



THE NEUROVASCULAR UNIT AS A POTENTIAL BIOMARKER AND THERAPEUTIC TARGET IN CEREBROVASCULAR DISEASE

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THE NEUROVASCULAR UNIT AS A POTENTIAL BIOMARKER AND THERAPEUTIC TARGET IN CEREBROVASCULAR DISEASE

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Editorial: The Neurovascular Unit as a Potential Biomarker and Therapeutic Target in Cerebrovascular Disease

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Keywords: neurovascular unit (NVU), cerebrovascular disease, biomarker, cerebral blood flow (CBF), therapeutic target

Editorial on the Research Topic

The Neurovascular Unit as a Potential Biomarker and Therapeutic Target in Cerebrovascular Disease

INTRODUCTION

The brain is one of the most metabolically demanding organs—it requires 20% of the body's energy supply yet forms only 2% of the body's weight. Therefore, brain activity is supported by a dynamically regulated supply of blood and, hence, energy substrates through the cerebrovascular network. Furthermore, the post-mitotic nature of neurons and their longevity makes the brain very sensitive to extracellular perturbations. Thus, the cerebrovasculature also has to be specialized to minimize perturbations of the brain microenvironment, a property mediated by the blood-brain barrier that is formed by the neurovascular unit (NVU).

The NVU is a multicellular structure formed by neurons, glia and vascular cells. The NVU is fundamental to the distribution of cerebral blood flow, regulation of extracellular homeostasis and, as recently shown, waste clearance from the brain. Therefore, the NVU represents a pivotal point of vulnerability for the brain. In this Research Topic, 12 contributing articles address the potential of the NVU and provide evidence demonstrating the importance of the NVU as a biomarker and therapeutic target for cerebrovascular disease.

PHYSIOLOGICAL PARAMETERS OF THE NVU AS BIOMARKERS AND TARGETS FOR THERAPEUTIC INTERVENTION

It is important to first understand the structure and function of the healthy NVU to identify abnormal events and therapeutic targets. Shaw et al. provide a thorough and detailed characterization of the cerebrovascular bed, highlighting differences in its structure and function as it branches from penetrating arterioles to capillaries. Using immunohistochemistry and by assessing vasomotion along the vascular tree in awake mice *via* two-photon microscopy, they provide novel data characterizing points of contractility, angiogenic capacity, and vasodilatory potential across the vascular tree and branch points (Shaw et al.). The differences in vasodilatory response across the vascular tree are also highlighted in the study by Rosehart et al., in which they show that stimulation of capillaries with prostaglandin E2 elicits vasodilation of upstream arterioles. This was blunted in a model of small vessel disease, suggesting a reduction in capillary-to-arteriole signal propagation as a potential mechanism for the pathologies associated with cerebral small vessel disease, while also suggesting a new biomarker for NVU dysfunction (Rosehart et al.).

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Specific modifiable physiological parameters of the NVU are also highlighted within this Research Topic. Inhibitors of carbonic anhydrases, the pH regulating enzymes, can protect the NVU in stroke and Alzheimer's disease, as reviewed by Lemon et al. Hayes et al. provide novel data on another parameter, the neuroprotective hormone insulin-like growth factor-1 (IGF-1), demonstrating a difference in vulnerability to glutamate-induced toxicity across different cell types of the NVU when IGF-1 is inhibited.

Together, these data highlight multiple structural and functional points within the NVU that may potentially serve as targets for therapeutic intervention. The differential regulation at the various vascular segments and the transitional zones between them in the vascular tree is also noteworthy; each segment could be differently affected during disease, or be targeted for therapy.

NVU RISK FACTORS AND BIOMARKERS ACROSS THE LIFESPAN

The biomarker and target potential of the NVU across the lifespan is another key focus of this topic. In an opinion piece, Beishon and Panerai emphasize the importance of mid-life therapeutic targeting and lifestyle alterations targeting known risk factors for dementia. The comprehensive review by Ouellette and Lacoste highlights the shared vascular abnormalities associated with neurodevelopmental and neurodegenerative disorders, despite their distinct clinical presentation at different life stages.

Lecordier et al. provide a detailed review of early risk factor events for NVU dysfunction leading to dementia, highlighting triggering pathological events. They specifically underscore the importance of air pollution as a risk factor for dementia, and provide a detailed summary of its impact on NVU components (Lecordier et al.). Exercise is suggested to reduce cerebrovascular disease and alleviate the risk for developing dementia. Ohtomo et al. report that, in a model of chronic hypoperfusion, exercise was able to alleviate fewer behavioral deficits in middle aged mice compared to their previous study on young mice, suggesting the value of early lifestyle interventions. Altogether, these reports showcase the potential of the NVU as a biomarker and the importance of protecting NVU function across the lifespan.

CLINICAL BIOMARKERS OF DISEASE OUTCOME AND PROGRESSION

The significance of using the NVU, or the pathologies associated with it, as a biomarker and therapeutic target in the clinical setting is also emphasized in this Research Topic. A systematic meta-analysis by Wang et al. demonstrates the role of cerebral small vessel disease in negative prognosis following intravenous thrombolysis treatment for acute ischemic stroke, highlighting the value of NVU pathology in predicting treatment outcomes.

In addition, Monteiro et al. report a reduction in neurovascular coupling in patients with hypertension while hypercapnic hyperemia is preserved, prior to detectable cerebral small vessel disease. This impairment was further exacerbated when hypertension was comorbid with diabetes,

suggesting that NVU dysfunction may contribute to the cerebral effects of chronic hypertension and diabetes (Monteiro et al.). Staszewski et al. provide further support for the NVU as a biomarker for disease progression, demonstrating that decreases in cerebrovascular reactivity are measurable over a 24 month follow-up period in patients with small vessel disease, irrespective of initial radiological disease assessment.

Lastly, Ren et al. suggest that disruptions in the glymphatic system, a hypothesized route of paravascular waste clearance, may be the cause of, and serve as a biomarker for, peri-operative neurocognitive disorder. Their review highlight anesthesia choice and management as potentially modifiable risk factors for peri-operative neurocognitive disorder (Ren et al.).

CONCLUSION AND FUTURE DIRECTIONS

This Research Topic provides both preclinical and clinical evidence of the importance of the NVU as a contributor to cerebrovascular disease and, thus, a therapeutically targetable fulcrum, while underscoring its value as an accessible biomarker for disease etiology and progression. The additional targeting of co-morbidities and detrimental lifestyle risk factors that lead to NVU dysfunction, which change over the lifespan, are also essential for disease management. It is clear that further characterization of the NVU, including its functions in health and different pathological contexts, and how it is modified by lifestyle choices and comorbid conditions, are all important avenues for future work. Findings from such studies may suggest potential therapeutic targets for cerebrovascular diseases, and may even be applicable to related dementia conditions.

AUTHOR CONTRIBUTIONS

All authors listed have made a substantial, direct, and intellectual contribution to the work and approved it for publication.

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Dysfunction of the Glymphatic System as a Potential Mechanism of Perioperative Neurocognitive Disorders

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Perioperative neurocognitive disorder (PND) frequently occurs in the elderly as a severe postoperative complication and is characterized by a decline in cognitive function that impairs memory, attention, and other cognitive domains. Currently, the exact pathogenic mechanism of PND is multifaceted and remains unclear. The glymphatic system is a newly discovered glial-dependent perivascular network that subserves a pseudo-lymphatic function in the brain. Recent studies have highlighted the significant role of the glymphatic system in the removal of harmful metabolites in the brain. Dysfunction of the glymphatic system can reduce metabolic waste removal, leading to neuroinflammation and neurological disorders. We speculate that there is a causal relationship between the glymphatic system and symptomatic progression in PND. This paper reviews the current literature on the glymphatic system and some perioperative factors to discuss the role of the glymphatic system in PND.

Keywords: glymphatic system, postoperative complications, postoperative neuropathy, postoperative cognitive dysfunction, perioperative neurocognitive disorders

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INTRODUCTION

Perioperative neurocognitive disorder (PND), which encompasses delirium and postoperative cognitive dysfunction, commonly occurs in the elderly after anesthesia and surgery. Previously, all forms of postoperative cognitive impairments were called “postoperative cognitive dysfunction,” but more recently, the use of “perioperative neurocognitive disorder” is recommended as an overarching term for cognitive impairment identified in the preoperative or postoperative period (Evered et al., 2018a). This change aligns PND with the phenotypically similar neurocognitive diagnoses listed in the Diagnostic and Statistical Manual of Mental Disorders, version 5, such as Alzheimer's disease (AD) (Evered et al., 2018a). Furthermore, PND has emerged as a significant global public health issue that leads to more extended in-hospital stays, higher hospitalization costs, and higher mortality rates (Yang and Terrando, 2019). The risk factors for PND are multifaceted, which might be related to anesthetic management, tissue damage, surgery duration, psychological stress, and genetic susceptibilities (Subramanian and Terrando, 2019; Eckenhoof et al., 2020). As the elderly population continues to increase, the number of cases of PND will continue to rise. The incrementally growing prevalence forces researchers to explore the mechanisms underlying the pathogenesis of PND and to seek optimal prevention and treatment measures (Evered and Silbert, 2018; Berger et al., 2019).

Preclinical and clinical studies support the notion that neuroinflammation plays a significant role in the pathogenesis of PND (Nathan, 2019; Yang et al., 2019, 2020). Trauma experienced during surgery triggers the release of endogenous factors known as damage-associated molecular

patterns that can activate immune cells such as neutrophils and monocytes, promote pro-inflammatory cytokines, and subsequently cause systemic inflammation (Huber-Lang et al., 2018). Uncontrolled systemic inflammation is associated with a compromised blood-brain barrier (BBB). BBB impairment promotes immune cell movement and pro-inflammatory cytokines from the blood into the brain parenchyma, triggering neuroinflammation and ultimately leading to postoperative cognitive impairment (Abrahamov et al., 2017; Yang et al., 2017, 2020). Microglia, the resident immune cells of the central nervous system (CNS), perform “immune surveillance,” and survey their assigned brain regions (Kettenmann et al., 2011). In addition, with prolonged systemic inflammation, microglia start to develop an activated phenotype characterized by an increase in pro-inflammatory mediators, such as interferon- γ , interleukin (IL)-1 β , tumor necrosis factor (TNF)- α , and reactive oxygen species (Liu L. R. et al., 2020). Pro-inflammatory factors released by microglia such as IL-1 α and TNF contribute to the subsequent activation of astrocytes and further promote neuroinflammation (Liddelow et al., 2017; Liu L. R. et al., 2020). Complement system activation is another essential inflammatory response that is activated by surgically triggered damaged-associated molecular patterns. For example, the deposition of the C-reactive protein, a biomarker for delirium, can activate and regulate the classical complement pathway, thereby contributing to dysregulated inflammation. Blocking the complement cascade (e.g., complement C3 gene deficiency and selective complement inhibitor for complement C3) improves neuroinflammation and functional outcomes of PND (Alawieh et al., 2018) also suggests the contribution of neuroinflammation in PND.

Several studies have shown that the pathological mechanism of PND is similar to that of AD (Evered et al., 2016; Gerlach and Chaney, 2018). It was also demonstrated that a significant accumulation of amyloid- β (A β) and tau in the brain parenchyma after anesthesia and surgery (Terrando et al., 2011). Moreover, increased A β and tau in the cerebrospinal fluid (CSF) have been reported in patients after surgery (Evered et al., 2018b), which are risk factors for PND (Xie Z. et al., 2013; Evered et al., 2016; Cunningham et al., 2019). The accumulation of A β and tau induces neuroinflammation, leading to glial activation, pro-inflammatory factor release, and neuronal damage (Calsolaro and Edison, 2016).

The glymphatic system is a recently discovered waste removal system that utilizes a unique perivascular channel system formed by astrocytes (Iliff et al., 2012; Benveniste et al., 2017; Zhang C. et al., 2018). Although the glymphatic system contributes to the delivery of nutrients, specifically glucose, its most influential and recognized function is to clear extracellular metabolites and waste products from the parenchyma into the CSF (Abbott et al., 2018; Nedergaard, 2013; Simon and Iliff, 2016). Since its discovery, researchers have proposed that waste or protein aggregates induced by glymphatic dysfunction are associated with neurodegenerative diseases, including AD (Tarasoff-Conway et al., 2015; Harrison et al., 2020; Nedergaard and Goldman, 2020). Given the characteristics of the glymphatic system and the complexity of possible changes during the perioperative stage, we herein speculate a causal relationship

between the glymphatic system and symptomatic progression in PND. Glymphatic dysfunction in the perioperative period, which leads to waste accumulation, could trigger or exacerbate neuroinflammation and eventually lead to cognitive dysfunction. Through this review, we discuss the function and driving mechanism of the glymphatic system, outline the current evidence to illustrate the impact of anesthesia and surgery on the glymphatic system, and emphasize the viewpoint that glymphatic dysfunction is involved in the pathogenesis of PND.

GLYMPHATIC PATHWAY

The glymphatic system is a perivascular network that subserves a pseudo-lymphatic function throughout the brain (Figure 1). Iliff et al. named this brain-wide fluid transport pathway the glial-associated lymphatic system or glymphatic system due to its dependence on glial water flux and the lymphatic function of the brain (Iliff et al., 2012). As the macroscopic waste clearance system of the CNS, the glymphatic system's fundamental role is to eliminate soluble proteins and metabolites, including A β and tau, based on the evidence that 40%–80% of neurotoxic compounds can be cleared from the CNS via this system (Iliff et al., 2012). The neurotoxic compound clearance process within the glymphatic system is described as a 3-step serial process (Benveniste et al., 2019b). First, CSF is continuously transported from the subarachnoid space and Virchow-Robin space to the peri-arterial spaces in a bulk-flow driven manner; subsequently, CSF is propelled from the peri-arterial compartment into the interstitial fluid (ISF) space. The convection and mixing of CSF and ISF are facilitated by aquaporin 4 (AQP4) in the dense and complex brain parenchyma. Ultimately, CSF-ISF fluid mixes with interstitial waste solutes and is subsequently transported to the perivenous space, from the meningeal lymphatic vessels to the lymphatic vessels and circulatory system (Plog and Nedergaard, 2018).

AQP4 is required to maintain the glymphatic function and is abundantly expressed in the end-feet of astrocytes that surround arteries and veins. It is also anchored to the astrocyte membrane by the carboxyl terminus of α -syntrophin. Syntrophin-dependent AQP4 mediates the bidirectional transport of water across the brain-blood interface (Amiry-Moghaddam et al., 2003). As a bidirectional channel, the effects of AQP4 on glymphatic function may be diverse and require further investigation. Although the role of AQP4 in promoting ISF clearance is debated, a large body of evidence suggests that it facilitates CSF movement from perivascular spaces into the interstitial space and activates flushing of ISF (Smith et al., 2017). Studies on different strains of AQP4-knockout mice have suggested that AQP4 is necessary for the rapid movement of CSF from the perivascular space into the ISF and through the brain (Mestre et al., 2018a). Moreover, AQP4 reduces the resistance to CSF movement from the periarterial space into the interstitium and subsequently from the interstitium into the perivenous space (Groothuis et al., 2007). The flow of CSF via the glymphatic pathway is an essential part of the CNS that eliminates metabolic waste products, such as A β and lactic acid, from deep inside the brain and delivers

