vasodilation of parenchymal arterioles in response to neuronal stimulation can be severely compromised after hemorrhagic stroke [74], an increase in the number of reactive astrocytes (astrogliosis) after injury can increase vasodilation by the release of vasoactive molecules from the astrocyte endfeet, thus helping to maintain blood flow. In demonstrating the essential role of astrocytes (neuron-astrocyte signaling) in mediating the cerebrovascular response to neurons, Zonta et al. found that when neuronal afferents were stimulated by an extracellular electrode placed in the cortex, arterioles showed a transient relaxation, and interestingly, this also triggered an increase in intracellular calcium ($[Ca^{2+}]_i$) in the astrocytic endfeet surrounding the arterioles; these events occurred in a temporally related manner and links synaptic activity to the control of vascular tone [75]. They also reported that when the metabotropic glutamate receptors (mGluR) (known to mediate the $[Ca^{2+}]_i$ increase in astrocytes) were blocked, there was reduced dilation of the arterioles in response to afferent stimulation of the neurons. Conversely, direct stimulation of the mGluR produced an increase in vasodilation, correlated to the timing of $[Ca^{2+}]_i$ increase in astrocytes, and direct stimulation of the astrocytes also triggered dilation of the arteriole region that was in contact with the astrocytes' processes [75, 76]. Interestingly, the effect of mGluR stimulation implicates the involvement of synaptically released glutamate, the most abundant

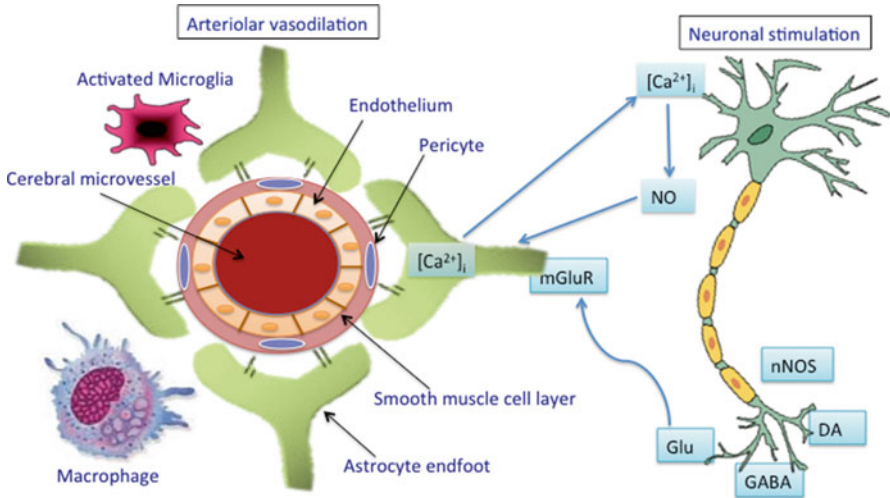


Fig. 4 Neurovascular coupling. Neuron-astrocyte crosstalk plays a crucial role in mediating neurovascular coupling and functional hyperemia. Astrocyte endfeet are layered around smooth muscle cells surrounding cortical microvessels, and upon afferent stimulation of neurons, there is an increase in intracellular calcium ($[Ca^{2+}]_i$) in the astrocyte endfeet, mediated by the metabotropic glutamate receptors (mGluR) upon binding of glutamate (Glu). The result is increased vasodilation of the arterioles ensheathed by the astrocyte endfeet processes. Glutamate release from neurons also induces calcium uptake, which activates neuronal nitric oxide synthase (nNOS), leading to release of nitric oxide (NO), a vasodilator. The NO released from neurons diffuses into the astrocytic endfeet where it modulates the release of vasoactive substances, and may also directly regulate smooth muscle cells. In addition, other neurotransmitters, such as dopamine (DA), and gamma aminobutyric acid (GABA) help to regulate cerebrovascular responses to neural activity. Microglia and macrophages are recruited to the damaged area following a stroke, inducing several inflammatory reactions. This can lead to microvascular injury. However, they also engulf the blood and damaged or dead tissues, providing an environment for tissue repair

excitatory transmitter in the CNS. The uptake of glutamate from the synapse by astrocytes, as discussed above, aids in reducing excitotoxicity by glutamate following a stroke, as the levels of the neurotransmitter are maintained by astrocyte activity [39, 77]. It also helps to drive glucose uptake from local blood vessels to fulfill the metabolic needs of the neurons via the astrocytic endfeet processes [78]. These data illustrate the unified action of cells of the VNN to ensure that the metabolic demand and blood flow requirements of the brain are met, and also that neuronal injury is reduced (Fig. 4).

Ionic imbalance is another major mechanism by which vascular smooth muscle constriction occurs after a hemorrhagic stroke. Intracellular calcium levels in VSMCs are elevated by either increased influx through voltage-gated calcium channels or impaired calcium reuptake in the sarcoplasmic reticulum [79–83]. After SAH, the reuptake is compromised by the lack of nitric oxide (NO), which, under normal conditions, would elevate cyclic guanosine monophosphate (cGMP) via calcium reuptake in smooth muscle cells [84]. Increased calcium levels induce opening of

calcium-activated potassium channels in astrocytes, which results in highly elevated extracellular potassium, leading to arteriolar vasoconstriction [46]. Additionally, increased expression of the voltage-gated potassium (Kv) channels in parenchymal arteriolar myocytes leads to depolarization and increased vascular tone via the heparin-binding epidermal growth factor-like growth factor [85]. Consequently, the Kv7 channels are common targets of vasoconstrictor spasmogens [86].

2.6 Pericyte-Endothelial Interactions and the Blood Brain Barrier

During injury, vascular function is regulated by various mediators that initiate the differentiation and migration of cerebrovascular pericytes, which are able to further modulate the structure and function of cerebral arteries and maintain BBB function [87]. These include VEGF, PDGF, nerve growth factor, angiopoietins, TGF- β , notch, etc. [4]. Extracellular TGF- β and intracellular Rho GTPase were previously reported to be involved in vasoconstriction. Several studies suggest that this contractile effect is due to pericytes [88–90]. Pericytes in the CNS have receptors for a large number of vasoactive mediators, thus they may play a role in cerebrovascular autoregulation. Electrical stimulation of cerebellar pericytes evoked localized capillary constriction in response to neural activity, indicating a role for pericytes in blood flow regulation during injury [91].

In a perforation-induced SAH model [92], electron microscopic evaluation revealed partially collapsed capillaries, which were associated with enlarged perivascular astrocyte foot processes and luminal endothelial cell protrusions, as well as defects in the basal lamina and endothelium [93]. Increased endothelial apoptosis has been shown in parenchymal vessels 10 min after SAH, up to 24 h [94, 95], and this can result in BBB disruption [96]. Several studies have reported decreased occludin and collagen IV in parenchymal vessels, associated with the disruption of endothelial tight junctions and widening of the inter-endothelial spaces [38, 97–99]. Additionally, BBB tight junction opening has been shown after hemorrhagic stroke, with the greatest opening occurring at 3 and 72 h, coinciding with the downregulation of the tight junction proteins zona occludens (ZO)-1 and occludin [100].

Pericyte–endothelial interactions are important for vascular integrity following a cerebral injury. Pericytes are located in precapillary arterioles, capillaries, and post-capillary venules, and they synthesize and deposit elements of the basal lamina [101]. In the mature capillary, they are located between endothelial cells and parenchymal astrocytes and neurons; thus, there is active crosstalk among these cells. The ratio of pericytes to endothelial cells is much higher in the cerebral microcirculation than in other tissues, and they play a role in microvascular blood flow as a final regulatory step after arterioles [102]. As such, pericytes on isolated retinal capillaries have been shown to constrict or dilate in response to neurotransmitters, due to an increase in intracellular calcium concentrations. Furthermore, pericyte numbers around endothelial cells have been found to inversely correlate with BBB function,

as loss of pericyte coverage leads to an increase in the gene expression of proteins known to increase vascular permeability (angiopoietin-2, intercellular adhesion molecule (ICAM)-1, etc.) [4]. Pericytes maintain BBB homeostasis by inducing the expression of occludin, maintaining the expression of other BBB-specific genes in endothelial cells, and by inducing polarization of astrocyte endfeet adjacent to the cerebral microvasculature [6, 103]. Antigen-presenting pericytes induce a local pro-inflammatory response mediated by the elevated expression of adhesion molecules (ICAM-1 and vascular cell adhesion molecule (VCAM-1)) near endothelial cells and circulating cytokines and chemokines, which are involved in neutrophil infiltration and tight junction degradation [104, 105].

2.7 Inflammation and Vascular Injury

Another contributor to microvascular injury after hemorrhagic stroke is inflammatory reactions, including leukocyte rolling and adherence to the walls of cerebral vessels [106, 107]. Coinciding with this, a progressive increase in leukocyte adhesion to venules was reported after SAH [58]. As a result, the adhesion molecule P-selectin was more expressed in the endothelial cell membrane following SAH, while no difference in cytosolic P-selectin expression was observed [47]. Other inflammatory mediators, apart from leukocytes, are also produced and elevated after a hemorrhagic stroke, including the cytokines interleukin (IL)-1 β , IL-6, tumor necrosis factor (TNF)- α , and high-mobility group box protein 1 [108, 109]. These cytokines, combined with the induced oxidative stress, also contribute to vascular injury and BBB disruption [110].

3 Mechanisms of Neurovascular Repair

3.1 Glial Cells in Hemorrhage Clearance

Resident microglia and peripheral macrophages are rapidly mobilized to the site of injury (around the hematoma), and recruit other immune cells, as well as initiate the release of effector molecules [111–113]. Microglial activation within the perihematomal region has been seen as early as 1 h after collagenase-induced experimental ICH, and within 4 h in the autologous blood double injection model [114, 115]. By 72 h post ICH, microglia reach their peak numbers in the perihematoma, then levels begin to fall back down by 7 days post-injury, and return to basal levels within 3–4 weeks [112, 116, 117]. This time course coincides with the hematoma volume over time in ICH (peaking at 72 h, remaining elevated up to 7 days, and returning to basal levels within 2–4 weeks) [118], suggesting that activated microglia and macrophages play a significant role in hematoma resolution [119]. Activated microglia recruit hematogenous phagocytes to the area of the hemorrhage; the phagocytes in

turn engulf the deposited blood, damaged and dead tissue, and then provide an environment for tissue repair [24]. Zhao et al. found that PPAR- γ treatment led to more efficient microglia-induced phagocytosis [31]. PPAR- γ treatment was also found to reduce IL-1 β , TNF- α , MMP-9, and inducible nitric oxide synthase (iNOS) expression, while increasing CD-36-mediated microglial phagocytosis of red blood cells [31, 118]. Another treatment targeted at microglial activation following ICH is minocycline, which was found to reduce microglia/macrophage numbers at 5 days when given 3 h post ictus [120]. It also reduced TNF- α , IL-1 β , and MMP-12 levels both in vivo and in vitro [121, 122]. These studies indicate an important role for glial cells in neurovascular repair.