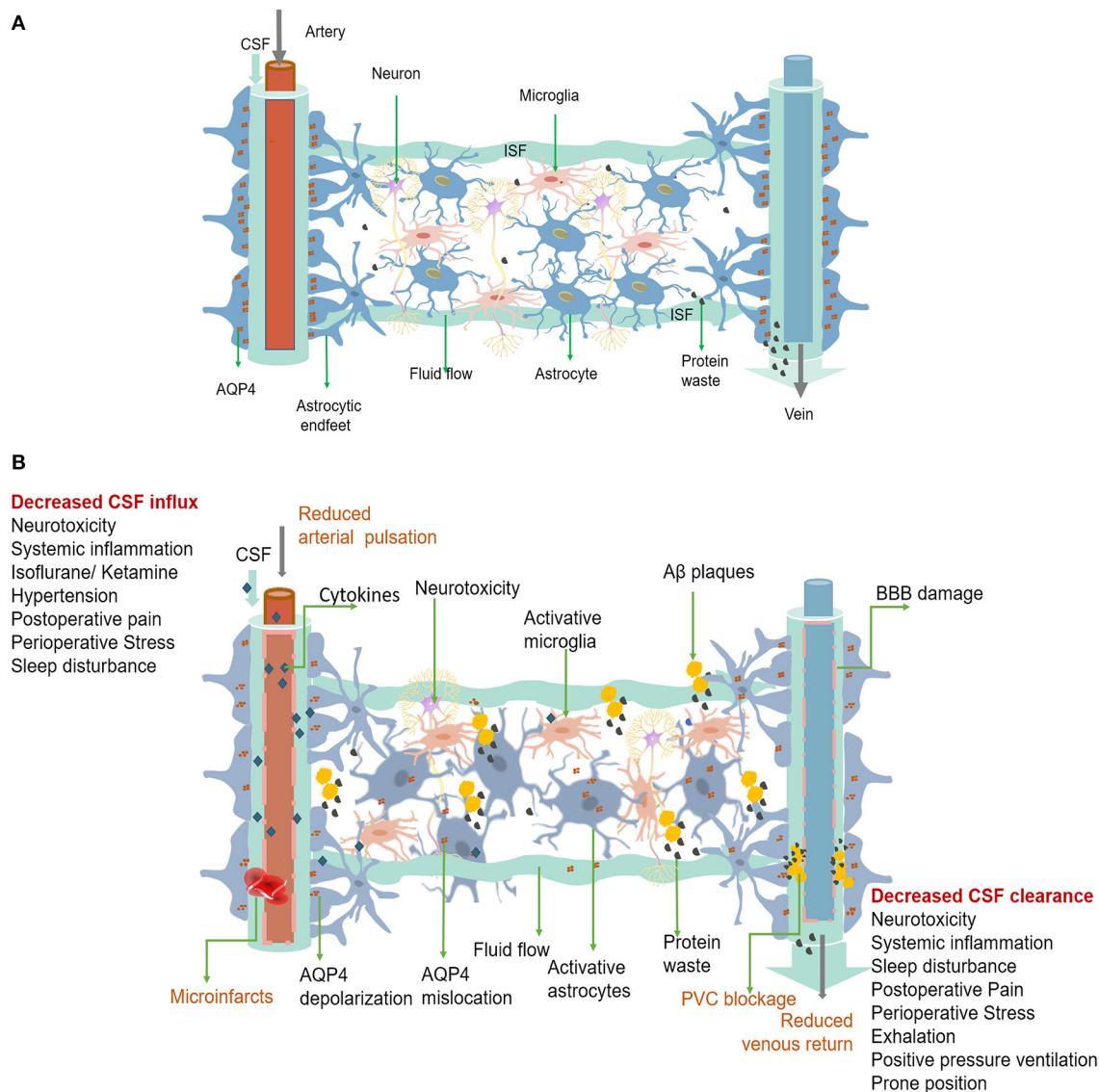


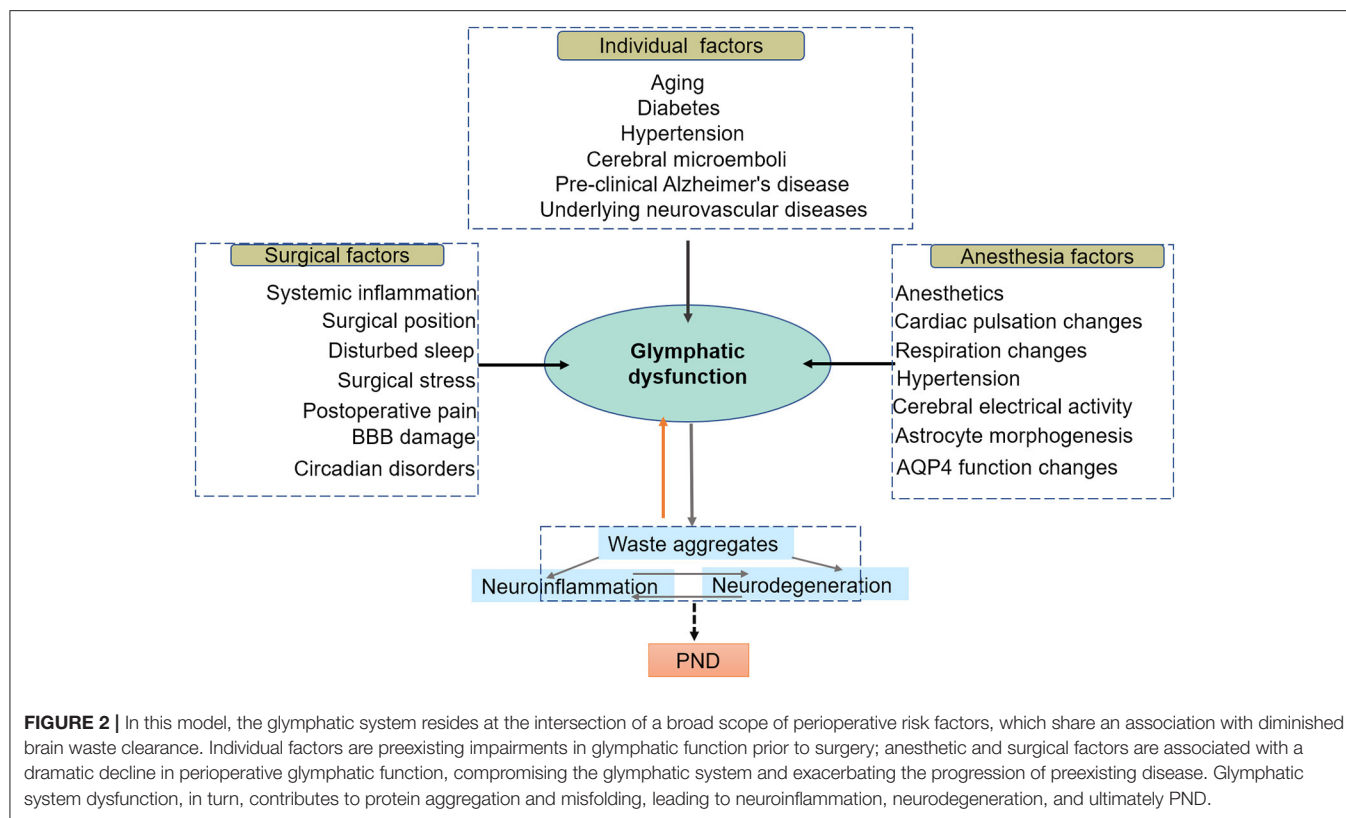
FIGURE 1 | (A) The glymphatic system is a perivascular channel formed by astrocyte end-feet throughout the brain. CSF enters the brain parenchyma through the periarterial space, exchanges with ISF, and finally exits through the perivenous space. Rapid exchange of CSF within ISF is facilitated by AQP4, which is anchored to the astrocytic end-feet. Interstitial solutes, including protein waste, are drained from the brain with CSF through the perivenous space and via the meninges and cervical lymphatics. **(B)** Dysfunction of the perioperative glymphatic system. Perioperative anesthetic drugs can cause hemodynamic changes that reduce arterial pulsation mechanism change and decrease the inflow of the glymphatic system. Surgically induced systemic inflammation can cause blood-brain barrier opening and glymphatic system damage, leading to neuroinflammation and decreased waste clearance. Both the entry of peripheral inflammatory substances and the accumulation of protein wastes in the brain, such as A β accumulation and folding, can activate astrocytes and microglia and trigger neuroinflammation. Neuroinflammation can worsen the damage to the function and structure of the glymphatic system. Forceful expiration, positive pressure ventilation, and prone position can cause a decrease in venous return, leading to a decrease in CSF clearance. Pain, preoperative stress, and sleep disturbances can affect both CSF inflow and clearance. Glymphatic dysfunction can lead to a more significant accumulation of protein and waste products, which can trigger neuroinflammation and lead to PND. PVC, Perivascular space.

nutrients, such as glucose, or therapeutic drugs to the brain parenchyma. This CSF circulation differs from typical lymphatic vessels and acts as a pseudo-lymphatic drainage within the CNS. Therefore, more recently, alternative CSF circulation in the brain has been proposed as a glymphatic circulation (Benveniste et al., 2019b).

THE DRIVING MECHANISM OF GLYMPHATIC SYSTEM

Arterial Vascular Pulsation

Unlike other peripheral organs, limited tissue compliance of the rigid skull facilitates the propagation of arterial pressure



pulsations throughout the brain, resulting in measurable pulsatile blood flow and venous outflow in the microvasculature (Shi et al., 2020). This cardiac pulsation along the entire vascular bed is believed to be the most critical physical mechanism for glymphatic propulsion, facilitating CSF movement in the glymphatic system (Kyrtos and Baras, 2015; Kiviniemi et al., 2016; Hablitz et al., 2020). Cardiovascular pulsation is also the fastest driving mechanism of the glymphatic system, originating in the basal peri-arterial spaces around the Willis's circle and extending centrifugally toward the cerebral cortex (Kiviniemi et al., 2016). The dynamic source is mainly due to the oscillation of the arterial wall caused by the heartbeat, which produces a net flow consistent with the blood flow direction and the same frequency as the cardiac cycle (Iliff et al., 2013; Kiviniemi et al., 2016). Each cardiac cycle involves a fast expansion of the artery wall, followed by a slow contraction. The bulk flow rate of CSF, calculated by CSF viscosity and the shape, cross-sectional area, and length of the vessels, is proportional to the hydraulic pressure drop along the vessels and inversely proportional to the hydraulic resistance (Thomas, 2019). The combined effects of diffusion and advection also play a role in clearing solutes from the brain (Thomas, 2019). The arterial wall pressure generates the hydraulic pressure of the CSF, which allows CSF to be delivered to the brain via the glymphatic system, a process in which the contractility of the vessel wall or the slow wave of vasomotor tension also plays an important role (Kiviniemi et al., 2016). Any reduction in vessel wall contractibility or

slow contraction in vasomotor tone increases backflow and decreases net flow in the glymphatic system (Mestre et al., 2018b). CSF flow may be mechanically regulated by pressure differences, which may be a neurophysiological regulatory mechanism. For example, astrocyte calcium activity has been shown to propagate along blood vessels as waveforms and regulate water permeability and ion exchange in the perivascular space (Rangroo Thrane et al., 2013). Additionally, intracranial pressure and heart rate are related to cardiac pulsation and affect glymphatic function. Increased intracranial pressure (ICP) decreases the mean arterial pressure and impairs the function of the glymphatic system (Chen et al., 2018), while glymphatic influx correlates negatively with heart rate (Hablitz et al., 2019).

Research has shown that the glymphatic cross-section is a non-concentric elliptical space outside the vessel, which provides the least hydraulic resistance (Tithof et al., 2019). This cross-section model is quite different from the usual circular annulus models, which assume that the small arteries are located in the center of the space (Mestre et al., 2018b; Thomas, 2019). Moreover, Mestre et al. have shown that the glymphatic system infrastructure around pial arteries is 10 times larger *in vivo* than previous estimates after fixation based on electron microscope images (Tithof et al., 2019). As such, the glymphatic system offers much less viscous resistance to CSF flow than previously thought. These results also demonstrate the importance of further *in vivo* studies on the glymphatic system.

Respiratory Forces

Advances in imaging and computational techniques have demonstrated that the respiratory mechanism is another important process that dominates the perivenous space (Kiviniemi et al., 2016). Although the respiratory force is not the primary driver of CSF flow, it is a modulating factor that supplements cardiac pulsation (Dreha-Kulaczewski et al., 2015). Respiratory dynamics are related to extreme low-frequency vasomotor oscillations during perivenous fluid movements (Kiviniemi et al., 2016). A study using magnetic resonance spin labeling in humans showed that CSF movement was enhanced during deep inhalation and was inhibited during deep exhalation (Yamada et al., 2013). A possible explanation is that forced inspiration causes blood to be drawn from the brain through the veins into the thorax, resulting in a decrease in intracerebral venous pressure, an expansion of the perivenous space, and an increase in CSF outflow from the perivenous space into the brain. Moreover, small but rapid CSF movement was observed during breath-holding, which is thought to reflect cardiac pulsations (Yamada et al., 2013). Alternatively, exhalation reduces venous outflow from the brain in a low-pressure venous drainage system (Kiviniemi et al., 2016). Additionally, respiration produces lower-frequency ICP oscillations and hydraulic pressure changes (Dreha-Kulaczewski et al., 2015).

Circadian Regulation

Sleep is essential for the maintenance of healthy brain functions. Hence, early research suggests that glymphatic clearance is mainly associated with the sleep-wake cycle (Xie L. et al., 2013). Studies in rodents have shown that CSF absorption in the perivascular space and ISF flushing increase during sleep (Xie L. et al., 2013). Glymphatic activity is regulated by circadian regulation, with a daily rhythm that peaks at midday when mice are mostly likely to sleep (Hablitz et al., 2020). Similarly, glymphatic activity was more significant at night than during the day in patients administered with gadolinium as a CSF tracer to evaluate glymphatic function (Eide and Ringstad, 2019). The mechanism by which the circadian rhythm of the brain regulates glymphatic activity remains unclear. Circadian neural activities can be detected with an electroencephalogram (EEG) device (Dijk, 1999). Overnight EEG has shown that delta (1–4 Hz) power and sigma (12–15 Hz) power are high during non-rapid eye movement (NREM) sleep and low during rapid eye movement (REM) sleep. In contrast, beta (23–30 Hz) power is higher in REM sleep than in NREM sleep, except for an artifactual peak during cycle 2. EEG activity that resembles sleep or wakefulness tightly correlates to the glymphatic influx (Hablitz et al., 2019). The brain's waste clearance mainly occurs during the NREM stage, while the CSF influx was strongly suppressed during wakefulness (Hablitz et al., 2019, 2020). Sigma power during NREM fluctuates reciprocally with delta, becoming high when delta is low and low when delta is high (Campbell, 2009). It has been demonstrated that either high delta rhythm or slow oscillation (<1 Hz), which is a characteristic of NREM sleep, is positively correlated with the glymphatic function, while beta power that resembles wakefulness is negatively correlated with the glymphatic function (Hablitz et al., 2019). The

underlying relationships between EEG power and glymphatic activity remains unclear. A recent study has revealed a coherent pattern of oscillating electrophysiological, hemodynamic, and CSF dynamics during NREM sleep (Fultz et al., 2019). These results suggest that neural activity may affect the function of the glymphatic system through hemodynamic circadian rhythms (Figure 2). Lactate concentration in the brain also closely corresponds to EEG activity and is considered the best metabolic biomarker of the sleep-wake cycle (Lundgaard et al., 2017). Lactate concentration decreases during the transition from wakefulness to sleep, leading to an expansion of the extracellular space and a decrease in intra-tissue resistance results, contributing to faster CSF influx into the interstitium and finally solute efflux.

GLYMPHATIC FUNCTION AFFECTED BY ANESTHETICS

Currently, most animal experiments on glymphatic function are performed by applying anesthetic drugs to simulate the sleep state. The effect of anesthetics cannot be ignored when studying the effect of sleep on glymphatic function. Currently, limited research has been conducted to investigate the effects of anesthetics or sedatives on the rodent glymphatic system. A study demonstrated that inhaled isoflurane (2–2.5%) inhibited CSF circulation and waste clearance in the brain (Gakuba et al., 2018). This result indicates that inhaled isoflurane inhibited glymphatic function, especially at a high dose (3%) (Gakuba et al., 2018). This research also showed that ketamine, a common intravenous anesthetic, also inhibited glymphatic activity at an intraperitoneal dose of 150 mg·kg⁻¹ in mouse models (Gakuba et al., 2018). The inhibitory effect of ketamine on glymphatic function could be reversed using a combination of ketamine and 10 mg·kg⁻¹ of xylazine (Gakuba et al., 2018). However, Xie et al. previously found that ketamine (100 mg·kg⁻¹) combined with xylazine (20 mg·kg⁻¹) significantly increased the influx of the CSF tracer in all mice analyzed (Xie L. et al., 2013). The opposite effects of the two different anesthetic regimens require further study. One possible explanation is whether ketamine inhibits or increases lymphatic inflow, depending on the dose of ketamine and its combination with xylazine, an α_2 -adrenergic agonist and an analog of clonidine used as a sedative and analgesic in animals. Xylazine increases glymphatic CSF influx and may remedy the impairment of ketamine use in the glymphatic system (Gakuba et al., 2018).

Dexmedetomidine (DEXM) is another α_2 -adrenergic agonist widely used for anesthesia or sedation in clinics and has also been shown to enhance glymphatic function in animal research (Lilius et al., 2019a). Recently, Ozturk et al. reported that DEXM supplemented with low-dose isoflurane increased glymphatic transport compared to that of isoflurane only (Ozturk et al., 2021). DEXM has also been shown to increase the delivery of intrathecally administered drugs, such as oxycodone and naloxone, by increasing glymphatic flow (Lilius et al., 2019b).

Propofol is a widely used intravenous anesthetic in humans with anti-inflammatory (Ren et al., 2014) and neuroprotective

properties (Miller et al., 2018). Recent studies have found that propofol can increase glymphatic activity (Gakuba et al., 2018). Pentobarbital has also been shown to significantly increase glymphatic function in animal models (Hablitz et al., 2019), although benzodiazepines have mainly been replaced by human anesthesia. Other anesthetics used in animal anesthesia such as α -chloralose and avertin inhibit CSF tracer influx in the glymphatic pathway (Hablitz et al., 2019). Therefore, glymphatic function is affected by the dose, type, and combination of anesthetics. Although the exact mechanisms underlying these preclinical experiments remain unclear, they may provide valuable information for studying the functional effects of the glymphatic system during clinical anesthesia (Table 1).

Interestingly, these animal studies on glymphatic dysfunction agree with those of clinical studies on PND. Anesthetics are a risk factor for postoperative cognitive decline or neuropathological changes (Schenning et al., 2016). It has been confirmed that inhaled anesthetics, such as isoflurane or sevoflurane, have more chance of causing postoperative cognitive dysfunction (Ologunde and Ma, 2011; Hu et al., 2014), whereas some intravenous anesthetics, such as DEXM and propofol, cause slight harm or even improve perioperative cognitive function (Miller et al., 2018). Anesthetics that have a significant inhibitory effect on the glymphatic system are more likely to cause perioperative cognitive impairment, whereas some anesthetics that enhance glymphatic function are less likely to cause cognitive impairment. The effects of ketamine on the glymphatic system and postoperative cognition have been controversial (Morrison et al., 2018). DEXM enhances glymphatic function and has also been shown to prevent postoperative delirium and cognitive dysfunction caused by inhaled anesthetics in elderly patients (Zhang et al., 2018). Therefore, anesthetics may affect postoperative brain function by affecting the glymphatic function. Currently, the mechanisms underlying the effects of anesthetics on glymphatic function remain unclear. Further clinical research is necessary to determine the optimal variety, dose, and combination of anesthetics and drugs to reduce glymphatic function impairment during anesthesia.

THE UNDERLYING MECHANISM OF ALTERED GLYMPHATIC FUNCTION DURING INTRAOPERATIVE ANESTHESIA MANAGEMENT

The mechanisms that regulate glymphatic function under normal physiological conditions include arterial pulsatility, respiration, and neural activity. However, these mechanisms may be disrupted during anesthesia, including general anesthesia and spinal anesthesia (Scott, 2018). During clinical anesthesia, the heart rate, blood pressure, and arterial pulsation are significantly changed, and spontaneous negative pressure respiration is replaced with positive pressure ventilation. The arousal mechanism is also significantly suppressed during general anesthesia (Schiff, 2020). Anesthesia-induced pathological processes, such as

the breakdown of glymphatic CSF-ISF exchange, have been linked to disease initiation and progression. As a result, the impact of anesthesia on glymphatic function needs to be considered.