Yet, there lies a paradox in the activity of microglia and macrophages: while they are clearing up the hematoma, they generate large quantities of oxidant by-products, which increase oxidative stress to the microglia, and may limit their functioning [25], and could also exacerbate tissue damage. This double-sided action on brain repair and tissue damage may be due to the polarization of microglia and macrophages toward different phenotypes at different stages of injury [123]. M1 microglia/macrophages are the classically activated phenotype, and are considered to be pro-inflammatory, secreting TNF- α , iNOS, IL-1 β , etc., while M2 microglia/macrophages represent the alternatively activated phenotype and are seen as healing cells involved in neuroprotection and repair via arginase activity and upregulation of neurotrophic factors [25, 124, 125]. Macrophages and microglia do not exist as terminally differentiated M1 or M2 states, but are able to switch phenotypes depending on the microenvironment [126, 127]. PPAR- γ plays an important role in modulating macrophage M2 polarization induced by IL-4 or IL-13 [128, 129].

3.2 Neurogenesis and Angiogenesis: The Neurovascular Niche

Neurovascular repair involves, among other mechanisms discussed, the synthesis of new neurons and vessels to replace the dead and damaged ones that result from cerebral injury. Neurogenesis is the process by which neurons are generated from the neural stem cells (NSCs) and progenitor cells, and angiogenesis is the process by which new blood vessels are formed from existing ones. Interestingly, the neural precursors, committed neuroblasts, glia, and endothelial precursors (angioblasts) are found in tight clusters grouped around small capillaries, referred to as the neurovascular niche, indicating a close association between neurogenesis and angiogenesis [130, 131].

Neurogenesis is mostly active in the prenatal period; however, contrary to initial beliefs, NSCs exist beyond embryonic development, into adulthood; thus, a large number of newly generated cells in the adult brain are neurons [132]. Under physiological conditions, adult NSCs differentiate predominantly to inhibitory granule/periglomerular interneurons in the olfactory bulb, and excitatory granule neurons in the dentate gyrus. Also, in adults, neurogenesis constitutively occurs only in the subventricular zone (SVZ) and the subgranular zone (SGZ) of the hippocampal

dentate gyrus and olfactory bulb [133, 134]. Insults to the CNS, such as hemorrhagic strokes, induce increased neurogenesis, which appear to also be generated in non-neurogenic sites, such as the cortex [135]. Immature neurons may be found in the tissue adjacent to the area of the stroke within the first 2–4 weeks after injury [131]. Neurogenesis induces the migration of neuroblasts into regions of degenerating striatum bordering the SVZ, the primary source of which are multipotent glial fibrillary acidic protein (GFAP)-expressing progenitors [132, 136]. The progenitor cells have the capacity to generate neurons, oligodendrocytes, and astrocytes.

After a hemorrhagic stroke, when neurogenesis is initiated, the formed neuroblasts migrate from the SVZ to the peri-infarct cortex, where there is also increased vascular remodeling, likely due to an increase in the metabolic demand following injury, again illustrating the presence of a neurovascular niche, as well as that neurogenesis may be dependent on environmental cues [137]. Additionally, the growth factors and chemokines secreted by vascular endothelial cells may support the survival of the newly formed cells. In line with this, Shen et al. found that coculturing neuroblasts with brain vascular endothelium significantly promoted neurogenesis, and that NSCs were in close proximity to the endothelial cells, where they interdigitated within the folds of the blood vessels and ensheathed the vascular endothelial cells [138]. They also observed that inhibiting angiogenesis resulted in a significant reduction in the number of new neural cells, and upon removal of the endothelial cells, the NSCs generate neurons, as well as astrocytes and oligodendrocytes, as the endothelial factors inhibited differentiation of the stem cells, and instead, promoted self-renewal [138]. Thus, it is suggested that growing NSCs with endothelial cells may be an effective approach to producing more NSCs and neurons for use in replacement therapies.

3.2.1 Astrocytes and Pericytes in Neurogenesis and Angiogenesis

The neural progenitors in the SVZ and SGZ are also in intimate contact with astrocytes, which helps to build a microenvironment that promotes neurogenesis and regulate fate specification [139]. The astrocytes produce signals, such as sonic hedgehog, which dictate the neurogenic potential of different CNS regions, such that neural progenitors are distributed beyond just the neurogenic SVZ and SGZ, into the cortex, cerebellum, and spinal cord. These cells, like those in the SVZ and SGZ, are GFAP-expressing cells [135]. This, therefore, indicates that the neurogenic and non-neurogenic areas of the brain differ not in the properties of their progenitor cells, but in their microenvironment, and this could therefore be therapeutically targeted in order to activate the regenerative potential in non-neurogenic regions of the injured adult brain [135].

Pericytes also play a role in these processes as they interact closely with endothelial cells during the early stages of angiogenesis and with neurons during the maturation of newly formed vessels [140, 141]. The recruitment of pericytes and astrocytes to the newly forming vessels is associated with the formation of tight

junctions, although the tight junctions can be formed without the ensheathment of astrocyte foot processes [101]. Pericytes are involved in the induction of endothelial activation and the augmentation of various proteases, adhesion molecules, and proteoglycans at the initiation of angiogenesis [142, 143], and are conversely involved in the silencing of MMP activities at maturation [144].

3.3 MMPs in Neurovascular Plasticity and Remodeling

While MMPs, which modulate the brain matrix, contribute to neurovascular disruption in the acute phase of hemorrhagic stroke, in the chronic phase, the neurovascular proteases play a role in repair and remodeling by regulating various factors such as VEGF and Kit-ligand [145]. Furthermore, inhibition of MMPs in the late stages results in reduced neuronal plasticity and vascular remodeling, and impaired functional recovery [146]. MMPs belong to a family of zinc-containing endopeptidases, most of which can be produced by endothelial cells. In the early stages after stroke, MMPs were found to be colocalized mainly with endothelial cells, but in the chronic stages, their expression is principally colocalized with neurons and astrocytes [146], which we know are involved in regulating blood flow and responding to increases in neural activity, a vital aspect of repair. MMP-2, MMP-9, and membrane type 1 MMP are known to play a role in the onset of angiogenesis, and matrix degradation is an important aspect of their ability to stimulate angiogenesis. The proteases are able to induce pro- and antiangiogenic effects, such as activating growth factors and cytokines, degrading inhibitors, and generating angiogenesis-inhibiting matrix-derived peptides [144]. For example, MMP-9 stimulates the onset of angiogenesis, but subsequently generates angiogenesis inhibitors, thus retarding the process [147].

It has also been shown that MMP-9 is colocalized with cells from the subventricular zone after stroke, and treatment with an MMP inhibitor significantly decreased migration of the cells to the striatum, indicating that MMPs play a role in neurogenesis after brain injury, and thus are important for the repair process [148]. Being able to modulate these endogenous matrix mechanisms may enhance neurovascular recovery following a hemorrhagic brain injury.

3.4 VSMCs and Neurovascular Coupling

Smooth muscle cells in the VNN, together with neurons, astrocytes, and endothelial cells, maintain homeostasis in the cerebral microcirculation (neurovascular coupling) [149, 150] by constricting or dilating the vessels when they contract or relax, respectively. VSMCs in arteries are able to migrate, proliferate, secrete extracellular matrix proteins, and contract, and they play an essential role in the cerebrovascular response to injury [87]. SMCs communicate with endothelial cells regarding the

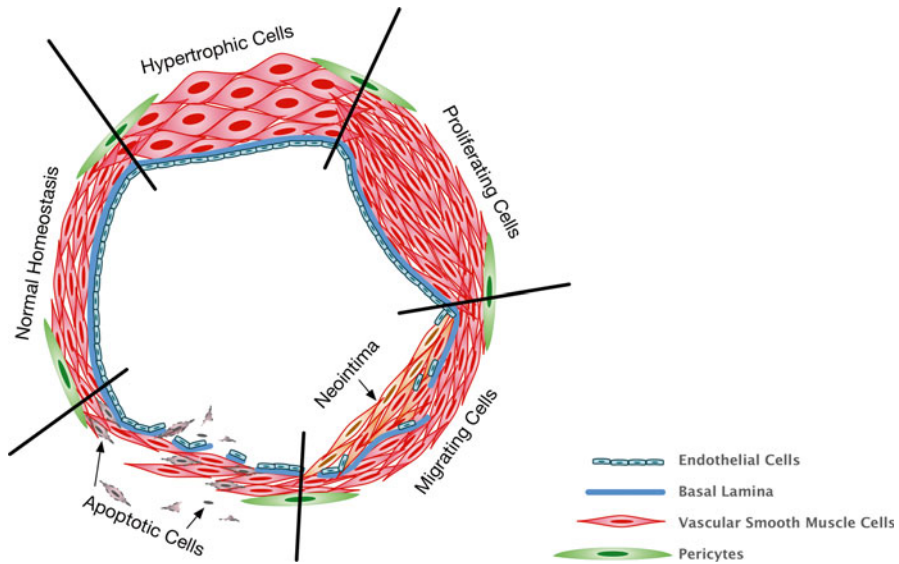


Fig. 5 Phenotypic changes of vascular smooth muscle cells in vascular remodeling after hemorrhagic stroke. Following hemorrhagic stroke, smooth muscle cells undergo phenotypic changes, leading to hypertrophy, proliferation, migration, and apoptosis. Hypertrophic and proliferating cells result in reduced vessel lumen, while apoptotic and migrating cells build a fragile neointima, leading to rupture and leakage. Inflammation, thrombin, and various growth factors may influence these changes, which affect autoregulation and neurovascular coupling

state of blood flow, via heterocellular connexin gap junctions, and also with astrocytes regarding neural activity; thus, they are considered to be the center of the neurovascular network [151]. After aneurysmal SAH, VSMCs undergo phenotypic transformation, proliferate, and secrete cytokines, which may result in loss of myogenic tone and autoregulation, compromising neurovascular coupling and the return of normal CBF (Fig. 5). This phenotypic change can be accelerated by inflammation, thrombin, and various growth factors [152, 153].

Originally, it was thought that neurovascular coupling was solely mediated by changes in the smooth muscle tone around the arterioles. However, it was discovered that pericytes, present at 50 μm intervals along the capillary, can markedly affect capillary diameter, and thus could possibly regulate CBF at the capillary level [150]. Perivascular nerve fibers are also able to promote vasoconstriction and vasodilation, as well as smooth muscle growth and differentiation [73]. SAH reduces both the density and cerebrovascular tone of the nerve fibers, which may lead to functional denervation of cerebral arteries, thus significantly impairing neurovascular coupling [154].

Longden et al. found that after a stress event, parenchymal arterioles did not dilate in response to neuronal stimulation as well as they did without the stress, and dilation of isolated arteries to external potassium was also reduced, suggesting that there was a defect in the functioning of smooth muscle inwardly rectifying potassium (Kir)

channels. As such, they reported that stress reduced Kir channel density, and that blocking the channels significantly inhibited neurovascular coupling in stressed, but not control, brain slices [74]. They conclude that the reduction in these potassium channels makes the arterioles less responsive to potassium released from the astrocytic endfeet, thus affecting neurovascular coupling. Moreover, Koide et al. found that, following SAH, neurovascular coupling was inverted, that is, parenchymal arterioles were dilated in response to elevation of calcium from astrocytic endfeet in control brain slices but were constricted in post-SAH brain slices, and that inhibiting the potassium efflux diminished the vessel response to neuronal activity [46]. These present targets for repair of neurovascular functioning after hemorrhagic stroke.

Another mechanism of neurovascular coupling is via nitric oxide. Neuronal nitric oxide synthase (nNOS) is activated upon calcium entry into neurons, which is induced by glutamate release [155]. This in turn triggers the release of NO, a potent vasodilator. NO can also be released by calcium-independent iNOS and from the endothelium via endothelial NOS (eNOS). In fact, all components of the neurovascular unit can produce NO. In the cortex, inhibition of nNOS reduces the increase in blood flow due to neural activity, indicating that NO plays an important role in neurovascular coupling [156]. Since NO is released upon neuronal activation, and it is known that CBF regulation is tightly bound to neuronal activity, it follows that NO plays a role in regulating CBF. The NO released from activated neurons diffuses through the cell membranes to the astrocytic endfeet (surrounding the arterioles) where it can modulate the release of various vasoactive metabolites; it may also be involved in vasodilation by acting directly on smooth muscle cells [157]. NO increases the release of ATP from astrocytes, leading to the formation of calcium waves via the 2PY purinergic receptors [158]. The calcium waves stimulate the production of arachidonic acid-derived vasodilation mediators such as epoxyeicosatrienoic acids (EETs) or activate large-conductance calcium-dependent potassium (BK) channels to induce arterial vasodilation. In line with this, epoxygenase inhibitors were reported to reduce cortical blood flow in response to vibrissal stimulation, indicating a role for EETs in neurovascular coupling [159]. Moreover, inhibiting potassium efflux via the BK channels in the astrocytic endfeet prevented both vasodilation and vasoconstriction in response to neuronal activity in SAH slices [46].

4 Current Therapies Targeted at Neurovascular Repair

The concepts of the neurovascular unit and VNN integrate neural and vascular cell types, which help to explain the failure of various neuroprotective strategies that target neurons only, without addressing vascular injury. As a result, over the last several years, there have been a lot of studies evaluating various therapies with the goal of improving neurovascular repair. Table 1 looks at some therapies that have been targeted at vascular repair after hemorrhagic stroke.

The BBB is the most effective deterrent to effective delivery of drugs to the CNS due to the tight junctions between the microvascular endothelial cells, as well as the support from surrounding astrocytes, microglia, and extracellular matrix [6]. This

Table 1 Experimental therapies for vascular repair

Therapy	Effects on vascular repair and blood flow	References
tPA	Improved cerebral blood flow and cortical perfusion, thus attenuating early brain injury after SAH	[10]
mGluR stimulation	Increased vasodilation, correlated to the timing of intracellular Ca ²⁺ increase in astrocytes	[75, 76]
PPAR- γ agonist	Attenuated the vasoconstrictive response by upregulating caveolin-1; increased efficiency of microglia-induced phagocytosis; reduced interleukin-1 β , TNF- α , MMP-9, and iNOS expression	[31, 118]
Nitric oxide donors	Dilated spasmic cerebral arteries, and reduced excitotoxicity and leukocyte migration	[90, 169]
Memantine	Attenuated delayed vasospasm by targeting eNOS	[167]
Dopamine D2-receptor agonist	Attenuated vasospasm by mediating increases in vascular and endothelial NOS activity	[168]
L-Arginine	Ameliorated secondary cerebral ischemic injury after SAH by enhancing cerebral blood perfusion, increasing nitric oxide, and decreasing endothelin-1	[171]
Kv7 channel openers	Attenuated basilar artery spasm	[172]
Soluble epoxide hydrolase gene deletion	Increased vasoprotective eicosanoids, and suppressed vascular inflammation and VCAM-1, leading to reduced hydrocephalus and edema	[177]
Sulforaphane	Activated Nrf2/ARE pathway in vascular smooth muscle cells—improved outcomes	[178]
Protein kinase C or Raf inhibitors	Reduced the vasospastic effect, and maintained neurovascular integrity	[179, 180]
PAR-1 inhibitor	Maintained microvascular integrity	[181]
Calcitonin gene-related peptide	Enhanced vasorelaxation by enhancing endothelial nitric oxide and cAMP-mediated endothelium-independent signaling	[182]
TNF- α proteinase inhibitor	Reversed the augmented myogenic tone in isolated vessels	[183]

Various therapies have been targeted at different components of neurovascular interactions. This table summarizes some current experimental treatments that are directed at vascular repair after a hemorrhagic stroke [177–183]

mGluR metabotropic glutamate receptor, *tPA* tissue plasminogen activator, *SAH* subarachnoid hemorrhage, *VCAM-1* vascular cell adhesion molecule 1, *Nrf2/ARE* nuclear factor erythroid 2-related factor 2/antioxidant responsive element, *PAR-1* protease-activated receptor, *Kv7* voltage-gated potassium channel, type 7, *NOS* nitric oxide synthase, *cAMP* cyclic adenosine monophosphate, *TNF* tumor necrosis factor, *PPAR* peroxisome proliferator-activated receptor, *MMP* matrix metalloproteinase

stringent barrier is necessary for normal physiological functioning: facilitating nutrient transport, regulating ion balance, and preventing toxic substances in the systemic circulation from entering the brain; restoring its function after stroke is the primary goal of many current therapies. The Src-family kinase inhibitor protein phosphatase-1 was found to decrease the phosphorylation of VEGF and mitogen-activated protein kinases (MAPK), thereby reducing BBB permeability, brain edema, and

mortality after hemorrhagic stroke [160]. Further, thrombomodulin preserved endothelial junction proteins and reduced cell apoptosis and inflammation in endothelial cells via blocking p38 and MAPK [161]. Norrin, a frizzled-4 receptor agonist that promotes β -catenin nuclear translocation, is also targeted at BBB preservation. Treatment with norrin attenuated BBB permeability by enhancing the expression of occludin, vascular endothelial (VE)-cadherin, and ZO-1 after SAH. [162].

Since MMPs and the by-products of reactive oxygen and nitrogen species are known to be involved in BBB breakdown during cerebral injury, several studies have also investigated interventions in these pathways in order to achieve neurovascular repair. A peroxynitrite inhibitor ameliorated neurovascular dysfunction and improved neurological function, in part by reducing MMP-9 after hemoglobin injection into the caudate nucleus [163]. Also, inhibition of c-Jun N-terminal kinases-induced MMP-9 activation was shown to protect the neurovascular unit and preserve the tight junction protein, ZO-1 [164]. Imatinib, a PDGF receptor alpha inhibitor, also preserved BBB integrity following experimental SAH by reducing MMP-9 [165].

Other therapies have been targeted at reducing vascular injury and the concomitant reduction in blood flow secondary to hemorrhagic stroke. For example, heparin, which blocks the activity of free radicals, including ROS, antagonizes endothelin-mediated vasoconstriction, smooth muscle depolarization, and inflammatory and fibrogenic responses via complex-formation with oxyhemoglobin [166]. Targeting eNOS, memantine attenuated delayed vasospasm after SAH [167]. Further, a dopamine D2-receptor agonist was found to mediate increases in vascular and endothelial NOS activity in vitro, and this was associated with attenuated vasospasm [168]. Notably, NO donors administered after experimental SAH dilated spasmic cerebral arteries, and reduced excitotoxicity and leukocyte migration [98, 169]. This proved consistent with the observation that acute ischemic injury can be reversed by the NO donor *S*-nitrosoglutathione [170]. Moreover, L-Arginine, which increases NO, and decreases ET-1 levels in blood, enhanced cerebral blood perfusion, and ameliorated secondary cerebral ischemic injury after experimental SAH [171]. Increases in intracellular calcium levels also induce vasoconstriction by activating potassium channels on astrocytes. As such, the voltage-gated potassium channel (type 7) openers (retigabine and celecoxib) were reported to significantly attenuate basilar artery spasm in rats with experimentally induced SAH [172].

Still other therapies have been targeted at oxidative stress and inflammation and their effects on neuronal survival and other outcomes after hemorrhagic stroke. Nuclear factor erythroid 2-related factor 2 (Nrf2) activation can induce antioxidation, reduce peroxide formation, and upregulate phagocytosis by inducing CD36 expression, thus promoting hematoma clearance after ICH [173]. Activating Nrf2 reduced the permeability of the BBB after ICH, and reduced microglial activation in the perihematoma, resulting in improved functional outcomes [174]. Nrf2 and other such therapies therefore protect the neurovascular components by reducing the toxicity induced by blood components and reducing vascular disruption. In fact, knock-out of Nrf2 led to impaired hematoma clearance, suggesting that Nrf2 in microglia plays an important role in augmenting this process [175]. Notably, resolution of the

hematoma also leads to a reduction in inflammation, as blood components are able to promote the initiation of inflammation following injury. This in itself can limit injury to the neurovascular network by decreasing the disruption of the BBB following hemorrhagic stroke [24].

Other anti-inflammatory therapies investigated include a P-selectin antibody and the nicotinamide adenine dinucleotide phosphate (NADPH) oxidase inhibitor, apocynin, which decreased rolling and adherent platelets and leukocytes in pial venules after SAH [58]. This is important as NADPH is one of the major sources of free radicals after subarachnoid bleeding. Vinblastine and anti-polymorphonuclear serum were also able to reduce the neutrophil adherence to parenchymal vessels [106]. Additionally, hyperbaric oxygen ameliorated neurovascular injury via several inhibitory signaling pathways with antioxidant, anti-inflammatory, and anti-apoptotic effects [176].

These are just a few of the many therapies being evaluated, and while advances have been made in the field, and these experimental drugs seem promising in experimental models, there is still the challenge of successful clinical translation. There may be a number of reasons for this, and of course, more studies are needed in this area. However, a closer look at combination therapies may be worthwhile, targeting neuronal and vascular injury simultaneously.

5 Conclusions

All the cells of the VNN play significant roles in brain injury and repair. Hemorrhagic stroke induces significant damage to neurons and the vasculature, impairing the functionality of the brain. Heme and heme breakdown products lead to an increase in neurotoxicity and vascular injury. Cerebral vessels and smooth muscle cells undergo phenotypic changes that alter the supply and demand system of the brain, whereby CBF is linked to neural activity. This can therefore result in significant damage to the VNN. Additionally, endothelial dysfunction leads to disruptions in the blood–brain barrier, vasoconstriction, and neuronal death. However, astrocytes help to protect neurons from excitotoxicity by modulating glutamate release and uptake, as well as in regulating CBF through calcium release; microglia and macrophages aid in repair by engaging in the clearance of toxic blood products; and pericytes and endothelial cells play vital roles in maintaining BBB function and in regulating inflammation. Additionally, there is increased neurogenesis and angiogenesis following a hemorrhagic stroke insult, and targeting these processes, within the neurovascular niche where they are codependent, could have significant impacts on neurovascular repair. Moreover, there is an increase in MMPs that initially enhances damage to the brain, but in chronic stroke, can be modulated to improve outcomes by interactions with the neurovascular matrix. Coordinately targeting the VNN, neuronal and vascular injury, may be the best approach for improving hemorrhagic stroke outcomes.