Cardiac Pulsation Mechanism Change

The sympathetic, stellate ganglion, and subcutaneous sympathetic nerves are all suppressed by most anesthesia types, including general, neuraxial, epidural, and lumbar anesthesia. Anesthesia suppresses blood pressure and heart rate by causing dose-dependent vasodilation or vasoplegia. Hypotension occurs due to anesthesia-induced reduction in heart rate, vasodilation, reduced afterload, and circulating volume. Furthermore, a lower heart rate has a strong positive association with glymphatic influx (Hablitz et al., 2019). Currently, there is no research on the relationship between hypotension and glymphatic activity. Mestre et al. found that the jerky and irregular reversal of microparticle flow caused by increased arteriole stiffness and pulsation amplitude in distal vessels during hypertension reduced perivascular pump dynamics and decreased net CSF flow and waste clearance in the perivascular space (Mestre et al., 2018b). Additionally, CSF production is related to sympathetic or adrenergic receptor activity. Increased sympathetic tone inhibits carbonic anhydrase associated with the choroidal epithelium and likely reduces CSF production and glymphatic circulation (Damkier et al., 2013; Benveniste et al., 2017). Recently, Liu et al. also reported that CSF production was increased by a combination of α_1 -, α_2 -, and non-selective β -adrenergic receptor antagonists, as well as some anesthetics, including isoflurane, ketamine, and xylazine (Liu G. et al., 2020).

Perioperative hemodynamic optimization is crucial for improving outcomes after surgery and anesthesia. Anesthesia management often includes the administration of catecholamine vasoactive drugs such as norepinephrine, dopamine, and phenylephrine to maintain stable hemodynamics throughout surgery (Foss and Kehlet, 2019). Norepinephrine, a central regulator, also inhibits glymphatic transport (Jessen et al., 2015). However, Iliff et al. demonstrated that dobutamine increased the pulsatility of the penetrating artery by 60% and enhanced the glymphatic influx (Iliff et al., 2013). Furthermore, catecholamines can modulate astrocyte function and signaling to regulate the glymphatic system (Fuxe et al., 2015). The astrocytic dopamine and adrenergic receptor subtypes are significant drug targets in neurological and psychiatric diseases (Fuxe et al., 2015). Norepinephrine and dobutamine also work together to affect glymphatic function by acting on fluid availability and convective fluxes through various mechanisms. Therefore, catecholamines may be the primary regulators of solute clearance from the brain.

Theoretically, any changes in cardiovascular parameters induced by anesthesia can impair the cardiac pulsation mechanism, resulting in waste accumulation. Hence, understanding the processes contributing to pulsatility and how these components drive waste clearance by the glymphatic system continues to evolve. Further research is needed to determine the most appropriate management of glymphatic function during clinical anesthesia.

TABLE 1 | The effect of verified factors on glymphatic system.

Classification	Perioperative factors	Glymphatic function	Classification	Perioperative factors	Glymphatic function
Anesthetic factors	Neurotoxicity	Impair (Rangroo Thrane et al., 2013)	Cardiovascular factors	Acute hypertension	Impair (Mestre et al., 2018b)
	Isoflurane (2–2.5%)	Inhibit (2–2.5%) (Gakuba et al., 2018)		Chronic hypertension	Impair (Mortensen et al., 2019; Koundal et al., 2020)
	Sevoflurane	Enhance (2.5%) (Gao et al., 2019)		Heart rate	Negative correlation (Hablitz et al., 2019)
	Dexmedetomidine	Enhance (20 mg.kg ⁻¹ ; ip) (Hablitz et al., 2019; Lilius et al., 2019a)	Surgical factors	Dobutamine	Enhance (Iliff et al., 2013; Yamada et al., 2013)
	Xylazine	Enhance (Hablitz et al., 2019)		Norepinephrine	Inhibit (Jessen et al., 2015)
	Ketamine	Inhibit (150 mg.kg ⁻¹) (with xylazine, 10 mg.kg ⁻¹) (Gakuba et al., 2018)		Neuroinflammation	Impair (Yu et al., 2019)
		Enhance (100 mg.kg ⁻¹) (with xylazine 20 mg.kg ⁻¹) (Hablitz et al., 2019)		Position	lateral > supine > prone (Lee et al., 2015)
	Propofol	Enhance (Gakuba et al., 2018)	Individual factors	Sleep disturb	Impair (Nedergaard and Goldman, 2020).
	Pentobarbital	Enhance (60 mg.kg ⁻¹ ; ip) (Hablitz et al., 2019)		BBB damage	Impair (Meng et al., 2019; Yu et al., 2019)
	α-chloralose	Inhibit (80 mg.kg ⁻¹ ; ip) (xylazine Hablitz et al., 2019)		Postoperative Pain	Impair (Chouchou et al., 2014)
Arouse stage	Avertin	Inhibit (120 mg.kg ⁻¹ ; ip) (Hablitz et al., 2019)		Perioperative Stress	Impair (Wei et al., 2019)
	Wake	Inhibit (Xie L. et al., 2013; Eide and Ringstad, 2019)	Other factors	Preclinical stage of AD	Impair (Masters et al., 2015)
	Sleep	Enhance (Xie L. et al., 2013; Alawieh et al., 2018)		Neurovascular diseases	Impair (Riba-Llena et al., 2018)
	EEG	Beta power inhibit (Hablitz et al., 2019)		Aging	Impair (Iliff et al., 2013)
Respiration		Delta power enhance (Hablitz et al., 2019)		Apo E gene	Impair (Mentis et al., 2020)
	Free breathing	N/A	Other factors	AQP4 gene	Impair (Hubbard et al., 2018).
	Deep inhalation	Enhanced (Dreha-Kulaczewski et al., 2015)		Chronic hypertension	Impair (Mestre et al., 2018b; Riba-Llena et al., 2018)
	Mechanical ventilation	N/A		Diabetes	Impair (Zhang et al., 2014; Jiang et al., 2017)
				Intracranial pressure	Increase will impair (Dreha-Kulaczewski et al., 2015)

Respiration Changes

During experimental testing in mouse or rat models described above, spontaneous respiration was inhibited but still preserved in research involving the glymphatic system. Spontaneous respiration in anesthetized animals is very different from that in controlled or assisted respiration under clinical anesthesia. During clinical anesthesia, positive-pressure mechanical ventilation is often used to replace spontaneous respiration. Mechanical ventilation differs from spontaneous breathing in terms of physiology, and its effect on glymphatic system transport in the brain is currently unknown. Ventilators are

inspiratory assist devices that deliver tidal breath under positive pressure. Positive pressure ventilation, including positive end-expiratory pressure, may increase intrathoracic pressure and reduce venous return, subsequently causing increased ICP (Chen et al., 2018). Positive pressure ventilation also reduces cerebral perfusion pressure due to increased ICP and decreased mean arterial pressure (Chen et al., 2018). It has been hypothesized that ventilation impairs respiration, causing waste accumulation in the brain.

Although the administration of local anesthetics through the spinal or epidural canal can provide anesthesia without the need

for mechanical ventilation, there are still additional effects on the sympathetic nervous system, parasympathetic nervous system, and motor control, resulting in respiratory impairment (Wink et al., 2016). Theoretically, these changes impair the function of the glymphatic system. Some respiratory parameters, such as tidal volume, frequency, and airway pressure, have not been studied in relation to glymphatic activity.

Preclinical and clinical evidence suggests that mechanical ventilation induces adverse neurocognitive effects (Wang et al., 2015). It is estimated that 70–100% of critically ill patients have cognitive impairment after mechanical ventilation, and 20% of patients still have cognitive impairment 5 years later (Herridge et al., 2016; Bilotta et al., 2019). In patients with COVID-19, delirium/neurological disorders are clearly correlated with prolonged mechanical ventilation (Alonso-Lana et al., 2020; Helms et al., 2020). However, the mechanism by which mechanical ventilation affects the incidence of postoperative cognitive disorder remains unclear. The discovery of this novel glymphatic pathway may provide clues for future research. Evaluating perioperative glymphatic function might provide new insights for understanding the pathophysiological mechanisms underlying cognitive decline during clinical anesthesia and surgery.

Arousal Regulation Mechanism Altered by Anesthesia

Some studies have demonstrated that arousal state correlates with the glymphatic system (Hablitz et al., 2019), while the arousal state also reflects “depth of anesthesia” in clinical settings (Shalhaf et al., 2015). The arousal state is closely related to hemodynamic rhythms such as blood pressure, heart rate, and blood oxygenation level-dependent signals, which affect glymphatic function (Fultz et al., 2019). The depth of anesthesia can affect not only the hemodynamics but also the arousal mechanism. Therefore, there may be a relationship between anesthesia depth and glymphatic function (Shalhaf et al., 2015). General anesthesia is characterized by the suppression of central arousal controlled by the locus coeruleus and noradrenergic tone (Brown et al., 2010). Norepinephrine release is reduced when anesthesia suppresses sympathetic nerve activity. It has been confirmed that noradrenergic blockade or reduction produces the same changes in ISF volume fraction in the glymphatic system (Xie L. et al., 2013).

However, accurate estimation of the depth of anesthesia remains an important issue in clinical or animal anesthesia. Non-invasive EEG or evoked potential indices, such as the bispectral index, narcotrend index, state entropy, and response entropy, for monitoring anesthesia depth are widely used in clinical research. High EEG delta power is positively associated with glymphatic activity, while beta power is negatively correlated in animal models (Hablitz et al., 2019). When bispectral index-guided deep anesthesia is used in geriatric anesthesia, postoperative delirium and cognitive decline in elderly surgical patients are reduced (Deschamps et al., 2019; Miao et al., 2019). Therefore, intraoperative indicators for monitoring anesthesia depth are clinically significant for research about the relationship between

glymphatic function and PND. Insufficient anesthesia can cause physiological and psychological injuries, whereas overdose anesthesia may induce hemodynamic disturbances. Therefore, monitoring the depth of anesthesia may also prevent impairment of the glymphatic system and cognition due to inadequate or excessive anesthesia.

Astrocytes and AQP4 Function Changes

The use of *in vivo* two-photon imaging demonstrated that calcium signaling in astrocytes is associated with the selective transport of small lipophilic molecules and rapid ISF movement throughout the glymphatic system (Rangroo Thrane et al., 2013). A recent study found that glymphatic flow is regulated by fluid shear stress produced by perivascular CSF or ISF dynamics, which are capable of mechanically opening NMDA receptors and producing increased Ca^{2+} currents in cultured astrocytes (Maneshi et al., 2017). General anesthetics, especially ketamine, inhibit NMDA receptors and reduce Ca^{2+} signaling (Duman et al., 2019), likely to regulate glymphatic function through this mechanism. The anesthetic sevoflurane induces changes in astrocyte morphogenesis by downregulating ezrin, an actin-binding membrane-bound protein, in addition to disrupting astrocyte Ca^{2+} currents acutely and chronically (Zhou et al., 2019). Moreover, brain surgery such as cisterna magna puncture can cause spontaneous frequent but asynchronous astrocytic Ca^{2+} signaling and impair glymphatic lipid transport, consequently increasing intracellular lipid accumulation (Rangroo Thrane et al., 2013).

AQP4 colocalizes the inwardly rectifying potassium channel Kir4.1 and contributes to the coupled influx of water and K^{+} after neuronal activity (Jo et al., 2015). Accordingly, some anesthetics can impair AQP4 function by blocking potassium channels (Ou et al., 2020), leading to glymphatic function changes. In recent studies, 2.5% sevoflurane has also been shown to increase glymphatic function by upregulating AQP4 expression in astrocytes (Gao et al., 2019). Changes in AQP4 expression and depolarization are functionally relevant to the glymphatic system and cognition (Hubbard et al., 2018).

GLYMPHATIC FUNCTION AFFECTED BY SURGERY

Persistent systemic inflammation caused by surgery can disrupt the integrity of the BBB (Vacas et al., 2014; Hughes et al., 2016; Bi et al., 2017; Wang et al., 2017; Ni et al., 2018; Verheggen et al., 2018). The BBB is a complex functional and anatomical system that prevents the entry of neurotoxic plasma elements, blood cells, and pathogens into the brain. It is comprised of endothelial cells, perivascular microglia, pericytes, neurons, and astrocytic end-feet (Sweeney et al., 2019). Tight junction proteins located between adjacent endothelial cells include claudins, tricellulin, occludin, and zonula occludens (Sweeney et al., 2019). Acute endothelial dysfunction caused by surgically induced systemic inflammation may lead to BBB disorders (Plog et al., 2015; Ekeloef et al., 2017). Inflammation after laparotomy can decrease the levels of tight junction

proteins such as claudin, β -catenin, occludins, and ZO-1, all of which can contribute to tight junctions and BBB permeability (Yang et al., 2017). Moreover, astrocytes are easily activated by postoperative inflammatory cytokines such as TNF- α , IL-1 β , IL-6, complement C3, and high mobility group box-1, which can subsequently damage the integrity of the BBB, especially in the hippocampus (He et al., 2012; Jin et al., 2014; Xiong et al., 2018). Activated astrocytes can also encourage tissue proliferation, resulting in scar-like perivascular barriers (Voskuhl et al., 2009). The endothelial cell membrane and the BBB close junctions comprise the vascular components of the glymphatic system, while the avascular components consist of astrocyte end-feet integrated with glia-line boundaries between the neuropil and axon tract regions (Plog and Nedergaard, 2018). Therefore, systemic inflammation influences the function of the glymphatic system based on the evidence that the structure and function of the glymphatic system are closely related to that of the BBB (Verheggen et al., 2018). Notably, a recent study found that lipopolysaccharide-induced systemic inflammation reduced perivascular CSF tracer flow and penetration into the parenchyma (Manouchehrian et al., 2021). These observations may be beneficial for our understanding of the role of systemic inflammation in glymphatic clearance.

How the Surgical Position Influences Glymphatic Transport

Researchers have studied the effects of body position on glymphatic function and A β clearance using optical imaging, CSF radiotracers, and fluorescent tracers in the brains of anesthetized rodents. According to previous studies, glymphatic transport is most effective in the lateral position, inferior in the supine position, and least efficient in the prone position (Lee et al., 2015). Thus, it is clear that postural or gravitational factors also exert regulatory control over the glymphatic pathway. Thus, we speculate that in postures with the head down, minor changes in CSF pressure, such as ICP and hydrostatic pressure, increase due to gravity; tissue pressure or vascular function could alter the shape of perivascular spaces and accelerate glymphatic drainage problems, impairing cognitive function.

Although PND was initially described as a complication after cardiac surgery, many recent studies have found that some patients undergoing non-cardiac surgeries, such as orthopedic surgery, also suffered cognitive decline (Evered and Silbert, 2018). Due to the various types of surgery, surgical posture may be an essential but overlooked factor in these studies. Some prone surgeries, particularly spine surgery, may cause postoperative cognitive impairment (Kim et al., 2016; Ezhevskaya et al., 2019). Furthermore, Trendelenburg, a supine position used in laparoscopic surgery in which the patient's head is lower than the feet, can result in postoperative cognitive decline. This position could increase intra-abdominal pressure, reduce cerebral venous outflow, and impair lymphatic function, similar to positive pressure ventilation or positive end-expiratory pressure (Maerz et al., 2017). Some studies have even demonstrated that sleeping in the supine position has been shown to reduce cognitive

function in healthy volunteers, while sleeping in the upright position did not affect the participants (Muehlhan et al., 2014). According to studies on posture and glymphatic function, the surgical position can worsen the cognitive impairment, especially in prolonged surgeries, including cardiac surgery. Understanding the effects of different body postures on the glymphatic clearance pathway may be necessary for interpreting the procedure of PND.

Perioperative Sleep Disorder Leads to Glymphatic Dysfunction

The clearance of the glymphatic system is closely related to the sleep-wake cycle, with sleep promoting faster metabolite removal (Xie L. et al., 2013). However, perioperative sleep is frequently disrupted by many factors, including postoperative pain, environmental and surgical stress, anesthesia, and other factors that lead to discomfort (Whitlock et al., 2017; Su and Wang, 2018). Postoperative sleep disruptions manifest as sleep fragmentation and reduced slow-wave and REM sleep durations (Chouchou et al., 2014). In particular, REM sleep disorder may significantly negatively affect postoperative cognition (Lazic et al., 2017). Subsequently, the glymphatic system is impaired and degrades with sleep disturbances (Nedergaard and Goldman, 2020). After a night of sleep deprivation, protein A β levels in the thalamus and medial temporal structures are also elevated in healthy individuals (Shokri-Kojori et al., 2018). Moreover, sleep disruption increases the neuronal activity and generates more waste products, including lactate, which are exported via glymphatic fluid transport (Lundgaard et al., 2017).

Postoperative pain remains a significant health care issue that disturbs sleep in the postoperative period, and sleep disturbances may, in turn, exacerbate postoperative pain (Chouchou et al., 2014). Anesthesia is substantially different from natural sleep, sometimes interfering with the circadian rhythm and disrupting the sleep cycle, subsequently affecting glymphatic clearance (Lazic et al., 2017). Additionally, surgical and environmental stress appears to be a significant cause of sleep disruptions during peri-operation (Xu et al., 2016), and chronic stress via glucocorticoid signaling impairs AQP4-mediated glymphatic transport (Wei et al., 2019). This evidence suggests that cognitive decline in patients with sleep disorders may be caused by decreased waste removal and increased waste accumulation in the brain due to glymphatic dysfunction.

Currently, in animal models of glymphatic function, sleep is simulated by anesthesia or sleep deprivation. Owing to technical limitations, it is difficult to perform such experiments on naturally sleeping mice (Xie L. et al., 2013). According to some studies, the effects of sleep and general anesthesia on the glymphatic system are equivalent (Xie L. et al., 2013; Benveniste et al., 2019a). However, anesthesia and natural sleep outcomes remain markedly different; anesthesia can affect hemodynamics, interfere with the biological clock, and disrupt the sleep cycle (Lazic et al., 2017). Thus, the impact of anesthesia on the glymphatic system needs to be investigated further.