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The Role of Nonneuronal Nrf2 Pathway in Ischemic Stroke: Damage Control and Potential Tissue Repair

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Ischemic stroke is one of the leading causes of disability and mortality in the world. Multiple pathological processes participate in the progression of stroke, including excitotoxicity, oxidative stress, inflammation, mitochondrial dysfunction, etc. Excessive oxidative stress is one of the most important pathogenic mechanisms in stroke. Therefore, therapeutic strategies against oxidative stress may be feasible for the treatment of stroke. Given that nuclear factor erythroid 2-related factor 2 (Nrf2) pathway is the predominant antioxidant system, we will focus on the promising protective role of Nrf2 pathway against ischemic stroke.

1 Introduction of Nrf2 Pathway

Nrf2 is a member of the basic leucine zipper (bZIP) transcription factor family featuring a Cap 'n' collar structure [1, 2], and it is essential for the transcriptional induction of phase II drug-metabolizing and antioxidant enzymes, such as glutathione s-transferase (GST), heme oxygenase 1 (HO-1), and NADPH quinone oxidoreductase 1 (NQO-1) [3]. Nrf2 regulates gene transcription of these enzymes by binding the antioxidant response elements (AREs) on their promoters.

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1.1 Regulation of Nrf2/ARE Pathway

Nrf2 has six functional domains known as Nrf2-ECH homologies (Neh), designated as Neh1-6. Each Neh domain has its own function [4]. Among them, the Neh1 domain is the CNC-bZIP domain, enabling Nrf2 to form heterodimer with the ZIP domain of small musculoaponeurotic fibrosarcoma (Maf) proteins [5]; and the Neh2 domain mediates binding with the cytosolic repressor of Nrf2. Keap1 encompasses three functional domains: a bric-a-brac (BTB) domain, an intervening region (IVR), and a Kelch domain (also named DGR domain). Under non-stressed conditions, Nrf2 forms a complex with Kelch-like ECH-associated protein 1 (Keap1), which promotes rapid proteasomal degradation of Nrf2 by proteasome, giving Nrf2 a half-life time of about 20 min [3]. As a result, Keap1 functions to inhibit Nrf2.

It has been reported that alterations in the structure of Keap1 leads to dissociation of the Nrf2–Keap1 complex, and this is essential for Nrf2/ARE activation [5]. Three cysteine residues (Cys151, Cys273, and Cys288) in Keap1 are essential for Keap1 repression of Nrf2 activity under unstressed conditions [6]. The oxidation of these cysteine residues causes conformational change of Keap1 and leads to dissociation of the Nrf2-Keap1 complex. Some oxidative products can free Nrf2 from the complex and increase the expression of phase II genes [7].

Transcription factors need to translocate to the nucleus in order to transactivate. As for Nrf2, this process is quite rapid: Nrf2 can accumulate in the nucleus within 15 min after tert-Butylhydroquinone (t-BHQ) treatment [8]. The key mediators that regulate nuclear import and export of transcription factors are the nuclear localization signals (NLS) and nuclear export sequences (NES). A number of such nuclear shuttling signals have been identified on Nrf2, including three NLS motifs and two NES motifs. The direction of Nrf2 movement is determined by a homeostatic balance between import and export driving forces. Under normal condition, Nrf2 stays in the cytosol because the import force is less than the export force while NES1 is functional. Under oxidative stress, Cys183 of NES1 is adducted and NES1 loses function. Then the import force is more than the export force, resulting in Nrf2 nuclear translocation [9]. Additionally, Nrf2 nuclear translocation can also be triggered when phosphorylated by kinases. The phosphorylation sites of protein kinase C (PKC), glycogen synthase kinase-3 and Fyn have been identified. The phosphorylation site of PKC is Ser40 in Neh2 domain [10], but its role in Nrf2 nuclear shuttling is controversial. Fyn phosphorylates Tyr568 in the Neh3 domain, controlling nuclear export of Nrf2 [11, 12].

After translocating into the nucleus, Nrf2 can form heterodimer with Maf proteins [13], which further enhances the specificity of Nrf2 to bind to a *cis*-acting enhancer ARE [14] located at the promoter of phase II genes [1, 15]. Subsequently, the transcription of phase II gene is initiated.

Recent studies have also proposed a Keap1-independent ubiquitination model of Nrf2 degradation [16]; in which, GSK3 β is a key player because it can phosphorylate Nrf2 at Ser342 and Ser347 located at Neh6. Phosphorylated Neh6 can bind with an ubiquitin ligase adaptor beta-transducin repeat-containing protein (β -TrCP). β -TrCP is a scaffolding protein that directly links Nrf2 to the Cullin1/Rbx1 ubiqui-

ination complex. Therefore, GSK-3 β -mediated phosphorylation of Neh6 causes the ubiquitination and degradation of Nrf2 via β -TrCP. This model is supported by the stabilization of Nrf2 by GSK-3 β inhibitors in Keap1^{-/-} mouse embryo fibroblasts (MEFs) [12]. Additionally, cancer-chemo preventive agent nordihydroguaiaretic acid can activate Nrf2 and increase HO-1 protein levels through inhibiting GSK-3 β phosphorylation in Keap1^{-/-} MEFs [17].

Additionally, there are some other regulatory mechanisms to prevent excessive activation of Nrf2. For example, AREs are located in the promoter region of Cul3, Rbx1, and Keap1 genes; while activation of Nrf2/ARE pathway boots the expression of phase II genes, it also upregulates these inhibitory proteins at the same time. This negative feedback loop is known as an autoregulatory arm of the Nrf2/ARE pathway [18]. In addition, the Keap1-Cul3-Rbx1 complex can enter the nucleus, mediated by prothymosin α (ProT α), a Keap1-binding protein with a NLS. As a result, 10–15 % of Keap1-Cul3-Rbx1 complex is localized in the nucleus. Once entering the nucleus, the Keap1-Cul3-Rbx1 complex drops ProT α and binds Nrf2, leading to the ubiquitination and degradation of nuclear Nrf2 [19].

1.2 Inducers and Effectors of Nrf2 Pathway

It is important to identify inducers and effectors of Nrf2 pathway, because they could be promising therapeutic targets. As discussed above, Keap1 is the major inhibitor of Nrf2 activation; changes in its structure and/or its dissociation from Nrf2 results in the release and activation of Nrf2, which is recognized as the target mechanisms of nearly all Nrf2 inducers. There are several ways to classify Nrf2 inducers. Based on their origin, inducers are divided into two classes, the exogenous and the endogenous; based on their chemical structure, inducers are divided into ten groups [20]; based on the Keap1 domains that the inducers react with, they can be divided into four categories [21].

Currently, over 200 Nrf2/ARE-driven proteins are described for detoxification and antioxidant defense. Among which, phase 2 enzymes initially caught the interest of scientists for their preventative action against carcinogens [22]. Later, their function has been expanded to neuroprotection [23, 24]. Table 1 lists the major cytoprotective enzymes that participate in protection against neuronal injury in common neurological diseases.

2 Nrf2 Provides Neuroprotection Against Ischemic Stroke Via Self-Defense and Cell-Cell Interaction

In ischemic stroke, several pathological mechanisms known as the ischemic cascade are triggered, and consequently cause rapid and irreversible neuronal death within the ischemic core due to severe and rapid loss of energy. In the surrounding hypoperfused brain tissues, known as the penumbral areas, the neurons may be

Table 1 Effectors of Nrf2 pathway

	Enzymes	Action mechanisms
Thioredoxin enzyme system	Thioredoxins	Reduction oxidizing proteins via the exchange between cysteine thiol and protein disulfides
	Thioredoxin reductase	Reduction of oxidized thioredoxins by consuming NADPH
	Peroxiredoxins	Catalyzing the reduction of peroxides by utilizing thioredoxins
	Sulfiredoxins	Reduction of the sulfinic acid phosphoric ester on oxidize peroxiredoxins
Glutathione system	GSH	Scavenging multiple oxidative species; Serving as a reservoir for cysteine
Transferase	GSH-S-transferase	Transference of GSH
	N-Acetyltransferase	Transfer of an acetyl group to an arylamine
	Sulfotransferase	Transferring a sulfo group to an alcohol or amine
Detoxifying enzymes	HO-1	Breakdown of heme; Generation of antioxidants
	NQO1	Catalyzing the two-electron reduction of quinone to the redox stable hydroquinone

potentially salvageable due to relatively moderate or mild energy loss. Therefore, salvaging the neurons in penumbra is of great translational significance. To do so, multiple protection and supports may be needed, because neuronal death in penumbra is not a stand-alone event, instead, it is the consequence of neuronal injury itself plus dysfunction of all kinds of cells within the neurovascular unit (NVU), including endothelial cells, astrocytes, and others [25–27].

2.1 Oxidative Stress and Nrf2 Activation in Neurons After Brain Ischemia

Following ischemia and reperfusion, the cascade of events that leads to neuronal deaths can be summarized as below [28–30]. Energy failure is the first event of the cascade caused by decreased ATP production, the latter then disrupts Na⁺/K⁺-ATPase, Ca²⁺/H⁺-ATPase, and Na⁺/Ca²⁺-transporter, leading to depolarization of cellular membrane. After depolarization, excitotoxicity amino acids, especially glutamate, are released into the synaptic cleft, which consequently activates *N*-methyl-D-aspartic acid (NMDA), α -amino-3-hydroxy-5-methylisoxazole-4-propionic acid (AMPA), and metabotropic glutamate receptors increasing intracellular Ca²⁺ load. A variety of Ca²⁺ dependent enzymes are then activated and hence induce protein phosphorylation, proteolysis, mitochondrial damage, and the generation of free radical, i.e., reactive oxygen species (ROS) and reactive nitrogen species (RNS). ROS and RNS eventually lead to oxidative stress, which cause lipid peroxidation, membrane injury, DNA damage and moreover, allowing the infiltration of

leukocytes and activating inflammation [31]. Multiple neuroinflammatory cascades are also activated, inducing secondary brain injury that leads to cell death [32].

During the procedure of neuronal death in the penumbra, oxidative stress triggered by ROS and RNS generation is considered the “key” event. The predominant ROS and RNS produced are superoxide anion ($O_2^{\cdot-}$), hydrogen peroxide (H_2O_2), hydroxyl radical ($\cdot OH$), nitric oxide ($\cdot NO$), peroxynitrite anion ($ONOO^-$), and nitrogen dioxide ($\cdot NO_2$) [28]. Ischemic insults produce an excessive amount of free radicals especially in the reperfused regions, and the imbalance in the formation and clearance of ROS and RNS leads to oxidative stress, and subsequently alters the cell dynamics [33] and impacts major cellular components [26]. Complex signaling network involving numerous survival/death pathways have been reported to be redox mediated [34]. ROS and perhaps RNS, may act directly as executioners of cell death [31]. ROS leads to increase of mitochondrial permeability transition pore which subsequently causes mitochondrial swelling and neuronal death [35]. ROS also initiates apoptosis through activating p53 and p38 MAPK [36]. Another damage effect acted by ROS to ischemic brain is that ROS can directly increase the permeability of the blood–brain barrier (BBB) [37]. Focal ischemia induces a potent inflammatory response within a few hours after onset of ischemia, and BBB disruption allows various blood immune cells infiltrate into ischemic area [38–40]. These migrated immune cells, along with injured brain cells, produce inflammatory mediators, enhancing neuronal death [41].

The brain is sensitive to oxidative stress. Constituting only 2% of the total body weight, the brain uses 20% of the total oxygen consumed by the whole body, making the brain the major oxygen user and probably a major source of ROS [28]. In addition, a high level of unsaturated lipid content, multiple chemical reactions in dopamine oxidation and high concentrations of iron also contribute ROS generation in the brain [42, 43]. To defense against oxidative stress, both direct and indirect antioxidants are needed [44–46]. Direct antioxidants denote a group of low molecular weight compounds that can directly undergo redox reactions and scavenge reactive oxidation products, ROS and RNS. This group includes glutathione (GSH), ascorbate, tocopherols, lipoid acid, carotenes, and ubiquinones. Direct antioxidants are consumed in the process, therefore need replenishment or regeneration. Indirect antioxidants, however, may not present the redox activity themselves. They exert their antioxidant effects through the upregulation phase 2 and antioxidant enzymes (Table 1), or so-called “phase 2 response” [5, 47, 48], which display a wide variety of antioxidant activities.

Several studies suggest that ROS is generated in the brain after ischemia. Peters et al. reported that rats subjected to middle cerebral artery occlusion (MCAO) first exhibited a short period (10–30 min) of decreased ROS level after the occlusion, followed by a significant increase (to $162 \pm 51\%$; baseline = 100%) from 100-min ischemia to permanent MCAO onward [49]. Moreover, reperfusion after 1 h of MCAO led to a burst-like pattern of ROS production (about fivefold), and lasted at least to 3 h post MCAO. Liu et al. studied the spatial distribution of ROS in ischemic rat brain and found that ROS was significantly increased in the infarct core during

both ischemia and reperfusion, but in penumbra ROS was only increased during reperfusion [50, 51].

Nrf2 pathway is activated in post-ischemic stroke. Tanaka et al. [52] explored the expression of Nrf2 and Keap1 in mice subjected to MCAO using immunohistochemistry and immunofluorescence. Their results showed that, Keap1 was significantly decreased from 24 to 72 h post MCAO, paralleled with increased expression of Nrf2. In consistency with previous results, rats subjected to MCAO also exhibited significantly upregulated Nrf2 expression 24 h later, predominantly located in the penumbra compared to the core [53]. Li et al. [54] detected Nrf2 levels in rat brain by immunohistochemistry and western blotting analyses after MCAO and reperfusion, and found that, Nrf2 and hemeoxygenase-1 (HO-1) were upregulated in ischemic cortex, beginning at 6 h, peaking at 48 h and declining at 72 h after reperfusion. With a novel immunohistochemical technique, Srivastava et al. [55] examined the temporal and spatial distribution of Nrf2 in rat brain following ischemia-reperfusion injury in a detailed way. They used the nuclear/cytoplasmic Nrf2 ratio to indicate the level of Nrf2 activation of Nrf2 pathway, and demonstrated that nuclear to cytoplasmic Nrf2 ratios in stroke-affected regions were increased after 24 h and then declined after 72 h reperfusion. Moreover, Nrf2 expression was significantly higher in the penumbra than the infarct core. Oligemia is another model used to study the stroke penumbra, in which lipid peroxidation induces an increased oxidative stress and an augmented \bullet OH production during the reperfusion phase. Liverman et al. [56] adopted this model in mice by lowering the mean arterial pressure to 30–40 mmHg, and reported a significant elevation of Nrf2 level in neurons in the Purkinje cells of the cerebellar cortex and pyramidal neurons of the cingulate cortex.

In short, piles of evidence verify the indispensable role of Nrf2 pathway as a neuronal endogenous protective mechanism through the induction of its downstream antioxidants, including the direct (small molecular antioxidants) and indirect (predominant phase 2 enzyme) ones. These antioxidants then provide a cytoprotective role against ischemia-induced neuronal death.

2.2 Nrf2 Activations in Nonneuronal Cells: The Contribution of Cell–Cell Interaction

As indicated previously, the neuronal death in ischemic stroke is a not a separate procedure; instead, it is caused by additional dysfunction of other types of cells within and beyond the NVU due to complicated cell–cell interactions. Diverse types of cells in the brain work as a unit to maintain the central nervous system (CNS) homeostasis. This can partially explain the reason why neuroprotection trials have mostly failed, as any disruption within the complicated system might lead to NVU dysfunction after brain ischemia [57]. In this regard, therapeutic strategies should not only aim to neurons, but also to salvage other cells and hence restore function of the NVU [26, 29, 57].

Dual-direction responses have been reported between neurons and endothelial cells in the brain. During brain development, microvessel and neuron are already arranged to grow together along the extracellular matrix paths [58–60]. In adult brain, the regulation of regional cerebral blood flow depends on the activity of neurons [61]. Zonta et al. reported that, in a rat hyperemia model, neuronal afferent stimulation mediates the dilation of cerebral arterioles, which is dependent on the glutamate-mediated $[Ca^{2+}]_i$ oscillations in astrocytes [25, 62]. On the other hand, some novel discoveries focusing on microvessel structure, endothelial cells, and astrocyte endothelial adhesion indicated an opposite regulatory direct, from microvessel to the neurons they supply [63, 64]. An interesting research by Mabuchi et al. showed that an ordered and sequential microvessel–neuron relationship existed in contralateral basal ganglia; in ischemic area, neurons more distant from their nearest microvessel are more sensitive to ischemia, indicating that neuronal survival is dependent on the microvascular function [65]. Besides, microglia and macrophages play specific roles in the ischemic brain, functioning as a “double-edged sword” by either cleaning up or inducing local inflammation [66, 67]. In the following context, we will discuss the contribution of non-neuronal Nrf2 pathways to neuronal survival.

2.2.1 Nrf2 Pathway in Astrocyte

Comparing to neurons, astrocytes produce large amount of antioxidants [45, 68], and several lines of evidence have demonstrated that astrocytic Nrf2 contributes to neuronal survival following brain ischemia. Kraft and colleagues reported that the activation of astrocytic Nrf2 protected neurons from hydrogen peroxide and glutamate in cell cultures [69]. This protection disappeared in cultures from Nrf2 knockout animals; and the protection could be restored by infecting cells with a replication-deficient adenovirus that carried Nrf2, indicating an important role of astrocytic Nrf2 in neuronal protection. Shih et al. also studied the astrocyte–neuron interaction using a co-culture model of neurons, naïve astrocytes, and infected astrocytes (Nrf2 overexpression cells) [70]. They reported that astrocytes have higher basal Nrf2 expression and ARE activity than neurons, with an increased expression of GSH. When stimulated, the co-culture with oxidative glutamate toxicity, they found that Nrf2 activation led to an increase in both media and intracellular GSH in astrocytes; experiments on selective inhibition of glial GSH synthesis indicated that an Nrf2-dependent increase in glial GSH synthesis was both necessary and sufficient for the protection of neurons [70]. It is not fully understood how astrocytic GSH protects neurons [71]. It has been reported that GSH can scavenge free radicals in the extracellular area [72]. In addition, GSH can be hydrolyzed on the external surface of astrocytes and thus provide high yield of cysteine and glycine, which can be taken up by neurons and used for intracellular GSH synthesis [73].

Astrocytic Nrf2 also contributes to neuronal protection in a mitochondria-dependent manner [74, 75]. 3-nitropropionic acid (3NP) and malonate are two mitochondrial

complex II inhibitors, and they can kill neurons probably by inducing ROS generation in mitochondria. Calkins et al. reported that Nrf2-deficient astrocytes and mice were more vulnerable to 3NP or malonate; and they also found that astrocytes showed increased ARE-regulated transcription [76]. If Nrf2-overexpressing astrocytes were transplanted into the brain before 3NP or malonate treatments, a dramatic protection was noticed against complex II inhibition [76]. To extend these findings, they developed a line of transgenic mice with astrocyte-specific overexpression of Nrf2, and found the transgenic mice was resistant to malonate insults, which was associated with elevation of Nrf2-driven genes such as NQO1, GCLM, and HO-1. Furthermore, they showed that malonate toxicity could also be reduced by striatal transplantation of neuroprogenitor cells overexpressing Nrf2, which differentiated into astrocytes after grafting [77].

Astrocytic Nrf2 pathway was also reported to play a positive role in brain protection following ischemic preconditioning [78]. Bell et al. reported that Nrf2 could be activated by mild oxidative stress in both rodent and human astrocytes, and that transient ischemic conditions *in vitro* and *in vivo* cause an increase in the expression of Nrf2 target genes, especially the GSH system [78]. Astrocytic Nrf2 also contributes to chemical preconditioning induced by resveratrol [79]. Nrf2 pathway was found activated in astrocytes by resveratrol, and loss of Nrf2 reduced resveratrol-mediated neuroprotection in mice. After resveratrol treatment, both wild-type and Nrf2^{-/-} cortical mitochondria produced ROS, and Nrf2^{-/-} cells showed decreased mitochondrial antioxidant expression and failed to elevate cellular antioxidants after preconditioning. These data suggest that astrocytic Nrf2 pathway is critical in preconditioning-induced neuroprotection [79].

It has also been reported that astrocytes can protect endothelial cells in a GSH-dependent manner. Schroeter et al. reported that astrocytic cultures showed higher antioxidative activity than endothelial cultures, indicated by increased levels of SOD, catalase, and glutathione peroxidase [80]. When cultured these types of cells together, they found an increased antioxidative capacity in endothelial cells. In another report, Laird et al. reported that hemin could induce apoptosis in mouse and human endothelial cells; and that astrocyte-conditioned media could rescue the apoptotic endothelial death, in which GSH played a key role as the protection was aborted by prior treatment of astrocyte with DL-buthionine (S, R)-sulfoximine, a GSH-depleting agent [81].