WHY SOME PATIENTS ARE SUSCEPTIBLE TO PND

Notably, the adverse cerebral effects of anesthesia were primarily reported in the elderly (Bedford, 1955; Berger et al., 2015). It is not yet fully understood why elderly patients are highly susceptible to PND, while young people who undergo surgery rarely develop it (Hu et al., 2020). Researchers have discovered some PND-susceptible patients with preoperative disease, such as preclinical AD (Xu et al., 2014), hyperglycemia (Zhang et al., 2014), cerebral microemboli (Wang et al., 2020), and other underlying neurovascular diseases or neurodegeneration (Wang et al., 2020). These preoperative diseases may share common underlying mechanisms and contribute to neurocognitive dysfunction or dementia. With advances in imaging technology, it has been found that these diseases have glymphatic impairment (Gaberel et al., 2014; Kress et al., 2014; Kyrtos and Baras, 2015; Jiang et al., 2017; Hadjihambi et al., 2019). A theory states that susceptible patients with PND and aging have preexisting glymphatic dysfunction but progresses slowly without significant cognitive impairment (Bolte et al., 2020). Perioperative risk factors exacerbate the preexisting glymphatic damage beyond the compensatory limit, resulting in significant cognitive changes in these individuals. AD is a progressive neurodegenerative disorder characterized by gradual cognitive decline, which is characteristic of extracellular A β deposition and intracellular accumulation of hyperphosphorylated tau (Long and Holtzman, 2019). Patients in the preclinical stage of AD, particularly those with APOE4 genetic risk factors (Mentis et al., 2020), already have A β accumulation and impaired glymphatic clearance, but no clinically detectable symptoms of cognitive impairment (Masters et al., 2015). Moreover, aging is associated with degenerative changes in the brain's structure and function, such as reduced cerebrovascular pulsatility, depolarized or downregulated AQP4, and sleep deprivation, which impair glymphatic function (Zeppenfeld et al., 2017). Glymphatic dysfunction may result in the deposition of toxic solutes (including amyloid) in the aging brain, which exacerbates glymphatic dysfunction and creates a vicious cycle (Iliff et al., 2013). Vascular dementia is caused by many neurovascular diseases, including hypertension, ischemic stroke, traumatic brain injury, and microinfarcts, which significantly impair arterial CSF pulsation and glymphatic clearance (Riba-Llena et al., 2018). Diabetes can also impair cardiac pulsation due to pathological changes in the vascular endothelium (Jiang et al., 2017).

Due to technical limitations, few previous studies have used non-invasive methods to investigate glymphatic function in humans, particularly hospitalized surgical patients susceptible to PND. Recent advances in imaging, such as real-time MRI (Ding et al., 2018), ultra-fast MRI (Kiviniemi et al., 2016), and gadolinium-based contrast agents enhanced MRI (Ringstad et al., 2018; Deike-Hofmann et al., 2019), will provide a sensitive and non-invasive tool for assessing glymphatic function in clinical trials. Notably, it is essential to develop intraoperative monitoring tools for glymphatic function. A new method of measuring human glymphatic function that uses

near-infrared spectroscopy has been developed for long-term monitoring of brain function, which is highly compatible with perioperative monitoring (Myllylä et al., 2018). These non-invasive methods may help translate glymphatic measurements from the laboratory to the clinic (Jiang, 2019). Advances in technology will yield new and valuable information about the glymphatic system, providing a quantitative map for the diagnosis, monitoring, and prognosis of neurological disorders, including PND.

CONCLUSION AND PERSPECTIVE

Recent discoveries concerning the glymphatic system anatomy and mechanisms have contributed to a better understanding of CSF circulation and waste removal. Furthermore, the glymphatic system has an immune function and may affect the onset and progression of PND, either directly or indirectly. Because clinical research on perioperative glymphatic changes is still lacking, scientists and clinicians, particularly gerontologists, neurologists, psychiatrists, anesthesiologists, and surgeons, can collaborate to uncover the etiological mechanisms of perioperative glymphatic changes. The glymphatic concept may provide a strategic breakthrough in rethinking, treating, and, most importantly, preventing PND.

Human studies exploring the physiological and pathological mechanisms between the glymphatic system and brain cognition over time have greatly improved our knowledge; however, some key issues still need to be addressed by future research. First, the definition of normal or healthy glymphatic function may be one of the biggest challenges. A simpler and quicker intraoperative monitoring method remains to be developed. A better description of glymphatic dynamics will help demarcate "normal" and "abnormal" glymphatic function in humans and identify preventive strategies for PND. Second, perioperative factors such as anesthetic type, intraoperative hemodynamics, surgery position, surgical inflammation, and previous health status may confound research on the glymphatic system and PND. Cross-sectional clinical studies have demonstrated specific changes in the glymphatic transport of patients susceptible to neurological diseases, which is helpful in predicting the incidence of PND. Third, more clinical studies should be conducted instead of purely animal studies. Although association analysis may provide critical information for cause-effect deductions, correlation does not necessarily mean causation. Fourth, basic research targeting the perioperative glymphatic system in PND merits optimization; specifically, appropriate model systems should be carefully selected. Finally, the translation of basic research results into clinically relevant effects in humans should be expedited. Most data on the role of the glymphatic system are based on animal studies. Preclinical animal studies frequently result in unexpected failures during clinical transition due to unidentified reasons. Alternatively, future therapeutic interventions will likely be based on individual factors due to significant differences in anatomical variation and disease complexity among human populations.

AUTHOR CONTRIBUTIONS

XR, GZ, LL, and CL contributed significantly to conception of the review and edited the review. XR undertook the literature, drafted, and corrected most

of the manuscript and tables. SL, KL, and HL assisted with part of the pictures and manuscript. CL contributed to the sections of glymphatic system and anaesthetics. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Neuroimaging Markers of Cerebral Small Vessel Disease on Hemorrhagic Transformation and Functional Outcome After Intravenous Thrombolysis in Patients With Acute Ischemic Stroke: A Systematic Review and Meta-Analysis

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Objective: The aim of this study was to perform a systematic review and meta-analysis to assess whether cerebral small vessel disease (CSVD) on neuroimaging of patients with acute ischemic stroke (AIS) treated with intravenous thrombolysis (IVT) is associated with an increased risk of hemorrhagic transformation (HT), symptomatic intracranial hemorrhage (sICH), and poor functional outcome (PFO).

Methods: A thorough search of several databases was carried out to identify relevant studies up to December 2020. We included studies of patients with AIS and neuroimaging markers of CSVD treated with IVT. The primary outcome was HT, and the secondary outcomes were sICH and 3-month PFO. The quality of the studies involved was evaluated using the Newcastle–Ottawa Scale (NOS). The meta-analysis with the fixed effects model was performed.

Results: Twenty-four eligible studies ($n = 9,419$) were pooled in the meta-analysis. All included studies were regarded as high quality with the NOS scores of at least 6 points. The meta-analysis revealed associations between the presence of CSVD and HT, sICH, and the 3-month PFO after IVT. Compared with no CSVD, the presence of CSVD was associated with an increased risk of HT (OR: 1.81, 95% CI: 1.52–2.16), sICH (OR: 2.42, 95% CI: 1.76–3.33), and 3-month PFO (OR: 2.15, 95% CI: 1.89–2.44). For patients with AIS complicated with CSVD, compared with a CSVD score of 0–1, a CSVD score of 2–4 was associated with an increased risk of HT (OR: 3.10, 95% CI: 1.67–5.77), sICH (OR: 2.86, 95% CI: 1.26–6.49), and 3-month PFO (OR: 4.58, 95% CI: 2.97–7.06).

Conclusion: Patients with AIS complicated with neuroimaging markers of CSVD are at increased risk of HT and 3-month PFO after IVT. However, it is still necessary to clarify the exact role of CSVD in the occurrence, development, and prognosis of AIS.

Systematic Review Registration: www.ClinicalTrials.gov, identifier CRD42021233900.

Keywords: cerebral small vessel disease, acute ischemic stroke, intravenous thrombolysis, hemorrhagic transformation, neuroimaging markers

INTRODUCTION

Stroke is the second leading cause of death worldwide, leading to death in 5.5 million people and affecting 13.7 million each year. Ischemic stroke accounts for 70% of all patients with stroke. The 2016 Global Burden of Disease Study data that were published in 2019 suggest that one in four adults is reported to be at risk of having a stroke in their lifetime (GBD 2016 Stroke Collaborators, 2019; Lindsay et al., 2019).

Cerebral small vessel disease (CSVD) is a widespread cerebrovascular disease with specific neuroimaging characteristics (Chen et al., 2019). With the development of neuroimaging technology, the brain imaging of more patients with acute ischemic stroke (AIS) has detected the neuroimaging markers of CSVD such as cerebral microbleed (CMB), white matter hyperintensity (WMH), lacunar infarction (LI), and enlarged perivascular space (EPVS) (Curtze et al., 2016; Chen et al., 2019). CSVD accounts for 25% of cases of AIS, and it affects cognitive function, gait disturbance, swallowing, and other functions (Pantoni, 2010).

Studies have shown that CSVD may be a risk factor for intracranial hemorrhage, but none of these studies have a clear determinism (Emberson et al., 2014; Jickling et al., 2014; Pang et al., 2019). Hemorrhagic transformation (HT) occurs in 10–40% of patients with ischemic stroke and is a major complication of intravenous thrombolysis (IVT) (Terruso et al., 2009; Beslow et al., 2011; Jickling et al., 2014). HT can be divided into symptomatic and asymptomatic according to the deterioration of neurological function, both of which worsen the prognosis of stroke, especially in cognitive and neurological functions (Dzialowski et al., 2007; Park et al., 2012). Most patients with AIS complicated with CSVD have no obvious clinical symptoms at the initial stage, which are easy to be ignored by doctors. Therefore, for patients with AIS complicated with CSVD, HT, symptomatic intracranial hemorrhage (sICH), and poor functional outcome (PFO) after IVT have gradually attracted the attention of medical researchers (Liu X.Y. et al., 2019). Different subtypes of CSVD may have different effects on HT and clinical prognosis after IVT in patients with AIS, and different subtypes of CSVD often coexist in the same patients with AIS.

Several recent studies have explored the relationship between these neuroimaging markers of CSVD and clinical outcomes after IVT in patients with AIS (Drelon et al., 2020; Liu X. et al., 2020). Previous meta-analyses have investigated the increased risk of CSVD for HT and PFO in patients with AIS (Charidimou et al., 2016; Tsivgoulis et al., 2016; Wang et al., 2017). However,

these studies are mainly limited to a particular subtype of CSVD, and some patients receive endovascular treatment. Whether the existence of neuroimaging markers of CSVD affects the HT, sICH, and PFO of patients with AIS after IVT is still a controversial issue. Therefore, we performed a systematic review and meta-analysis to evaluate whether CSVD on neuroimaging of patients with AIS treated with IVT is associated with an increased risk of HT, sICH, and PFO.

METHODS

This systematic review and meta-analysis was performed in accordance with the Preferred Reporting Items for Systematic reviews and Meta-Analyses (PRISMA) guidelines.

Search Strategy

We systematically searched the MEDLINE, Cochrane Library, Embase, CNKI, VIP, and WANFANG databases from inception to December 2020 to find relevant studies. The articles were not restricted based on the language of publication. The details of the search strategy are presented in **Supplementary Appendix 1**. Two authors (i.e., WYQ and YXT) scanned the titles and abstracts to find the articles that were most relevant to this study; then, the full texts of the relevant articles were examined, and the final decision on inclusion was made by consensus.

Selection Criteria

Types of Studies

We included cohort studies (i.e., prospective and retrospective) in which patients with AIS or suspected AIS were treated with IVT. The relationship between neuroimaging markers of CSVD and clinical outcomes was assessed. All eligible trials were published in full text without language restrictions.

Types of Participants

We considered trials that included IVT-treated patients with AIS or patients who were treated with IVT for suspected ischemic stroke. The diagnosis of AIS meets the WHO diagnostic criteria (No authors listed, 1989), and participants were confirmed by CT or MRI. The diagnosis points of AIS are as follows: (1) acute onset, (2) focal neurological deficit, (3) responsible lesions appearing on imaging or symptoms lasting more than 24 h, and (4) excluding cerebral hemorrhage by brain CT/MRI. Trials involving patients treated with endovascular therapy were excluded. Cerebral imaging had to be performed for the

visualization of CSVD. CMB, WMH, LI, and EPVS can all be detected on MRI (Chen et al., 2019). CMB is defined as a small, round, or oval hypointense lesion that can be shown on T2*-weighted gradient recalled echo (GRE) and susceptibility-weighted imaging (SWI) (Wardlaw et al., 2013). WMH is an imaging description of white matter demyelination, it is hyperintense on T2-weighted image (T2WI) and fluid-attenuated inversion recovery (FLAIR) sequences on MRI, and it can also be shown on CT, but the range of lesions shown on CT may not be ideal (Wardlaw et al., 2013; van Leijssen et al., 2018). LI is a hyperintense area with the largest lesion diameter less than 20 mm on the axial plane of the FLAIR sequence (Wardlaw et al., 2013). Generally, the diameter of the perivascular space is less than 2 mm, the EPVS can extend to a diameter of 2–4 mm, and it can be detected on MRI (Chen et al., 2019). To expand the scope of the study, WMH detected on CT is also included (Charidimou et al., 2016).

Primary Outcome Assessments

The primary outcome for the systematic review was the occurrence of HT. There are many types of HT diagnostic criteria, such as the European Cooperative Acute Stroke Studies (ECASS) (Fiorelli et al., 1999; Larrue et al., 2001; Hacke et al., 2008) and the National Institute of Neurological Disorders and Stroke (NINDS) (No authors listed, 1997). The diagnosis points of ECASS are as follows: (1) hemorrhagic infarction (HI): there was a small spot-like hemorrhage along the edge of the infarct and sheet-like non-massive bleeding or multiple fused spot-like hemorrhages in the infarct area and (2) parenchymal hematoma (PH): hematoma, i.e., bleeding with slight or obvious space-occupying effect or bleeding away from the infarct. The main points of the diagnosis of NINDS are as follows: (1) HI: different low-density/high-density foci with punctate or blurred borders in the acute infarct can be tested on CT and (2) PH: typical homogeneous high-density lesions with clear boundaries, with or without cerebral edema or space-occupying effects, can be tested on CT.

Secondary Outcome Assessments

The secondary outcomes included the occurrence of sICH, and sICH was defined by the scores on the ECASS (Fiorelli et al., 1999; Larrue et al., 2001; Hacke et al., 2008), NINDS (No authors listed, 1997), and Safe Implementation of Thrombolysis in Stroke-Monitoring Study (SITS-MOST) (Rha et al., 2014). The 3-month PFO was defined as a Modified Rankin Scale (mRS) score > 2. The relationships among HT, sICH, 3-month PFO, and the total burden of CSVD (by using the Total Burden Rating Scale of CSVD) were also examined. The diagnosis points of ECASS are as follows: bleeding is seen on CT, and the increase in the National Institute of Health Stroke Scale (NIHSS) score is ≥ 4 points. The diagnosis points of NINDS are as follows: bleeding on CT, accompanied by neurological decline. The main points of diagnosis of SITS-MOST are as follows: the infarct area or remote area PH is seen on CT, the NIHSS score increased by 4 points or more compared with the minimum level of 24 h after admission, and bleeding caused death. The mRS can assess the complete independent living ability of patients with stroke, where a score of 0 means asymptomatic and the higher scores indicate

a worse prognosis (i.e., a score of 6 indicates death). The concept of the total burden of CSVD was proposed by Staals et al. (2014), who developed a scale to quantitatively evaluate the cumulative effect of CSVD throughout the whole brain. The total score of the scale ranges from 0 to 4 points, and the higher the score, the more serious the CSVD.

Data Extraction

We used a preset electronic collection form to extract the basic characteristics of the studies, such as study design, first author, year of publication, country, sample size, age, sex, imaging method, type of neuroimaging markers of CSVD (i.e., CMB, WMH, LI, and EPVS), the definition of HT and sICH, the total burden of CSVD, and clinical outcome. The number of patients with neuroimaging markers of CSVD and the number of outcome events in each study were extracted, and conversion was performed when studies were reported as a percentage. When a study reported the relationship between different subtypes of CSVD and outcomes, the data were extracted separately.

Quality Assessment

Two reviewers independently assessed each study for quality using the Newcastle–Ottawa Scale (NOS) (Wells et al., 2013), which mainly contains three domains as follows: (1) selection, (2) comparability, and (3) outcome. The selection domain included the representativeness of the exposed cohort, selection of non-exposed cohort, ascertainment of exposure, and demonstration that the outcome of interest was not present at the beginning of the study. The comparability domain included the comparability of cohorts based on the design or analysis. Outcomes included the assessment, long enough follow-up for outcomes to occur, and adequacy of follow-up of cohorts. The maximum score of the NOS was 9 points, and a score ≥ 6 indicated a high-quality study. Disagreements in the studies were resolved by senior researchers. The details of the quality assessment can be found in **Supplementary Appendix 2**.

Statistical Analyses

We used RevMan software version 5.3 and Stata MP software version 14.0 for all the statistical analyses. The meta-analysis was used to calculate the combined odds ratio (OR) and the corresponding 95% CI to quantify the strength of the association between the presence and severity of CSVD and HT, sICH, and 3-month PFO after IVT. The total burden of CSVD was divided into two groups, namely, 0–1 and 2–4 points. We tested for heterogeneity between trial results using the I^2 statistic (Higgins et al., 2003) (when $P \leq 0.1$ and $I^2 \leq 25\%$ indicated low heterogeneity, $25\% < I^2 \leq 50\%$ indicated moderate heterogeneity, and $I^2 > 50\%$ indicated significant heterogeneity). The fixed effects model was used when heterogeneity between studies was not detected; otherwise, the random effects model was used.