2.2.2 Nrf2 Pathway in Endothelial Cell

In term of structure, endothelial cells are the center of NVU, as they play important roles in nourishing surrounding cells and keeping the integrity of BBB and NVU. Under pathological conditions, however, endothelial cells may generate ROS that may induce neuronal cell death [82]. When their ROS reacts with nitric oxide (NO), peroxynitrite, a much more powerful oxidant than ROS, will form [83, 84]. In addition to damaged surrounding structures, those ROS and RNS will cause the

disruption of BBB integrity following ischemia, in the forms of increased passive diffusion or massive cellular infiltration [85, 86].

As a way of self-defense, endothelial cells equip themselves with Nrf2 system, and several studies suggest that Nrf2 and its target enzymes protect endothelial cells from oxidative insults and sustain BBB integrity following brain ischemia. Alfieri et al. [87] pretreated rats with sulforaphane and then subject them to MCAO followed by reperfusion. They found sulforaphane significantly increased HO-1 expression in brain microvessel and that BBB disruption was attenuated after stroke. Bénardais et al. detected the protective effects of dimethylfumarate (DMF), an Nrf2 activator, on BBB integrity and found that both DMF and its primary metabolite monomethylfumarate (MMF) activated Nrf2 pathway and upregulated NQO1 in brain endothelial cells [88]. Moreover, DMF can partially attenuate TNF- α -induced downregulation of tight junction (TJ) protein [88]. DMF may also protect endothelial cells by stabilizing the BBB via preventing TJ disruption and suppressing the activity of matrix metalloproteinase (MMP) [89]. A similar protective effect was reported by Wu et al. [90] with an oral administration of procyanidin B2, a dietary Nrf2 inducer, 3 h after MCAO. They found that procyanidin B2 significantly reduced the infarct volume, brain edema, and neurological deficits; moreover, they also noticed increased levels of TJ proteins in the brain microvessel and sustained BBB integrity indicated by Evans blue leakage. Nrf2 activation may underlie the observed protections as these protections are associated with increased expression of HO-1, NQO1, and GST α [90].

In addition to ischemic stroke, BBB protection by Nrf2 pathway is also reported in other CNS disorders, such as subarachnoid hemorrhage (SAH) and traumatic injuries to brain and spinal cord.

SAH is another form of stroke with noticeable vasospasm and subsequent ischemia for several days, and BBB disruption is common after SAH that can result in further brain injury. It has been reported that Nrf2-ARE pathway was activated after SAH, and administration of Nrf2 inducers, such as sulforaphane [91] or t-BHQ [92, 93] can significantly attenuate BBB disruption induced by SAH. In addition to these Nrf2 inducers, some other neuroprotective agents can ameliorate BBB disruption through activating Nrf2 and upregulating HO-1, NAD(P)H, NQO-1; such agents include astaxanthin [94], propofol [95], melatonin [96], and recombinant human erythropoietin [97]. On the other hand, Nrf2 knockout exacerbated brain injury after SAH with increased brain edema, BBB deficits, and decreased GSH [98].

In traumatic brain injury, sulforaphane was reported to preserve BBB function in association with reduced loss of TJ protein and endothelial markers [99]. This protection disappeared in Nrf2 knockout mice or when the animals were pretreated with decoy oligonucleotides containing the binding site of Nrf2 [99]. Similar phenomenon was reported in animals subjected to spinal cord injury. For instance, t-BHQ could alleviate spinal cord edema and suppress the expression of tumor necrosis factor alpha (TNF- α), interleukin-6 (IL-6), and interleukin-1 beta (IL-1 β) [100]. Nrf2^{-/-} mice showed more severe edema after spinal cord injury, which is associated with increased levels of MMP-9 and TNF- α probably caused by an

increase in BBB disruption [101]. These studies suggest that Nrf2 pathway contributes to endothelial protection in broad-spectrum conditions.

2.2.3 Nrf2 Pathway in Microglia/Macrophage

For a long time, the brain has been regarded as an “immune privileged” organ for its lack of classical immune response [102, 103]. However, it has become increasingly clear that the brain is actually an immune organ, especially with the discovery of lymphatic vascular system in the brain [104–106], suggesting that complicated interactions may exist between CNS and the immune system.

Microglia are CNS residential immune cells which share a lot of features with monocytes and macrophages and provide innate immune responses. They need to consume energy for a broad range of activities, and are thus highly sensitive to energy deficits and local changes in blood perfusion [67]. It is not surprising that microglia are among the first cells to respond to CNS injuries. For example, they are mobilized within 1 h after ischemic stroke, start to proliferate within 72 h, and continue to accumulate for >1 month. The function of microglia in stroke is recognized as a “double-edged sword” [66, 104]; they release cytotoxic cytokines that can kill the other cells including neurons at acute stage after ischemia, but they clear cellular debris via phagocytosis and autophagy and restrict the inflammation at late stage. Therefore, it may be critical to fine-tune the function of microglia and macrophages in order to limit brain damage and enhance restoration functionally and partially structurally following stroke. We will briefly discuss the perspectives of activating microglial Nrf2 pathway in the modulation of inflammatory response, autophagy, and phagocytosis after ischemic stroke.

Nrf2 Suppresses Microglial Inflammatory Response

It is widely accepted that inflammatory processes play a pivotal role in neuronal death, especially the delayed one that may last for days and even weeks after ischemic stroke [107]. Microglia and macrophages mediate inflammatory response by releasing pro-inflammatory cytokines such as IL-1 β and TNF- α and by releasing other factors such as cyclooxygenase-2 (COX-2), NO and MMPs [107, 108].

Foresti et al. screened 56 small molecules that are potent Nrf2 activators, including t-BHQ, and carnosol, and showed that activation of Nrf2 could significantly inhibit LPS-induced inflammation in mice microglia-like cell line BV2 cells, which could be abrogated by transfection of Nrf2 shRNA and HO-1 shRNA [109]. Dilshara and coworkers reported that α -viniferin could reduce LPS-induced NO and COX-2 production in BV-2 cells, which could also be abrogated by Nrf2-siRNA transfection [110]. In addition, several other Nrf2 activators, including adenosine [111], β -lapachone [112], KCHO-1 [113] and tissue inhibitor of metalloproteinase-2 [114], could also reduce inflammatory responses in association with Nrf2 activation and HO-1 upregulation. It is not clear how Nrf2 activation can inhibit inflammatory

response. While Foresti [109] and Dilshara's findings [110] may suggest a direct link between Nrf2 pathway and the inhibition of inflammation, findings of other studies may need to be further investigated.

Nrf2 Pathway in Microglial Autophagy

Autophagy is a lysosome-mediated degradation process for non-essential or damaged cellular constituents, which involves more than 30 autophagy-related proteins (ATGs) and 50 lysosomal hydrolases [115]. Physiologically, autophagy also contributes to preserve homeostasis through the removal of unwanted or damaged mitochondria as well as other organelles. Three classes of autophagy have been described based on how protein substrates for degradation reach the lumen of lysosome [115]. (i) Macroautophagy, which is generally referred to as autophagy, is mediated by the formation of double-membrane vesicles known as autophagosomes. When autophagosomes mature, they fuse with and deliver their contents for degradation. (ii) Microautophagy is a process in which lysosomes directly engulf the cytoplasmic material to degrade. (iii) Chaperone-mediated autophagy is more complex and specific, which involves the recognition by the heat shock cognate protein 70 (HSC70)-containing complex. In brief, the target protein must contain the recognition site for HSC70 complex, which will allow it to bind to this chaperone, leading to the formation of the substrate/chaperone complex. The complex will then move to the lysosomal membrane and enter the cell, whereby get degraded by the lysosome. Among the three classes mentioned above, macroautophagy is the most common one and will be our focus in this section.

The overall mechanism of autophagic regulation is almost clear now. Generation of the autophagosomal structure requires the beclin-1-class III PI3K complex and the generation and insertion of light chain 3 (LC3, also known as Atg8)-II complex into the autophagosomal membrane. The Atg genes control autophagosome formation through Atg12-Atg5 and LC3-II complex [116]. Atg12 is conjugated to Atg5 in a ubiquitin-like reaction, which requires Atg7 and Atg10 as enzymes of the reaction. The Atg12-Atg5 conjugate then interacts with Atg16, and together form a large complex with beclin-1-class III PI3K. The generation of LC3-II complex requires Atg3, Atg4, and Atg7 as enzymes of a ubiquitin-like reaction. Sequestosome 1 (SQSTM1, also known as p62), an "adaptor" molecule in selective autophagy, helps the attachment of LC3-II complex to the autophagosome membrane.

There are at least two mechanisms via which Nrf2 pathway regulates autophagy. First, the Atg proteins mentioned above are sensitive to redox signaling, thus redox status regulates autophagy [117]. It is not surprised that Nrf2 can regulate autophagy through regulating the redox status. In addition, Nrf2 has also been reported to upregulate the expression of Sestrin 2 which acts to scavenge ROS and promote autophagy, through enhancement of Sestrin 2 promoter activity [118]. The other mechanisms involve the interaction of Nrf2 with p62. As a matter of fact, positive feedback loop exists between p62 and Nrf2 pathway. It is reported that Nrf2 pathway directly enhances macrophage autophagy through promoting p62 expression

[119, 120], because p62 is a target gene of Nrf2 [121, 122]. Nrf2 is also necessary in p62 aggregation. Using mouse RAW264.7 macrophages, Fujita et al. reported that treatment with LPS or *E. coli* could induce LC3-II and p62 expression, as well as the formation of selective autophagy of aggresome-like induced structures; and this effect is aborted in Nrf2-deficient macrophages, indicating the necessity of Nrf2 in p62 aggregation and p62-mediated autophagy [122]. Similar phenomenon is also shown in liver cells [123] and in the septic lung [124]. In this positive feedback pattern, p62 also contributes to Nrf2 activation. There is a Keap1-interacting region (KIR) in p62, and various stresses can induce the phosphorylation of the serine residue in the KIR, which markedly increases its binding affinity to Keap1. Therefore, phosphorylated p62 can completely abrogate the interaction between Nrf2 and Keap1, thus activating Nrf2 [125]. In addition, Keap1 in the complex with phosphorylated p62 is degraded by selective autophagy [121, 126], contributing to the continuous activation of Nrf2 pathway.

So far, there is little information about the role of Nrf2 in regulating microglial autophagy in stroke, although two reports showed that hypoxia promote both Nrf2 activation and microglial autophagy [127, 128], without knowing the potential link between these two events. Nevertheless, given that phagocytosing microglia produce large amount of ROS, and that ROS is a strong inducer of Nrf2 activation and autophagy, it is reasonable to speculate that the activation of Nrf2 in microglia may promote its autophagic capacity. It is worth to note that autophagy may also be a “double-edged sword,” with side effects of further induction of inflammation and delayed cell death, etc. The degree and time of autophagy would determine the overall prognosis [129]. In this regard, further research could be helpful and focus on how Nrf2 regulates microglial autophagy after stroke, and how to precisely control the degree and time of Nrf2 activation to achieve protective effects and avoid detrimental ones.

Nrf2 Pathway Promotes Microglial Phagocytosis

Phagocytosis is a Greek-derived term. Literally, it means the cellular processes of eating, which includes the recognition, engulfment, and degradation of large (>0.5 μm) particulated organisms or structures [130]. Using a co-culture system and live imaging technique, it has been determined that microglia can eliminate an apoptotic cell in 25–95 min *in vivo* [131]. *In vitro* under physiological conditions, the microglial clearance time of apoptotic cells has been estimated from 70–90 min to 1–2 h [132, 133]. Besides clearance of apoptotic cells and debris, which is critical for the CNS homeostasis [133], phagocytosis has other functions [134], which includes: (1) antigen presentation, (2) activation of respiratory burst, which can be triggered by hypoxia/reoxygenation, inducing ROS generation that contributes to killing engulfed microorganisms and degradation of other cargo, and (3) modulation of inflammatory responses.

In stroke, phagocytosis by microglia and macrophages plays a critical role in the clearance of cellular bodies and debris, thus limiting ischemic injury [135, 136] and

promoting tissue repair [137]. As demonstrated by Lalancette-Hébert et al., selective ablation of proliferating microglia/macrophages 72 h after MCAO led to a 2.7-fold increase in the number of apoptotic cells, predominantly neurons, and exacerbated ischemic brain injury [136]. In another research carried out by Faustino et al., they reported that depletion of microglia significantly enhanced focal inflammatory responses and ROS expression [135].

Few researches focus on the role of Nrf2 pathway in regulating microglial phagocytosis in the acute or sub-acute phase post the onset of stroke, though Nrf2 pathway has been shown to promote the phagocytosis of macrophages and microglia in several other circumstances. For example, curcumin could enhance the phagocytosis of parasitic protozoans by microphage, playing an antimalarial role; and this process was mediated by the activation of Nrf2 signal pathway but not PPAR- γ [138]. Sulforaphane and benzyl isothiocyanate, two Nrf2 activators, could increase the uptake of 2- μ m diameter polystyrene beads by RAW 264.7, a line of murine macrophage-like cells, and this effect was aborted in peritoneal macrophages from Nrf2^{-/-} mice [139]. In the lung, alveolar macrophages demonstrated a decreased ability to recognize and phagocytose bacteria in chronic obstructive pulmonary disease; sulforaphane treatment could restore bacteria recognition and phagocytosis of alveolar macrophages, which was only observed in wild-type mice but not in Nrf2-deficient mice [140]. Gene expression and promoter analysis revealed that Nrf2 increased the phagocytic ability of macrophages by direct transcriptional upregulation of the scavenger receptor MARCO (macrophage receptor with collagenous structure) [140]. Within the CNS, microglial phagocytosis is also under the regulation of Nrf2 pathway. Using an in vitro model of amyloid β -induced toxicity in microglia, Li et al. reported that milk fat globule-EGF factor 8 (MFG-E8) accelerated microglial phagocytosis, associated with enhanced Nrf2 and HO-1 expression [141]. In murine intracranial hemorrhage model, activating Nrf2 pathway with sulforaphane could induce the increased expression of CD36, a phagocytosis-mediating scavenger receptor, and increased hematoma clearance, which is abrogated in Nrf2^{-/-} mice [142]. Adopted primary microglia and red blood cells in a phagocytic study, Zhao and colleges also found that activating Nrf2 pathway could upregulate CD36 expression and enhance red blood cell phagocytosis [142]. It is worth noting that the harmful condition of self-producing ROS and pro-inflammatory mediators makes microglia hard to survive; Nrf2 pathway might enhance the tolerance and survival of microglia under this detrimental milieu. As mentioned above, phagocytosis of microglia could regulate inflammation through releasing anti-inflammatory cytokines, one of which is transforming growth factor (TGF)- β , and reduced production of pro-inflammatory cytokines such as TNF- α [143]. It has been reported that TGF- β is critical for survival of phagocytizing microglia through autocrine suppression of TNF- α and ROS [144], and thus enhance microglial phagocytosis. Therefore, enhancing microglial Nrf2 may promote its phagocytosis through downregulating ROS and prolong the lifetime of phagocytizing microglia, expediting debris cleanup in ischemic brain.

Like autophagy being a “double-edged sword” in microglia/macrophage, phagocytosis in microglia/macrophage could also be recognized as such a “double-edged

sword.” Under physiological and regulated conditions, the phagocytosis helps to scavenge the apoptotic and necrotic cells, to clear the debris and thus restrict the inflammation; under pathophysiological circumstances, however, dysregulated microglial phagocytosis may contribute to excess neuronal death after acute or chronic ischemia, leading to delayed neuronal loss, brain atrophy even vascular dementia. In a MCAO research carried out by Neher et al. [145], for example, genetic deficiency of phagocytic protein MFG-E8 or Mer receptor tyrosine kinase (MerTK) could completely prevent long-term functional deficits of motor neurons, and the phagocytic deficiency strongly reduced brain atrophy as a result of inhibiting phagocytosis of neurons. The mechanism might be associated with suppressed phagocytosis of neurons, as neurons reversibly exposed the “eat-me” signal phosphatidylserine (PS) to microglia or macrophages after stroke [145].

Taken together, several lines of evidence suggest that activation of Nrf2 pathway may contribute to the inhibition of inflammatory responses after stroke. Considering that cell death, either in the form of apoptosis or necrosis, is an inevitable event in ischemic stroke, enhancing phagocytic capacity of microglia and macrophage will expedite the clean-up process and enhance tissue repair.

3 Conclusion

In summary, Nrf2 is sequestered under physiological conditions and degraded through ubiquitination. Ischemic stroke induces severe oxidative stress that activate Nrf2 pathway by disassociating Nrf2 from Keap1. After nuclear translocation, Nrf2 induces the expression of antioxidants, mainly phase 2 enzymes, which then protect neurons against oxidative injury. It is worth noting that Nrf2 may protect not only through cellular self-defense, but also through cell–cell interaction. Therefore, Nrf2 pathway could be a promising therapeutic target to treat ischemic stroke.

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Stem Cell Therapy for Ischemic Stroke

Hung Nguyen, Naoki Tajiri, and Cesar V. Borlongan

1 Therapeutic Action of Stem Cell Transplantation in Stroke

The therapeutic utility of stem cells for treating neurological disorders has been demonstrated either by stimulating these cells endogenously [1–4] as well as by transplanting them exogenously within injured brain tissues [5–11]. Stem cell transplantation after a brain insult, such as stroke, is a major tenet of regenerative medicine [1–13]. Stem cell therapy may offer a glimpse of hope for stroke patients because very scarce successful clinical trials have emerged from preclinical laboratory research in this area [14–16]. Traditionally, the postulated therapeutic mechanism underlying stem cell repair for brain diseases implicates that the ensuing cellular regeneration occurs due to either the direct replacement of necrotizing cells with transplanted stem cells or the indirect repair of damaged tissue through the secretion of trophic factors by the stem cells [17, 18]. That stem cells act via cell replacement and growth factor release regenerative processes have accompanied the translation of stem cell therapy for stroke from the laboratory to the clinic over the last 25 years. Nonetheless, these two major mechanisms behind stem cells' ability to restore the stroke brain only partially explain the paradoxical robust functional recovery in stroke animals despite low graft survival of transplanted stem cells. Recent evidence from a relevant cerebrovascular disease, traumatic brain injury (TBI), suggests that transplanted stem cells are capable of forming a cellular migratory pathway as an active process for recruiting endogenous stem cells from the

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neurogenic niche to the damaged area within the cortex [19]. This transplanted stem cell-generated “biobridge” has been visualized immunohistochemically and by laser capture microdissection to reside in the region between the neurogenic sub-ventricular zone (SVZ) and the largely non-neurogenic TBI-damaged cortex. Because endogenous stem cells have limited capacity to migrate long-distances, as in the case of traversing from the SVZ to the injured cortex of non-transplanted TBI animals, the recognition of this biobridge should facilitate the migration of endogenous stem cells to damaged brain areas that are remotely inaccessible from the neurogenic niche [19, 20].

The overlapping pathologies between TBI and stroke indicate the potential of biobridge formation in stroke following stem cell transplantation. We discuss below the progress of stem cell therapy for stroke, highlighting this possibility of extending the concept of transplanted stem cell-mediated biobridge as a therapeutic mechanism underlying stroke recovery.

2 Secondary Cell Death in Stroke as Therapeutic Target for Stem Cell Grafts

Stroke is the consequence of blood flow restriction to a region of the brain. Its signature pathological characteristic is a region of dead neuronal cells known as the infarct core, and this region is encased within a zone of injured and dying tissue called the penumbra. This cardinal feature of vascular disablement associated with ischemic stroke results in a regional deficiency of glucose and oxygen, which then results in the stimulation of intricate secondary cell death processes [21]. Such blood flow interruption to the brain is commonly referred to as ischemic stroke, whereas blood vessel bleeding is termed hemorrhagic stroke. We focus our discussion of stem cell transplantation on ischemic stroke as this pathological condition, especially the secondary cell death, offers a far wide-ranging application of stem cell therapy than hemorrhagic stroke.

The wider therapeutic window, which can last for days, weeks, months, and even years, associated with the secondary cell death of ischemic stroke has been the target of stem cell therapy. Within a matter of minutes after stroke onset, the primary necrotic core becomes fixed and unsalvageable. In contrast, the secondary cell death that ensues in the tissue surrounding the core corresponds to the evolving ischemic penumbra [22], and may be resuscitated with proper treatment intervention such as stem cell therapy.

In view of the supracute window in the formation of the ischemic core, as opposed to the subacute and even chronic progression of the penumbra, a much better prognosis for arresting the secondary cell death seems to be indicated. Sufficient circulation must be re-established in a very short amount of time, preferably less than 3 h, in order to minimize the necrotic core and to limit the evolution of the penumbra. In this regard, tPA needs to be initiated during the initial 4.5 h after

stroke onset, but beyond this acute window, the drug carries significant adverse effects in particular bleeding [23]. Conversely, when targeting the evolving penumbra, treatment interventions over a much protracted window post-stroke should allow sequestration of secondary cell death events. Indeed, stimulation of endogenous neurogenesis, angiogenesis, and neuroplasticity has been shown to minimize and even reverse penumbra-associated functional and neurostructural deficits.

3 Engaging the Endogenous Regenerative Process in Stroke

The adult mammalian brain has long been considered to lack regenerative capacity, but scientific evidence has shown that neurogenesis occurs in discreet regions of the adult brain, namely in the SVZ of the lateral ventricles and the subgranular zone (SGZ) of the dentate gyrus in the hippocampus [24, 25]. Such endogenous regeneration even in adulthood [21], which has been documented to be further stimulated during injury, suggests that the adult brain is able to mount a reparative process against stroke. However, the proliferation level of newly formed cells, their neural lineage commitment, and their ability to migrate towards the injured brain areas may be limited, altogether indicating the insufficient regeneration by the host brain to exert clinically meaningful improvement after stroke. Current investigations into finding strategies to enhance this endogenous neurogenesis following stroke has been a major research area of interest in the field of regenerative medicine. Treatment regimens designed to boost proliferation, migration, and differentiation of neural progenitor cells by modifying the host microenvironment to be more conducive for synaptic integration of the newly generated neurons have been explored, but to date, compelling evidence suggests that the host neurogenesis remains relatively confined within the SVZ and SGZ, relegating most of the applications of this endogenous repair mechanism within these neurogenic niches or only to their neighboring tissues [25, 26]. Because cerebrovascular diseases, such as stroke and TBI, present with major tissue damage to brain areas remote to the neurogenic niches, utilizing endogenous neurogenesis against these neurological disorders will need consideration.