To explore the factors associated with heterogeneity, the subgroup analysis according to different types of CSVD (i.e., CMB, WMH, LI, and EPVS) was performed to explore the impact of different types on the outcome. The sensitivity analysis was used to analyze whether the conclusion was stable, and we

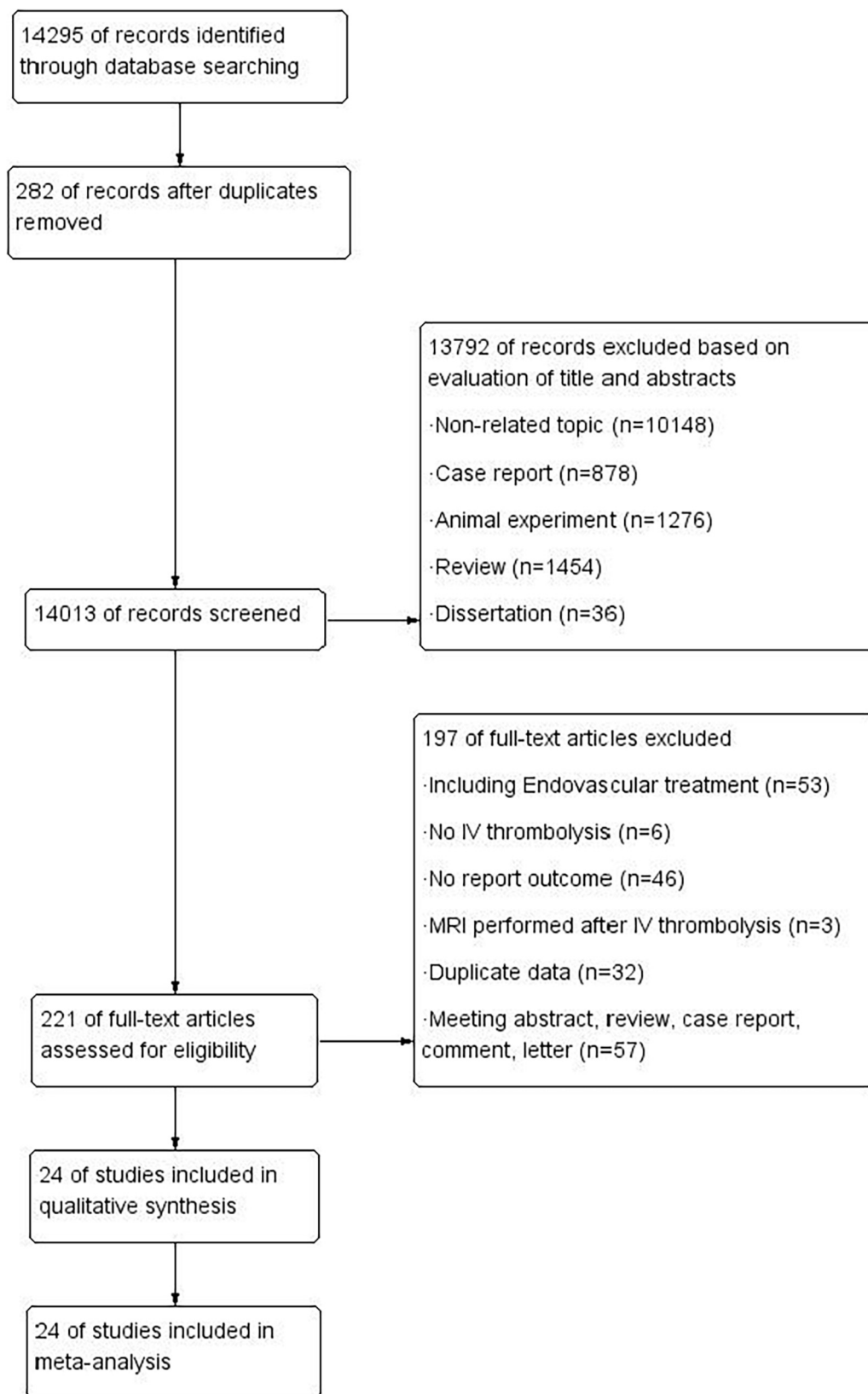


FIGURE 1 | Flow diagram of the search process and study selection.

TABLE 1 | Characteristics of the studies in the meta-analysis.

Author (year)	Period	Country	Study design	Sample	Mean/Median Age(y)	Gender (Female/male)	CVSD	Clinical outcomes	Imaging method	Time-HT/sICH	NOS
Moriya et al. (2013)	2007.11–2011.12	Japan	Prospective	71	73 ± 10	21/50	CMB	HT(NINDS)	T2*-GRE(1.5T), FLAIR, DWI, T1WI, T2WI	Between days 4 and 7 after admission	6
Capuana et al. (2020)	NA	Italy	Retrospective	434	68.3 ± 13.5	170/264	CMB	HT(ECASS II) sICH(ECASS)	T2*-GRE(1.5T), T1WI, T2WI, FLAIR, DWI, PWI, 3d-TOF-MRA	Within 24 h after IVT	8
Liu et al. (2019b)	2016.08–2018.07	China	Prospective	218	65.79 ± 11.99	144/74	The total burden of CSVD, WMH	PFO	T2*-GRE (-), FLAIR, T2WI, T1WI, DWI, 3d-TOF-MRA	/	8
Wen et al. (2019)	2016.09–2019.01	China	Retrospective	326	Good outcome: 65 ± 12 Poor outcome: 70 ± 12	117/209	WMH	PFO	T1WI, T2WI, FLAIR	/	8
Liu L. et al. (2019)	2017.01–2019.01	China	Retrospective	175	67 ± 11	62/113	The total burden of CSVD	PFO, HT (ECASSII), sICH (ECASSII)	DWI, FLAIR, SWI, MRA	Within 24 h after IVT	8
Liu X.Y. et al. (2019)	2016.08–2018.01	China	Retrospective	154	66.00 (59.00, 74.25) (Median; Interquartile range)	51/103	The total burden of CSVD, CMB, LI, CMB, LI	HT(ECASSII), sICH(NINDS)	T1WI, T2WI, FLAIR, DWI	Within 24 h after IVT	6
Chacon-Portillo et al. (2018)	2011.01–2015.12	America	Retrospective	292	62.8 ± 15.3	151/141	CMB	sICH (NINDS), HT(ECASSII)	T1WI, T2WI, DWI, SWI	Within 24 h after IVT	6
Nagaraja et al. (2018)	2009.01–2013.12	America	Retrospective	366	67 ± 15	168/198	CMB	HT (ECASSI)	T2*-GRE(1.5T), DWI, FLAIR, CT (HT)	At 18–36 h after IVT	6
Liu et al. (2018)	2013.06–2017.05	China	Retrospective	97	66.6 ± 9.1	29/68	WMH	PFO, HT(ECASSII)	DWL, FLAIR, MRA-, CT (HT)	Within 24 h after IVT	8
Liu et al. (2017)	2014.01–2017.03	China	Retrospective	78	WMH: 73.7 ± 6.7 No WMH: 61.3 ± 10.6	24/54	WMH, LI	PFO, HT(ECASSII), sICH (ECASSII)	T1WI, FLAIR, DWI, MRA	Within 24 h after IVT	8
Yan et al. (2015a)	2009.06–2015.06	China	Retrospective	449	66.8 ± 12.9 SICH: 73 (64, 79)	151/298	CMB	PFO, HT(ECASSII), sICH(ECASSII)	SWI	Within 24 h after IVT	7
Curtze et al. (2015a)	2001.12–2014.02	UK	Retrospective	2481	No SICH: 69 (60, 77) (Median; Interquartile range)	1,074/1,407	WMH	sICH(ECASSII)	CT	At 24 h post IVT	6
Dannenberg et al. (2014)	2008.01–2013.08	Germany	Prospective	326	76 (68, 84) (Median; Interquartile range)	167/159	CMB	sICH(ECASSIII)	T2*-GRE(3T), DWI	Within 36 h after IVT	8

(Continued)

TABLE 1 | Continued

Author (year)	Period	Country	Study design	Sample	Mean/Median Age(y)	Gender (Female/male)	CVSD	Clinical outcomes	Imaging method	Time-HT/sICH	NOS
Yan et al. (2014)	2009.06–2013.05	China	Retrospective	225	66.29 ± 13.01	73/152	CMB	HT(ECASSII)	FLAIR, DWI, PWI, SWI, MRA	Within 24 h after IVT	8
Zheng et al. (2012)	2006.07–2011.10	China	Retrospective	175	68 ± 10	70/105	WMH	HT(ECASSII)	CT, T2WI, FLAIR	Within 24 h after IVT	7
Kakuda et al. (2005)	2001.04–2005.01	America, Canada, Belgium	Prospective	70	71 ± 29	39/31	CMB	HT(ECASS) sICH(ECASS)	GRE(1.5T), DWI, PWI, MRA, T1WI	At 3–6 h after IVT and at day 30	8
Dereix et al. (2004)	2001.05–2002.08	France	Retrospective	44	63.2 ± 14.1	21/23	CMB	HT(NINDS), sICH(NINDS)	T2*-GRE(1.5T), CT(HT, sICH)	At day 7	6
Yan et al. (2015a)	2009.06–2014.02	China	Retrospective	333	66.15 ± 13.02	110/223	CMB	HT(ECASSII), PFO	GRE(3.0T), DWI, SWI	At 24 hours post IVT	6
Dong and Xia (2018)	2015.01–2016.12	China	Prospective	56	69.26 ± 2.25	25/31	CMB	HT(ECASS)	T1WI, T2WI, T2FLAIR, DWI, MRA, SWI	At 24 hours post IVT	6
Zhang and Zhang (2018)	2015.07–2016.07	China	Retrospective	206	CMB: 63.2 ± 9.5 NO CMB: 61.5 ± 9.0	83/123	CMB	sICH(SITS-MOST), PFO	T2*-GRE(-), T1WI, T2WI, DWI	NA	6
Huang (2017)	2010.01–2016.01	China	Retrospective	100	18–80	42/58	CMB	HT(ECASSI)	SWI, CT(HT)	At 24 hours post IVT	6
Zhuo et al. (2020)	2012.03–2018.01	China	Retrospective	178	62.3 ± 10.5	53/125	CMB, LI	PFO	T1WI, T2WI, DWI, FLAIR, SWI	/	8
Xue et al. (2017)	2012.01–2015.06	China	Prospective	80	56 ± 12	22/58	CMB	HT(ECASSII)	DWI, MRA, SWI	At 24 ± 12 hours post IVT	6
Curtze et al. (2015b)	2001.12–2014.02	UK	Retrospective	2485	78 (72–83)	1076/1409	WMH	PFO	CT	/	6

CMB, cerebral microbleed; WMH, white matter hyperintensity; LI, lacunar infarction; The total burden of CSVD, the total burden of cerebral small vessel disease; NA not available; HT, hemorrhagic transformation; sICH, symptomatic intracranial hemorrhage; PFO, poor functional outcome; ECASS, European Cooperative Acute Stroke Studies; SITS-MOST, safe implementation of thrombolysis in stroke-monitoring study; NINDS, the National Institute of Neurological Disorders and Stroke; T1WI, T1 Weighted Image; T2WI, T2 Weighted Image; FLAIR, Fluid attenuated inversion recovery; SWI, susceptibility-weighted imaging; T2*-GRE, T2*-weighted gradient-recalled echo; DWI, Diffusion-Weighted Imaging; PWI, Perfusion-Weighted Imaging; CT, Computed Tomography; MRA, Magnetic Resonance Angiography; 3d-TOF-MRA, 3D time-of-flight Magnetic Resonance Angiography; CT (HT, sICH), CT was used to detected HT, sICH.

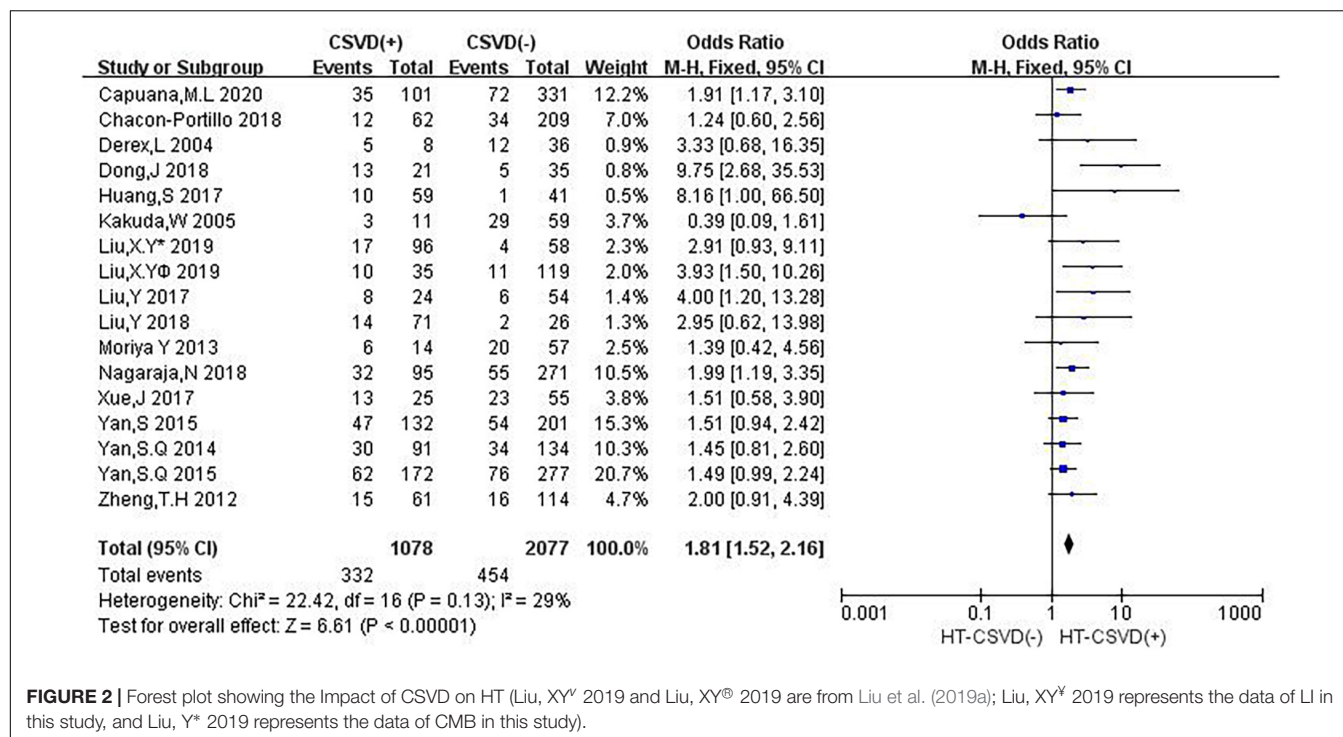


FIGURE 2 | Forest plot showing the Impact of CSVD on HT (Liu, XY[®] 2019 and Liu, XY[®] 2019 are from Liu et al. (2019a); Liu, XY[®] 2019 represents the data of LI in this study, and Liu, Y* 2019 represents the data of CMB in this study).

eliminated each study one by one to explore its influence on the conclusion. We planned to use funnel plots to visually observe the possibility of reporting bias and combined the Egger's test to detect publication bias (Egger et al., 1997). Two-sided $P \leq 0.05$ was considered statistically significant.

RESULTS

Review of the Literature

Our initial search yielded 14,295 potentially related studies. Of note, 282 studies were excluded because they were duplicates, 13,792 were excluded based on titles and abstracts, and 197 were excluded after a full-text reading revealed that the studies did not meet the inclusion criteria. Finally, for a total of 9,419 patients, 24 studies (Dere et al., 2004; Kakuda et al., 2005; Zheng et al., 2012; Moriya et al., 2013; Dannenberg et al., 2014; Yan et al., 2014, 2015a,b; Curtze et al., 2015a,b; Liu et al., 2017, 2018; Huang, 2017; Xue et al., 2017; Chacon-Portillo et al., 2018; Nagaraja et al., 2018; Dong and Xia, 2018; Zhang and Zhang, 2018; Liu L. et al., 2019; Liu et al., 2019a; Liu L. et al., 2019; Wen et al., 2019; Capuana et al., 2020; Zhuo et al., 2020) meeting the criteria were included. The selection process is shown in **Figure 1**. These studies were cohort studies published from the year 2004 to 2020 and included 6 prospective and 18 retrospective studies. Among them, 17 studies evaluated HT, 11 studies mentioned sICH, and 10 studies assessed 3-month PFO. Three of these studies reported separate associations with the total burden of CSVD. In the included studies, the neuroimaging markers of CSVD included cerebral CMB, WMH, and LI, and no studies related to EPVS were found. The characteristics of these studies are shown in **Table 1**.

Neuroimaging Markers of CSVD and HT

Sixteen studies evaluated the relationship between neuroimaging markers of CSVD and HT. The pooled overall rate of HT after IVT was 24.9% in the entire population. The pooled rate was 30.8% of patients in the CSVD presence group vs. 21.9% of patients in the group without CSVD presence. The total number of participants was 3,155 patients. Compared with no CSVD, the presence of CSVD was associated with an increased risk of HT (OR: 1.81, 95% CI: 1.52–2.16) (**Figure 2**). The heterogeneity between studies was not significant ($I^2 = 29\%$, $P = 0.13$). The risk of HT after IVT was higher in patients with CSVD than in patients without CSVD on neuroimaging.

In subgroup analyses, the influence of CSVD on HT showed a few changes when the studies were stratified according to the type of CSVD (**Figure 3**). Twelve studies (50%) reported the relationship between CMB and HT, three studies (12.5%) evaluated the association between WMH and HT, and one study (4.2%) mentioned the relationship between CMB and HT as well as LI and HT. Compared with no CMB, the presence of CMB was associated with an increased risk of HT (OR: 1.72, 95% CI: 1.43–2.08), and moderate heterogeneity was observed ($I^2 = 38\%$, $P = 0.08$). Compared with no WMH, the presence of WMH was associated with an increased risk of HT (OR: 2.54, 95% CI: 1.39–4.64), and there was no heterogeneity ($I^2 = 0$, $P = 0.62$). An estimate specific for LI and HT was provided in only one study (OR: 2.91, 95% CI: 0.93–9.11).

We performed the sensitivity analyses to explore whether the change in the inclusion criteria of studies influenced the robustness of the combined results. We recalculated the pooled OR by excluding each individual study in turn. The range of the combined ORs was from 1.75 (95% CI: 1.46–2.09) to 1.89

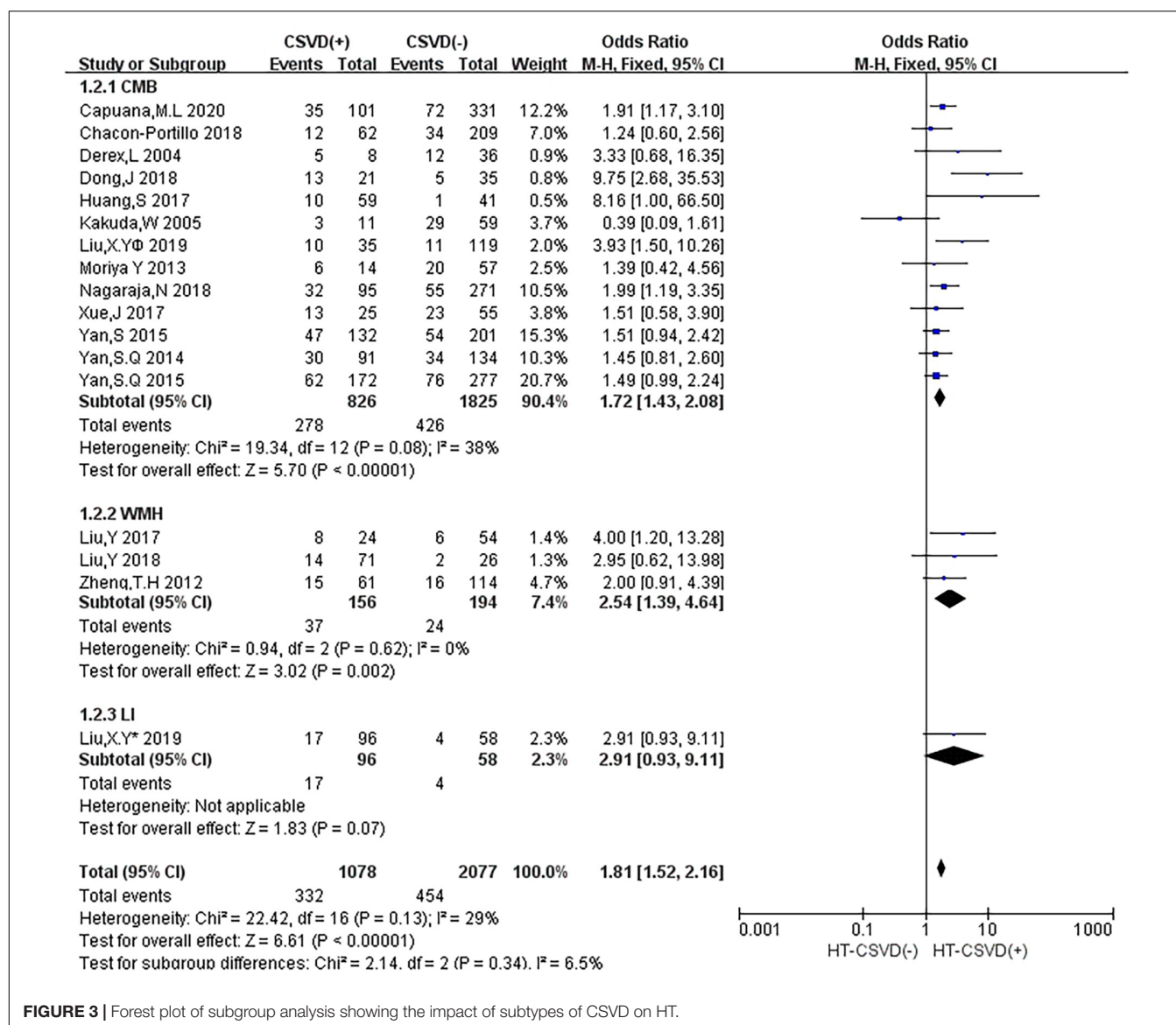


FIGURE 3 | Forest plot of subgroup analysis showing the impact of subtypes of CSVD on HT.

(95% CI: 1.56–2.30) when the studies of Dong and Xia (2018) and Yan et al. (2015a) were excluded. The results showed that no individual study significantly affected the pooled effect size. Furthermore, the funnel plots (Figure 4) and the Egger's test ($t = 1.88$, $P = 0.079$) indicated no publication bias.

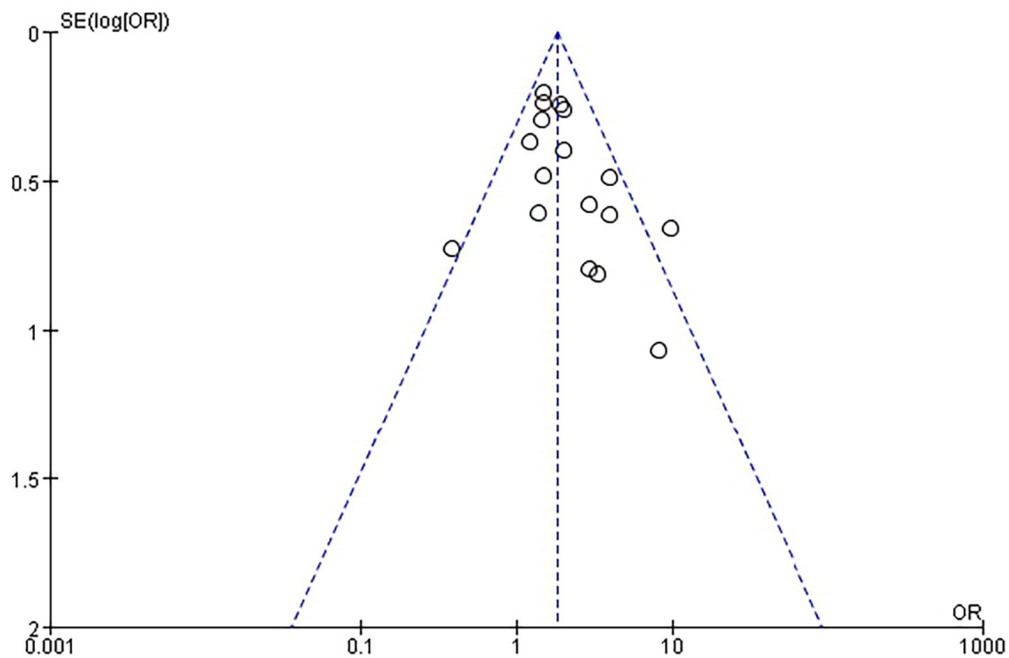
Neuroimaging Markers of CSVD and sICH

Nine studies reported the relationship between neuroimaging markers of CSVD and sICH. The pooled overall rate of sICH after IVT was 4.3% in the entire population. The pooled rate was 6.4% of patients in the CSVD presence group vs. 2.7% of patients in the group without CSVD presence. The total number of participants was 4,285 patients. Compared with no CSVD, the presence of CSVD was associated with an increased risk of sICH (OR: 2.42, 95% CI: 1.76–3.33) (Figure 5). The heterogeneity was

not detected ($I^2 = 0\%$, $P = 0.49$). The risk of sICH after IVT was found to be higher in patients with evidence of CSVD than in patients without CSVD on neuroimaging.

In subgroup analyses, different types of CSVD had slightly different effects on sICH (Figure 6). Seven studies (29.2%) reported the relationship between CMB and sICH, only two studies (8.3%) evaluated the association between WMH and sICH, and no study mentioned the correlation between LI and sICH. Compared with no CMB, the presence of CMB was associated with an increased risk of HT (OR: 2.86, 95% CI: 1.63–5.02), and low heterogeneity was observed ($I^2 = 5\%$, $P = 0.39$). Compared with no WMH, the presence of WMH was associated with an increased risk of sICH (OR: 2.27, 95% CI: 1.54–3.33), and there was no heterogeneity ($I^2 = 0\%$, $P = 0.61$).

We performed the sensitivity analyses to explore whether the outcome was stable. We recalculated the pooled OR by excluding each individual study in turn. The range of the combined OR



Egger's test

	Std_Eff	Coeff	Standard Error	t	p > t	[95% Confidence Interval]
Slope		.1967596	.2299342	0.86	0.406	-.2933335 .6868527
bias		1.15107	.6117395	1.88	0.079	-.1528215 2.454962

FIGURE 4 | Funnel plot and Egger's test of studies evaluating the association between CSVD and HT.

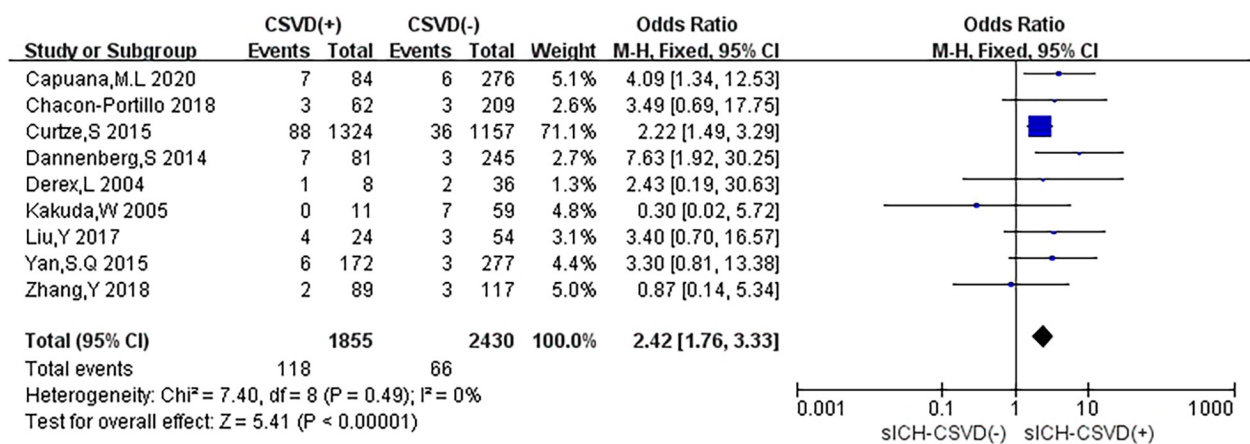


FIGURE 5 | Forest plot showing the impact of CSVD on sICH.

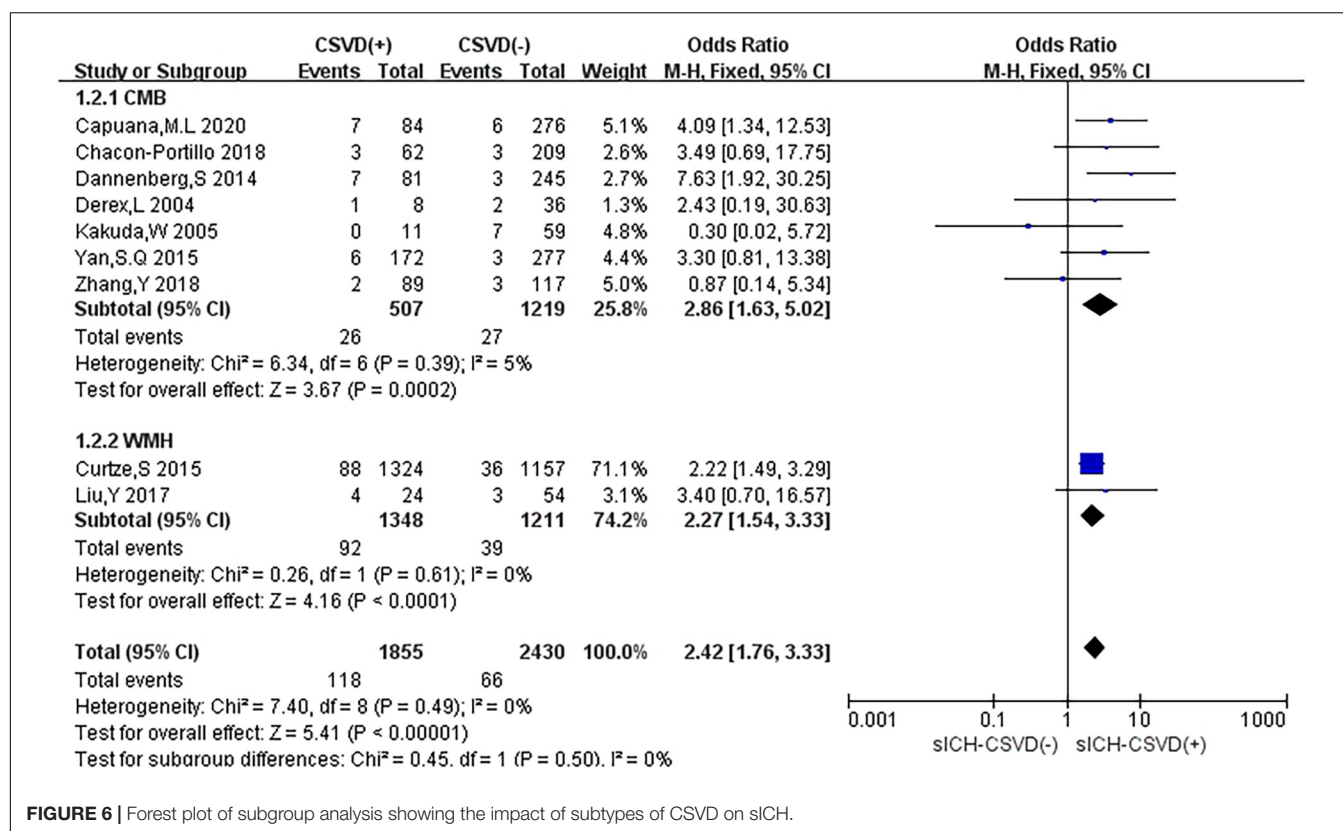


FIGURE 6 | Forest plot of subgroup analysis showing the impact of subtypes of CSVD on sICH.

was from 2.28 (95% CI: 1.64–3.16) to 2.92 (95% CI: 1.72–4.96) when the studies of Dannenberg et al. (2014) and Curtze et al. (2015a) were excluded. The results suggested that no individual study significantly affected the pooled effect size. The funnel plots (Figure 7) and the Egger's test ($t = 0.22$, $P = 0.833$) indicated no evidence of publication bias.

Neuroimaging Markers of CSVD and 3-Month PFO

Nine studies investigated the relation between neuroimaging markers of CSVD and PFO. The pooled overall rate of 3-month PFO after IVT was 38.8% in the entire population. The pooled rate was 45.8% of patients in the CSVD presence group vs. 31.6% of patients in the group without CSVD presence. The total number of participants was 4,626, and compared with no CSVD, the presence of CSVD was associated with an increased risk of 3-month PFO (OR: 2.15, 95% CI: 1.89–2.44) (Figure 8). The heterogeneity between studies was moderate ($I^2 = 40\%$, $P = 0.08$). The results showed that patients with evidence of CSVD were at higher risk for PFO after IVT than patients without neuroimaging evidence of CSVD.