Along the lines of mobilizing endogenous neurogenesis, ischemic stroke has been demonstrated to upregulate specific molecules that stimulate neurogenic niches, in particular increasing neural progenitor cell survival, proliferation, migration, and differentiation [21]. Moreover, these mobilized neural progenitor cells (albeit a small population) have been shown to migrate from the SVZ to the post-ischemic striatum in rat stroke models [27, 28]. In tandem, resident stem cells which remain quiescent during homeostasis, but are awakened after stroke, have been detected in the hippocampus and the cerebral cortex. In the clinic, however, post-mortem analysis of the stroke brain has not offered evidence of newly generated cortical neurons during the time period of 3 days to 13 years after stroke onset [29]. A rough estimate of host neurogenesis secondary to stroke has been calculated at

0.1%, or 1 out of every 1000 neurons [29], suggesting that relying merely on the scarcity of newly born cells and their capacity to migrate from the neurogenic niche to the injured area may not be feasible when contemplating endogenous neurogenesis as a regenerative process against stroke.

4 Biobridge: A Migratory Pathway Between Neurogenic Niche and Non-neurogenic Brain Areas

As noted in the preceding section, the two generally accepted mechanisms of action of stem cell transplantation which provides the therapeutic effects are cell replacement and modification of the microenvironment through trophic factors [17, 18]. The new biobridge concept was first described in a study of transplanting modified mesenchymal stromal cells (MSCs) in TBI model [19]. Adult rats were exposed to controlled cortical impact (CCI) TBI and were treated with MSCs. The MSCs formed a biobridge between the neurogenic niche of the host and the injured site, and served as a highway for the host endogenous stem cells to travel to the injured area [19]. The biobridge which spanned from the SVZ to the damaged cortex was visualized using immunohistochemistry and laser capture assay. Upon closer examination of the biobridge, the levels of extracellular matrix metalloproteinases (MMPs) were significantly increased. In particular, MMP-9 was found significantly increased within the biobridge, suggesting that this MMP may play an important role in facilitating the migration of neural progenitor cells in response to a brain insult, which likely released specific chemoattractant signals from the injured area. This study further suggests that an inflammation-enriched (albeit cytokines, chemokines, MMPs) mechanism closely mediates that formation of biobridge. Additional experiments are needed to better understand the effects of various inflammatory cues on the formation of biobridge. Interestingly, the grafted MSCs are eventually replaced by the host cells after a period of time [19], which is a welcomed translational safety outcome because the prolonged and unregulated survival and proliferation of stem cells may be associated with tumor formation. Moreover, that long-term graft survival per se is not a prerequisite robust and stable stroke recovery addresses the paradoxical dilemma on the reported studies demonstrating solid functional improvements despite mediocre engraftment.

A series of analyses to further characterize the biobridge reveals that MMPs assisted the grafted MSCs to form the biobridge. Once the biobridge was formed, the transplanted cells were replaced by the endogenous neurogenic niche-derived stem cells [19]. The biobridge guided the endogenous stem cells to successfully migrate from the SVZ and to home at the TBI injured site. The stem cells then populated the peri-impacted area and exerted their effects in this area. This novel concept of biobridge formation after transplantation is not limited to TBI but also can be extended to other neurological disorders, such as stroke. The damage areas in stroke are generally larger compared to TBI thus it may be more challenging for endogenous stem cells to migrate across tissues toward the injured site. The biobridge by

providing an extracellular matrix on which the endogenous stem cells could attach and migrate should improve in directed transport of neurogenic niche-derived cells to the remote injured areas [19].

5 The Molecular Composition of the Biobridge

The formation of the biobridge provides an alternative and complementary repair mechanism in stem cell therapy, in addition to the cell replacement and growth factor release regenerative processes. The biobridge allows a favorable microenvironment for cell migration and axonal growth. Three months after transplantation in the TBI brain, an increased level of proliferation and differentiation was observed at the peri-impacted area and a stream of cells labeled with nestin and DCX, markers for maturing neuron was maintained [19]. In contrast, control animals which received vehicle did not express significantly, the level of migration nor differentiation. In addition, the level of expression and activity of MMP-9 was upregulated ninefold which directly correlated with the endogenous stem cell migration compared with the animals that received the vehicle infusion [19]. Recent studies have demonstrated the significance of MMPs and extracellular matrices (ECMs) in stroke pathology [2, 12] which further validates the importance of biobridge formation in stem cell therapy for stroke. In the same vein, we envision that the grafted stem cells via a similar biobridge may promote the migration of stem cells from endogenous niche to the stroke core and infarct area. Stem cells, such as peripheral blood, umbilical cord blood, and adult brain stem cells, are able to modulate the levels and functions of MMPs and ECMs [30–32]. With MMPs and ECMs involved in stroke pathology, the ability of stem cells to generate a biobridge in stroke warrants investigations. A study has shown that inhibition of MMPs sequestered the migration of endogenous stem cell from the niche to the damaged site [33], likely due to the depletion of neurovascular network remodeling. On the other hand, the formation of a biobridge may stimulate the production of MMPs, which in turn can enhance the restoration of the neurovascular unit and the migration of host endogenous stem cells. It is worth noting that the physiochemical homeostasis between cell deformation, tissue barrier restriction, and migration rates correspond to limiting factors that determine the capacity of cell migration. In turn, this physiochemical homeostasis is regulated by the ability to degrade the ECM by proteolytic enzymes such as MMPs, and integrin- and actomyosin-mediated mechanocoupling [34].

Based on the concept of biobridge in the TBI model [19], we hypothesize that a similar biobridge will be realized after an intracerebral stem cell transplantation in the stroke brain. The biobridge will similarly guide the endogenous stem cells to migrate from the neurogenic niches, such as the SVZ and the SGZ, to the damaged areas in stroke brain, and eventually contributing to functional improvement [35]. Of note, laboratory studies have reported that transplantation of MSCs aids in stroke recovery, with modulation of inflammation and immune response as a postulated

therapeutic mechanism [36]. Indeed, it has been shown that transplanted stem cells can secrete transforming growth factor-beta (TGF- β) that can dampen the spreading of the inflammatory MPC-1 cells [36]. To this end, transplanted MSCs via formation of the biobridge can also secrete anti-inflammatory or immunomodulatory cytokines, termed as MSC secretome, which can mediate the therapeutic benefits in the stroke brain [37]. The MSC secretome may consist of insulin-like growth factor (IGF)-1, stromal cell-derived factor (SDF-1 α), and glial cell-line-derived neurotrophic factor (GDNF) [38], which are growth factors that are elevated in the stroke brain after intracerebral transplantation of MSCs [38].

6 Translational Caveats of Biobridge Application to the Clinic

Despite the demonstration of the biobridge in mediating stem cell therapy TBI animal models, there are limitations in translating this concept to the clinic. Additional studies are needed to further characterize the biobridge facilitation of endogenous stem cell migration. In particular, the formation of such biobridge after peripheral transplantation of stem cells is a missing endeavor that will be of practical use in the clinic, as a minimally invasive procedure may be more preferred instead of a direct intracerebral transplantation approach when targeting the subacute phase of TBI and stroke models.

Another limitation of the biobridge is that the formation itself depends on the severity of the injury. Severe TBI or chronic stage of stroke presumably presents with more extensive damage tissue which may influence the biobridge formation. The pathology of TBI and stroke are also complicated and varied from patient to patient based on the severity of the insult, requiring tailoring of stem cell regimen, e.g., timing of cell transplant initiation, number of transplantations over the progressive period of secondary cell death, among others that may affect the creation of a biobridge. Moreover, the formation of biobridge might be influenced by the non-conductive nature of the microenvironment in the brain, necessitating the need to treat the severe cases of TBI and stroke in the early and acute stage to immediately halt the progression of the disease and arrest the evolution of a non-favorable host microenvironment.

The inflammatory responses accompanying TBI and stroke also warrant investigations, in order to assess the effects of this secondary cell death on biobridge formation. While prolonged inflammation has detrimental effect, acute inflammation may support the brain (i.e., biobridge). The timeline when inflammation produces beneficial effects then switches into a destructive mode remains debatable. Arguably, transplanting at an earlier period post-insult may aid in the biobridge formation and may enhance the pro-survival over the pro-death properties of inflammation. Additional studies are needed to understand the critical time point to maximize the therapeutic effect of inflammation on stem cell transplantation and its modulation of biobridge formation.

7 Towards a Safe and Effective Stem Cell Therapy for Stroke

According to the American Heart Association, approximately 800,000 Americans have stroke every year, with about 17 % mortality or 1 in every 18 deaths in the US is a result of stroke [39]. Stroke survivors also experience disability ranging from moderate to severe. The total estimated healthcare costs due to disability related to stroke are \$18.8 billion annually. The number even approaches \$34.3 billion if an additional \$15.5 billion is added for loss of productivity and premature deaths [40]. Despite the fact that the mortality rate of stroke has fallen 33.5 % in 10 years period from 1996 to 2006, the cost of healthcare associated with stroke increases significantly for the past decades [39].

As the cost continues to increase, there is an urgent need for more effective treatment for neurological disorders such as TBI and stroke. Regenerative medicine has put stem cell therapy as a promising treatment for CNS disorders. The traditional mechanisms of action of stem cell therapy include cell replacement and bystander effects [41–44]. Our recent study has shown another novel mechanism of stem cell transplantation therapy, the formation of biobridge, which provides a cellular pathway for the exogenous transplanted cells to attract endogenous stem cells to migrate from the neurogenic niche towards remote damaged brain areas. This concept of biobridge may be extended in transplant studies to other CNS disorder, such as Parkinson's disease and spinal cord injury, whereby endogenous stem cells may require long-distance migration from the neurogenic niche to the injured tissues.

In conclusion, the multi-pronged therapeutic actions of stem cell therapy, such as cell replacement, secretion of growth factors, and biobridge formation may work in concert to provide beneficial outcomes in cerebrovascular diseases [19]. The formation of biobridge provides a cellular link between the exogenous repair mechanism and the endogenous regenerative processes. That stem cell transplantation may involve different biological pathways in affording its therapeutic effects caters to the multiple cell death events associated with stroke. In the end, an understanding of the mechanism of action that mediates stem cell transplantation may aid in optimizing the stem cell dose, timing of administration, and route of cell delivery, and the eventual laboratory-to-clinic translation of a safe and effective stem cell therapy for stroke.

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