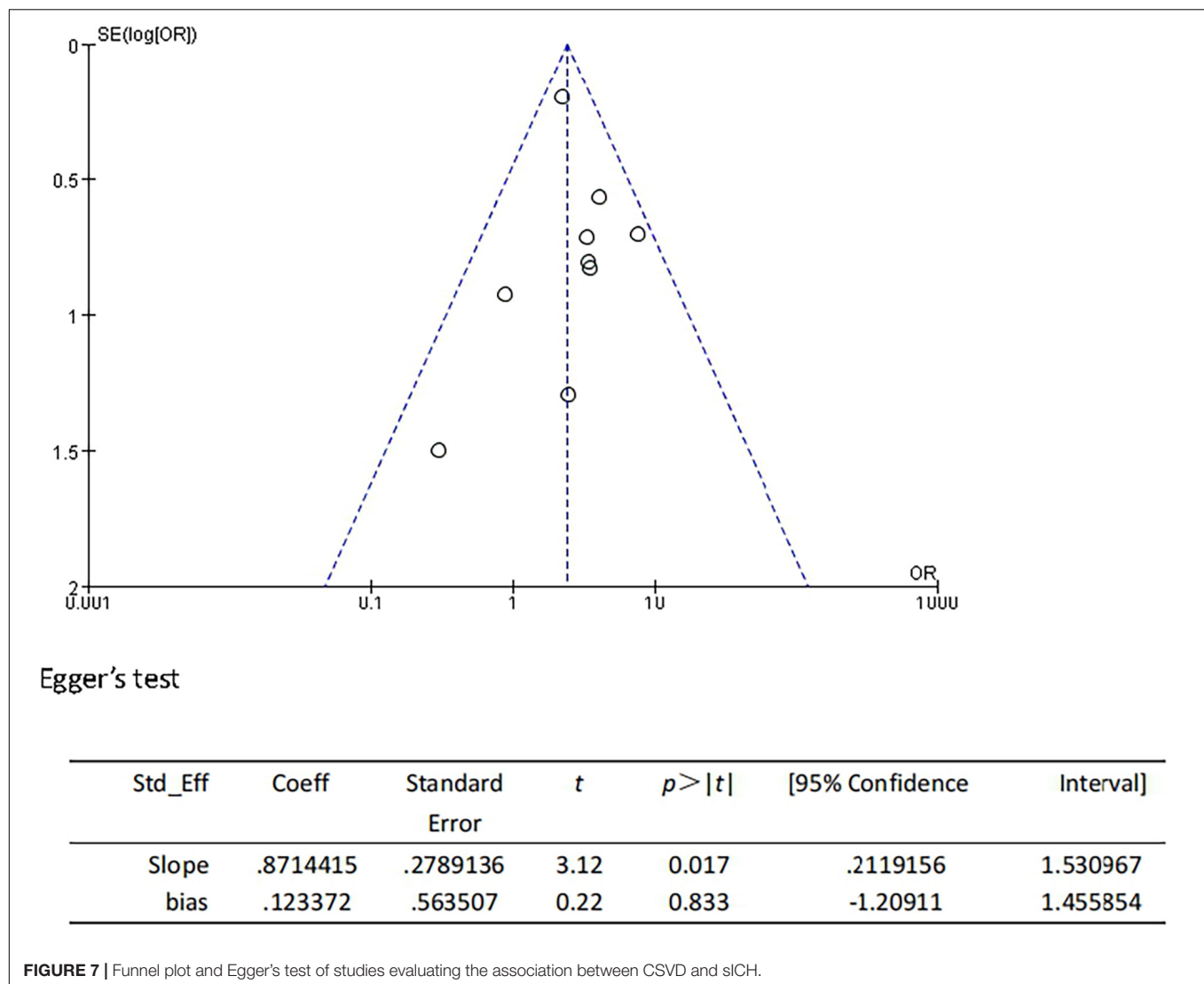
In subgroup analyses, the influence of CSVD on PFO showed some changes when studies were stratified according to the type of CSVD (Figure 9). Three studies (12.5%) reported the relationship between CMB and PFO, four studies (16.7%) evaluated the association between WMH and PFO, one study (4.2%) mentioned the relationship between CMB and PFO as well as LI and PFO, and one study (4.2%) reported the association

between WMH and PFO as well as between LI and PFO. Compared with no CMB, the presence of CMB was associated with an increased risk of 3-month PFO (OR: 1.65, 95% CI: 1.29–2.10), and no heterogeneity was observed ($I^2 = 0\%$, $P = 0.83$). Compared with no WMH, the presence of WMH was associated with an increased risk of 3-month PFO (OR: 2.32, 95% CI: 1.99–2.71), and there was significant heterogeneity ($I^2 = 55\%$, $P = 0.07$). Compared with no LI, the presence of LI was associated with an increased risk of 3-month PFO (OR: 3.18, 95% CI: 1.74–5.83). There was no evidence of substantial heterogeneity among studies ($I^2 = 0\%$, $P = 0.74$).

The sensitivity analysis was carried out to explore the robustness of our analysis, and we recalculated the pooled OR by excluding each individual study in turn. The range of the combined ORs was from 2.06 (95% CI: 1.69–2.52) to 2.21 (95% CI: 1.93–2.52) when the studies of Curtze et al. (2015b) and Yan et al. (2015a) were excluded. The results showed that no individual study significantly affected the pooled effect size. The funnel plots (Figure 10) and the Egger's test ($t = 0.69$, $P = 0.508$) showed no evidence of publication bias.

The Total Burden of CSVD

Three studies were related to the total burden of CSVD, among which one explored the relationship between the total burden of CSVD and HT, sICH, and PFO (Figure 11). One study (4.2%) reported the relationship between the total burden of CSVD and HT as well as sICH, and one study (4.2%) investigated the association between the total burden of CSVD and PFO. We



calculated pooled ORs of HT, sICH, and PFO for 2–4 CSVD scores vs. 0–1 CSVD scores. Compared with a CSVD score of 0–1, a CSVD score of 2–4 was associated with an increased risk of HT (OR: 3.10, 95% CI: 1.67–5.77), and no heterogeneity could be found ($I^2 = 0\%$, $P = 0.80$). Compared with a CSVD score of 0–1, a CSVD score of 2–4 was associated with an increased risk of sICH (OR: 2.86, 95% CI: 1.26–6.49), and there was no heterogeneity ($I^2 = 0\%$, $P = 0.52$). Compared with a CSVD score of 0–1, a CSVD score of 2–4 was associated with an increased risk of 3-month PFO (OR: 4.58, 95% CI: 2.97–7.06), and there was no evidence of substantial heterogeneity among studies ($I^2 = 0\%$, $P = 0.44$).

DISCUSSION

By using the systematic review and meta-analysis of these studies on the evidence of neuroimaging markers of CSVD, we observed that neuroimaging markers of CSVD (i.e., CMB, WMH, LI, and EPVS) represented an important indicator of a higher risk of

HT, sICH, and 3-month PFO after IVT in patients with AIS. In addition, we found that a CSVD score of 2–4 (i.e., the Total Burden Rating Scale of CSVD) was a potential risk indicator for incidents. Our meta-analysis confirmed the association between the presence of neuroimaging markers of CSVD and the risk of HT and PFO after IVT. At present, there are many reports on the relationship between various subtypes of CSVD and HT and clinical prognosis after IVT in patients with AIS, but no definite conclusion has been reached. To the best of our knowledge, our study is the first to systematically evaluate the risk of HT and clinical prognosis in patients with AIS combined with neuroimaging markers of CSVD after IVT.

Cerebral small vessel disease is a widespread cerebrovascular disease that mainly manifests in CMB, WMH, LI, and EPVS on imaging. It mainly affects the perforation of small arteries, capillaries, and venules. The increased risk of HT and PFO in patients with AIS with CSVD after IVT may be related to the following mechanisms: First, the changes in blood pressure variability may have an effect on the increased risk. Studies

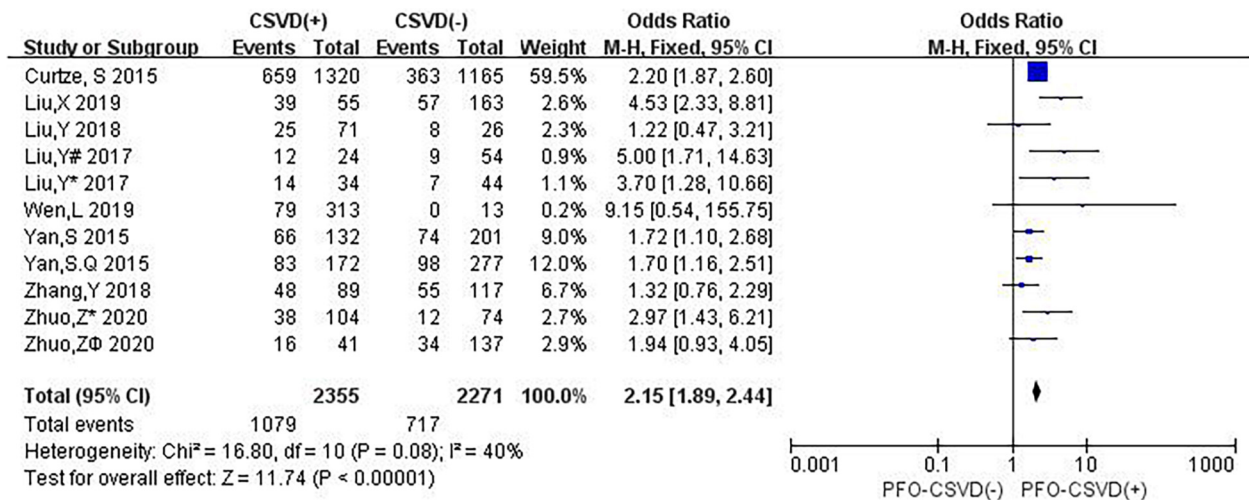


FIGURE 8 | Forest plot showing the impact of CSVD on 3-month PFO. (Liu, Y# 2017 and Liu, Y* 2017 are from Liu, Y 2017 (Liu et al., 2017); Liu, Y# 2017 represents the data of WMH in this study, and Liu, Y* 2017 represents the data of LI in this study; both Zhuo Z* 2020 and Zhuo Z Φ 2020 are from Zhuo et al. (2020); Zhuo Z* 2020 represents the data of LI in this study, and Zhuo Z Φ 2020 represents the data of CMB in this study).

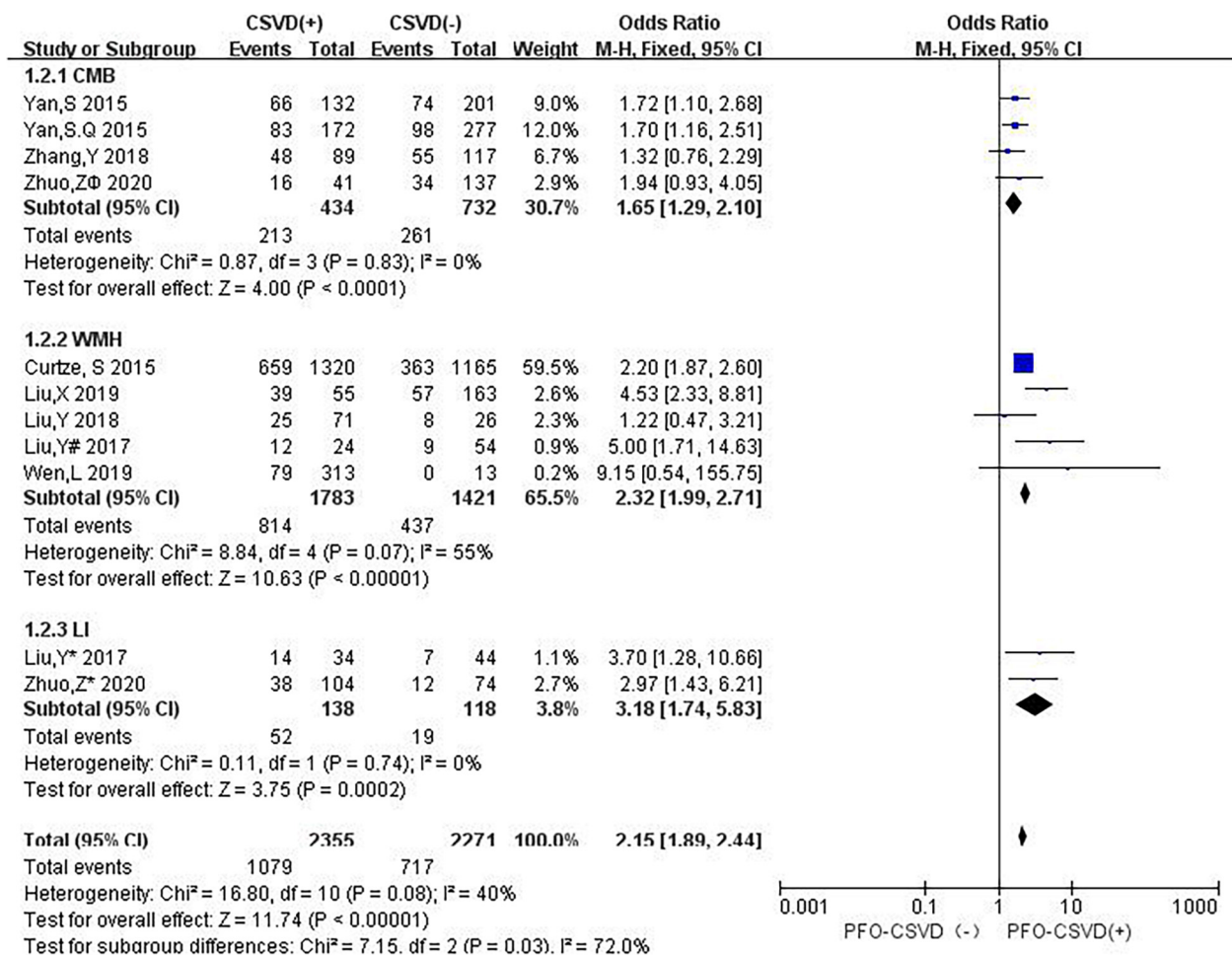


FIGURE 9 | Forest plot of subgroup analysis showing the impact of subtypes of CSVD on 3-month PFO.

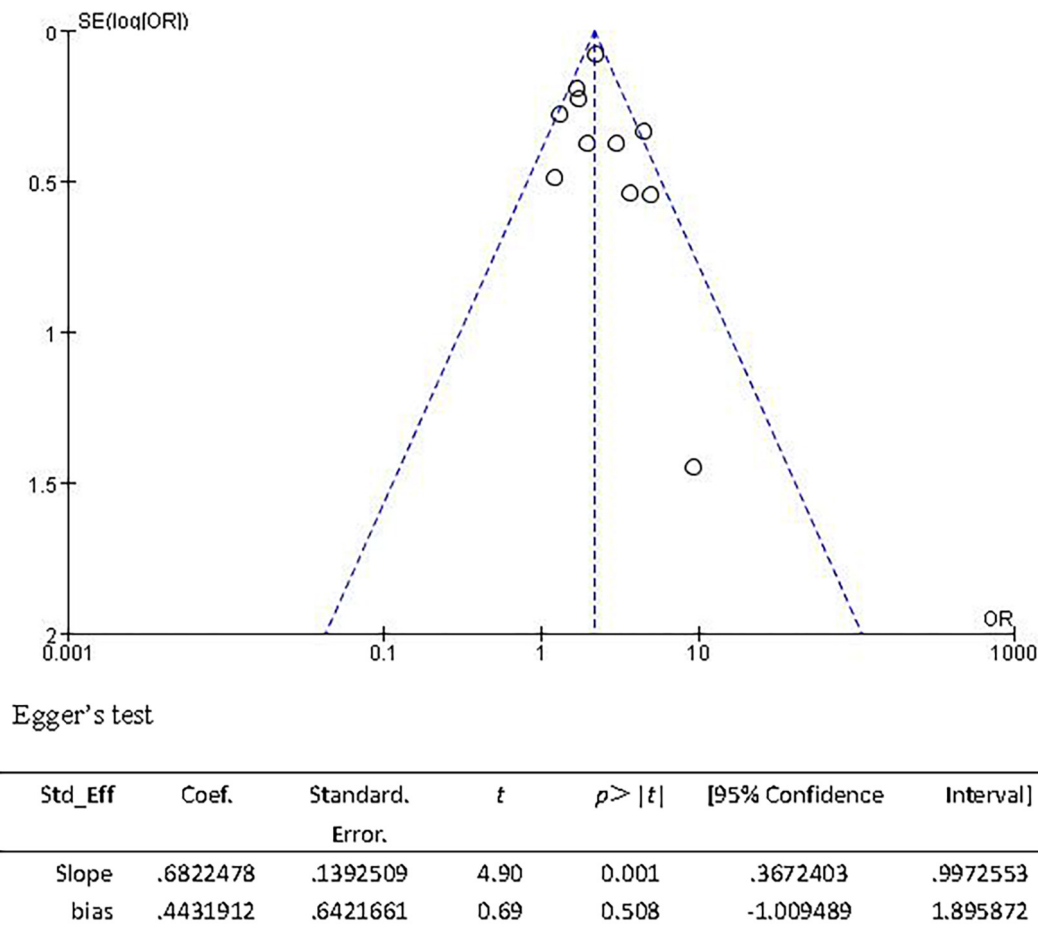
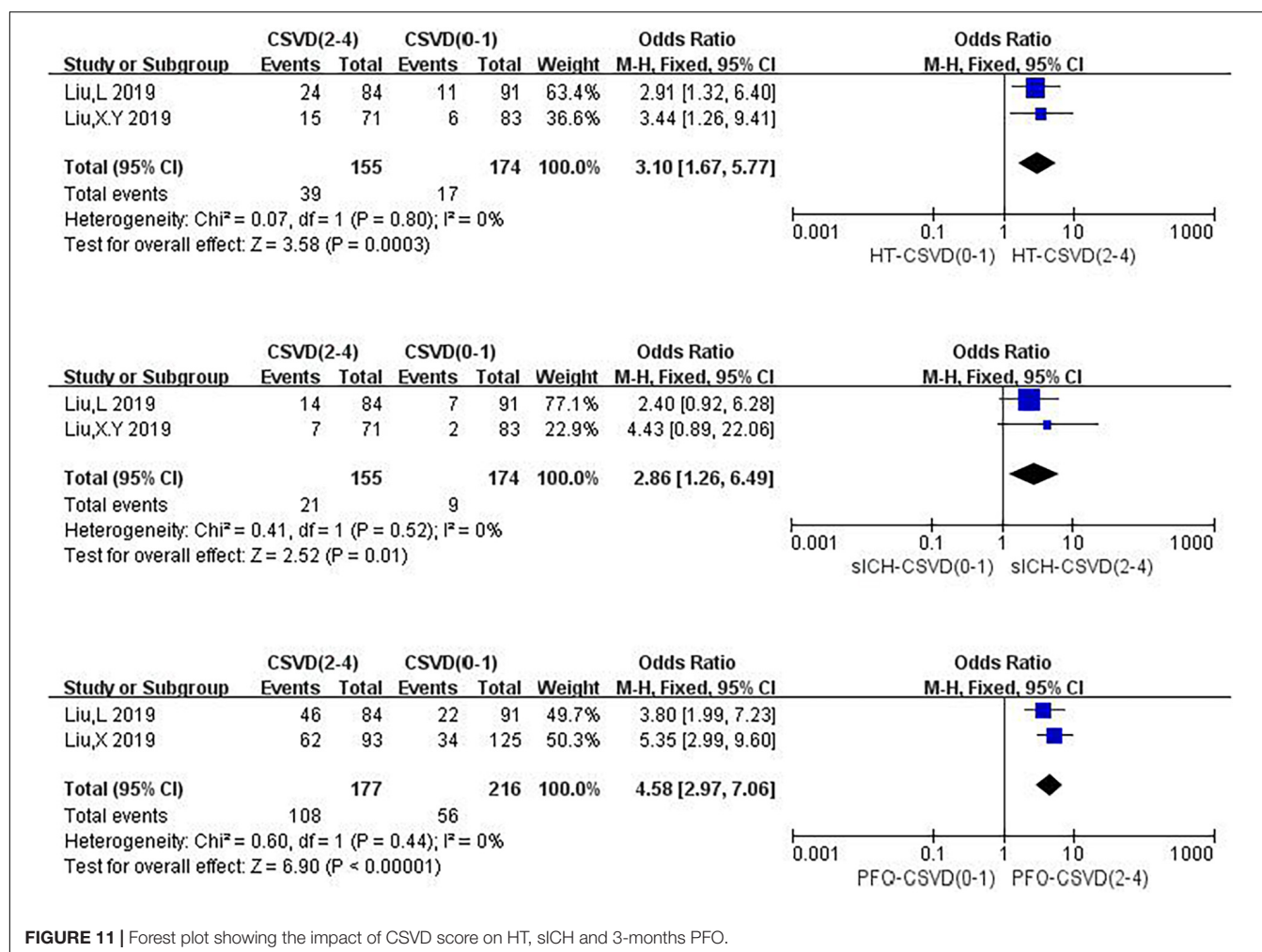


FIGURE 10 | Funnel plot and Egger's test of studies evaluating the association between CSVD and 3-month PFO.

have shown that increased blood pressure variability is closely related to the progression of CSVD, and increased blood pressure variability means increased blood pressure fluctuations, which increases the difficulty of maintaining the steady state of blood supply to the brain. The occurrence of hypoperfusion is more frequent (de Heus et al., 2020), which may be related to the occurrence of HT. Second, the destruction of the permeability of the blood–brain barrier, neuroimaging, and cerebrospinal fluid analysis indicate that leakage of the blood–brain barrier is a common feature of CSVD (Farrall and Wardlaw, 2009). When the permeability of the blood–brain barrier is broken, serum proteins and toxic effects will penetrate brain tissue, which may explain the complicated brain edema and poor functional prognosis in patients with AIS after IVT. Finally, the decrease in cerebral blood flow and vascular reactivity, the decrease in cerebral blood flow, and the decrease in vascular reactivity are related to CMB, WMH, LI, and EPVS (Gregg et al., 2015; Blair et al., 2016; Staszewski et al., 2019; Wang et al., 2020; Zhang et al., 2020). A previous meta-analysis of the cross-sectional studies revealed that patients with WMH had decreased cerebral blood flow (Shi et al., 2016), which may explain the PFO of patients with AIS after IVT. However, It is worth noting that IVT is still

the standard treatment for patients with AIS within 4.5 h of onset (Powers et al., 2019). Although there is a 2–7% risk of sICH after IVT treatment, the risk of disability or death is absolutely reduced by 10% at 3 months (Turc et al., 2014). There is currently no research to prove that CSVD is an absolute contraindication for IVT in the treatment of AIS, several studies have shown that LI will benefit from IVT (Eggers et al., 2017; Matusevicius et al., 2019). Therefore, we cannot arbitrarily regard the neuroimaging markers of CSVD as an absolute contraindication for IVT based on the results obtained in our study.

Different subtypes of CSVD may have different effects on HT and the clinical prognosis of patients with AIS after IVT. Several meta-analyses have studied the prognosis of patients with CMB and WMH. Our results are consistent with a recent meta-analysis including 11 studies, which reported that WMH significantly increased the incidence of sICH after IVT and showed that the combined incidence of WMH and sICH was 1.55, and the OR for WMH and 3- to 6-month PFO was 2.02 (Charidimou et al., 2016). However, a previous updated meta-analysis on a similar topic was performed by Wang et al. (2017) including 11 studies with 2,702 patients and showed that CMB presence was not significantly associated with an increased risk of sICH.



Compared with the previous meta-analysis, our results did not find such a connection. There are three possible reasons for this difference in the results. First, our study has a large sample size, including 24 articles (9,419 patients), and strict inclusion and exclusion criteria, while the earlier study only included 11 articles (2,702 patients). Second, in our study, the included patients with neuroimaging markers of CSVD could be tested by a variety of imaging methods, such as T1-weighted image (T1WI), T2WI, diffusion-weighted imaging (DWI), SWI, T2*-GRE, while in a previous study (Wang et al., 2017), only SWI and T2*-GRE were included in the imaging methods. Third, our results had low heterogeneity compared with previous studies (Charidimou et al., 2016; Wang et al., 2017).

Our study has the following strengths. First, our study is the earliest to systematically evaluate the risk of HT and PFO in patients with AIS combined with the neuroimaging markers of CSVD after IVT. Second, we applied rigorous methodological standards, performed a systematic search, included a large sample size, and conducted an in-depth analysis of different subgroups. Our study also has some limitations as follows. First, there are differences in neuroimaging methods, both CT and MRI, which may affect the diagnosis of CSVD. Although the

range of lesions shown on CT may not be ideal to expand the sample size, CT is also used to detect WMH. This may have a slight impact on the results of our study. However, we found that in previous studies, the application of CT to detect WMH was also be used for research (Charidimou et al., 2016). Second, the IVT protocol is different, and the baseline characteristics of the patient, such as previous use of anticoagulant drugs and repeated strokes, will affect the outcome. Third, the definitions of HT and sICH are different in each study. Although all we included are defined by standard criteria, they may still affect the results. However, in previous studies, the different definitions of HT and sICH, respectively, were combined for analysis (Charidimou et al., 2016; Wang et al., 2017). Fourth, in our study, in the subgroup analysis of CSVD and 3-month PFO, the presence of WMH was associated with the increased risk of 3-month PFO, and there was a significant heterogeneity. We found that the heterogeneity was observed from the study of Liu L. et al. (2019) probably due to the inclusion of participants, they excluded patients who had a pre-morbid mRS score > 2 . Finally, there were no studies with EPVS, and only three studies mentioned the overall burden of CSVD.

CONCLUSION

Our results indicate that patients with AIS complicated with the neuroimaging markers of CSVD are at an increased risk of HT and 3-month PFO after IVT. Our study strengthened the correlation between the neuroimaging markers of CSVD and the adverse outcome of AIS after IVT. However, it is undeniable that these patients can still benefit from IVT in some aspects. Therefore, it cannot be simply considered that the neuroimaging markers of CSVD are a contraindication for IVT in patients with AIS. It is still necessary to clarify the exact role of CSVD in the occurrence, development, and prognosis of AIS.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author/s.

AUTHOR CONTRIBUTIONS

YW and LL were responsible for conception and design of this systematic review. YW drafted the manuscript. LL and NX revised the manuscript. YW, JZ, and SG designed the search strategies. YW, XY, and PZ conducted the electronic search. YW and XY extracted the data. YW and GZ assessed the risk of bias. YW, HW, and LW analyzed and interpreted the data. LL arbitrated any disagreements in the process of systematic review. All authors approved this manuscript.

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The Neurovascular Unit in Dementia: An Opinion on Current Research and Future Directions

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INTRODUCTION

Dementia is a burgeoning public health crisis, with 50 million people currently affected worldwide (Prince et al., 2015). As the population ages, this figure is set to rise dramatically by 40% over the next 12 years (Prince et al., 2015). Dementia is an umbrella term for several disorders which result in the progressive loss of memory or other cognitive functions (Scott and Barrett, 2007). It remains an incurable disease, and current therapeutics have limited efficacy at slowing disease progression for one third of patients (Rockwood et al., 2008). Of the dementia sub-types, Alzheimer's disease (AD) remains the most prevalent, accounting for ~60–70% cases (Alzheimer's-Society, 2016). Vascular dementia (VaD) is the second most common form and is responsible for ~20% of cases, with a further 10% being a combination of these two diseases (Alzheimer's-Society, 2016). However, in practise these distinctions are somewhat arbitrary given the significant overlap in altered vascular structure and function in both of these major sub-types (Kalaria and Ballard, 1999). At least 30% of patients with AD have evidence of cerebrovascular disease on post-mortem examination, and almost all have evidence of cerebral amyloid angiopathy, microvascular degeneration, and white matter lesions (Kalaria and Ballard, 1999). Similarly, one-third of patients with VaD exhibit pathology consistent with AD (e.g., hippocampal or temporal lobe atrophy) (Kalaria and Ballard, 1999). Longitudinal studies have demonstrated that vascular risk factors (e.g., hypertension), significantly increase the risk of both AD and VaD (Rius-Pérez et al., 2018). In genetically at-risk individuals positive for apolipoprotein E4 (APOE4), atherosclerosis can increase the risk of AD by three-fold (Hoffmann et al., 2010). This article provides an opinion on the current evidence on the role of the neurovascular unit in dementia, for further information, several recent reviews are available on this topic (Nelson et al., 2016; Kisler et al., 2017).

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AMYLOID CASCADE HYPOTHESIS

A number of mechanistic models have been proposed to understand the pathological basis of AD. The amyloid cascade hypothesis gained increasing traction over the last few decades, having dominated the research sphere (Morris et al., 2014). Amyloid-based biomarkers have been incorporated into a number of diagnostic guidelines (Jack et al., 2018), and the histopathological (gold standard) diagnosis of AD includes the presence of amyloid plaque and neurofibrillary tangles (Deture and Dickson, 2019). However, despite decades of research into this hypothesis, and several large trials of amyloid based drugs, none have demonstrated efficacy warranting their widespread use in clinical practise (Morris et al., 2014). Only tramiprosate, a selective anti-oligomer agent, has demonstrated potential benefit for a sub-group of APOE4 positive individuals with early AD and is currently under investigation in a phase three trial (Tolar et al., 2020). These

findings have raised several questions around the amyloid cascade hypothesis. Firstly, the lack of efficacy for amyloid-based targets may suggest amyloid is a by-product rather than causative agent of the disease process. This is supported by the finding that amyloid deposition commonly occurs in cognitively healthy older adults, and plaque burden does not correlate well with the level of cognitive deficit (Morris et al., 2014). In contrast, synaptic loss, microglial activation, neurofibrillary tangles, and cerebral blood flow correlate better with disease severity in AD (Rius-Pérez et al., 2018). Secondly, the potential efficacy in a subgroup of early AD (Tolar et al., 2020) suggests that amyloid is a late occurrence in the disease process, at which stage irrevocable damage and cognitive decline has ensued. Furthermore, only patients with a strong genetic risk may benefit from these therapies (Tolar et al., 2020), limiting the wider applicability of these drugs. These unanswered questions have thus stimulated the search for earlier potential therapeutic targets, particularly those which are identifiable at earlier stages, preceding the development of cognitive decline and amyloid deposition.

VASCULAR CASCADE HYPOTHESIS

The vascular cascade hypothesis postulates that early disruption of vascular mechanisms as a result of sustained vascular risk factors and poor lifestyle habits, results in a state of chronic hypoperfusion (Rius-Pérez et al., 2018). This leads to the development of blood brain barrier breakdown, tau hyperphosphorylation, and amyloid deposition (Nelson et al., 2016). The blood brain barrier is essential to maintain a tightly controlled environment, and contributes to the clearance of amyloid-beta (Rius-Pérez et al., 2018). BBB dysfunction has been demonstrated to occur in the hippocampus with normal ageing (Montagne et al., 2015), early AD (Nation et al., 2019), and in APOE4 positive individuals (Montagne et al., 2020). As a result, amyloid deposition damages the cerebrovasculature, both structurally and functionally, therefore worsening hypoperfusion in a cyclical fashion (Nelson et al., 2016). These findings led to the development of the two-hit hypothesis, where the vascular insult represents the first “hit” to the system, followed by the amyloid or second “hit,” with the two processes subsequently interacting in a dynamic manner to worsen hypoperfusion, increase tau hyperphosphorylation, and promote amyloid deposition (Nelson et al., 2016). Importantly, the vascular hit is thought to occur earlier in the disease process (Hays et al., 2016). This notion is supported by longitudinal studies of ageing demonstrating that alterations in cerebral haemodynamics are detectable in cognitively intact older adults, and are predictive of future dementia risk (Wolters et al., 2017).

THE NEUROVASCULAR UNIT IN DEMENTIA

The neurovascular unit is formed by the neurone and its supporting cells (astrocytes, endothelial cells, pericytes, and smooth muscle cells) (Iadecola, 2017). They are closely related both structurally and functionally to ensure the tight coupling of

neuronal activity and cerebral blood flow, termed neurovascular coupling (NVC) (Iadecola, 2017). This is achieved through feedforward and feedback mechanisms as a result of the release of active metabolites and chemical mediators (Hosford and Gourine, 2019). De-coupling of these processes has been shown to occur in animal models of AD (Girouard and Iadecola, 2006). Human studies have demonstrated conflicting findings of both increased (Corriveau-Lecavalier et al., 2019), and decreased (Beishon et al., 2018) vascular responses to cognitive stimulation. These opposing findings may reflect compensatory mechanisms occurring early in the disease process, vs. the failure of these mechanisms at later stages (Merlo et al., 2019).

Therefore, deficiencies occurring in one or more components of the NVU threaten this tightly coordinated system. Inadequate matching of perfusion to neuronal activity will fail to clear the active metabolites generated by a resource intensive process, the accumulation of which can result in neurotoxicity (Girouard and Iadecola, 2006). Furthermore, inadequate perfusion will limit the provision of oxygen and glucose, essential for optimal neuronal function and cell signalling, thus limiting the capacity for cognitive function (Girouard and Iadecola, 2006).

THE NVU AS A BIOMARKER AND THERAPEUTIC TARGET IN DEMENTIA

As a result of these findings, increasing interest in the NVU as both a biomarker and therapeutic target has emerged. A number of neuroimaging based methods have been used to detect abnormalities in cerebral haemodynamics occurring in healthy, mildly impaired, or established dementia (Hays et al., 2016). A number of neuroimaging based biomarkers have been investigated, and can be broadly divided into portable and non-portable based techniques. Portable techniques have the advantage of providing a simple, bedside measurement with excellent temporal resolution and continuous monitoring of haemodynamic measures (Panerai, 2009; Balardin et al., 2017). Studies measuring metabolic changes, as a proxy for perfusion, have demonstrated good sensitivity and specificity to differentiate stable and progressive forms of mild cognitive impairment (MCI) (Henderson, 2012; Marcus et al., 2014). However, many of these techniques remain confined to the research domain, and are only recommended where the diagnosis remains uncertain (National Institute for Health Care Excellence, 2018).

In terms of vascular targets, the majority of research has focussed on currently available treatments to modify vascular risk, such as antihypertensive drugs (Bhat, 2015). Given the extensive evidence supporting a role for vascular mechanisms in the development of AD, modification of vascular risk is an attractive and amenable target. However, to gain benefit, these factors are likely to need controlling in mid-life given that these risks translate into cognitive decline over a sustained and longer period (Livingston et al., 2020). Furthermore, the role for vascular risk, and particularly blood pressure reduction, remains uncertain for people with established dementia (Harrison et al., 2016). A recent Cochrane review found limited evidence to support antihypertensive withdrawal in dementia, and may

result in increased cardiovascular events (Jongstra et al., 2016). Data from observational studies suggest cerebral autoregulation remains intact in MCI and dementia (De Heus et al., 2018), and a recent study demonstrated improved hippocampal CBF in patients with dementia treated with nilvadipine (Jong et al., 2019). The RADAR trial is currently ongoing, and will investigate the effects of losartan in mild to moderate AD on brain atrophy, white matter hyper intensities, and cerebral blood flow (Kehoe et al., 2018). Recently, interest has been gaining momentum on the effects of lifestyle interventions (exercises, diet, cognitive intervention) on cerebrovascular function, and whether multi-modal interventions can promote vascular brain health. In two recent systematic reviews (Beishon et al., 2020), cognitive training has been demonstrated to alter brain volumes and functional connectivity in MCI and dementia, but few studies have specifically investigated their effects on vascular mechanisms. Finally, novel therapeutic targets have been proposed around the various components of the NVU (Zlokovic, 2011). Vasculoprotective agents that target blood brain barrier function (e.g., activated protein C) and promote integrity are promising (Zlokovic, 2011). Similarly, mediators that promote angiogenesis (vascular endothelial growth factor) or improve amyloid-beta clearance (insulin like growth factor) may also be beneficial (Zlokovic, 2011).

DISCUSSION

In summary, vascular mechanisms play a key role in development and progression of cognitive dysfunction. Importantly, disruption to vascular physiology occurs early in the disease process, providing a potential target to prevent or delay the onset of dementia. Despite this breadth of evidence demonstrating both structural and functional damage to the cerebrovascular system in early dementia, few vascular targets have been the subject of large-scale randomised controlled trials. Disappointingly, in a recent review, few trials employed agents or targets of vascular dysfunction (Huang et al., 2020). This suggests more work is needed in both animal models to identify potential targets, and in patients to take these targets to clinical trials. Importantly, the identification of new targets has been hampered by a lack of translation between animal models and clinical trials (Cavanaugh et al., 2014). Current transgenic animal models of AD most closely represent inherited forms of AD, which are not the dominant phenotype seen in clinical practise

(Cavanaugh et al., 2014). These models will have a bias towards amyloid-based pathology, and may not reflect the alterations to vascular structure and function seen in humans, particularly with late-onset AD. Furthermore, the amyloid pathology in animal models does not correlate well with that seen in humans, suggesting there are key differences in the pathological basis of AD development between species (Cavanaugh et al., 2014). BBB dysfunction has been demonstrated in animal models of AD (Montagne et al., 2017), but amongst genetic-based models which may be pathologically distinct from late onset AD seen clinically. In addition to drug-based targets, research is urgently needed to clarify the role of lifestyle interventions on cerebrovascular disease in dementia risk reduction and treatment. Lifestyle interventions are resource intensive, and can be physically and mentally demanding for people with dementia to undertake. The Finnish Geriatric Intervention Study to Prevent Cognitive Impairment and Disability (FINGER) randomised at-risk older adults to an intensive programme of diet, exercise, cognitive training, and vascular risk monitoring, lasting 2 years (Ngandu et al., 2015). The trial found small benefits to cognitive function in the intervention group, with a drop-out rate of ~12% (Ngandu et al., 2015). Given that benefits to cognitive function tend to be small, and the long trajectory to cognitive decline, cerebrovascular biomarkers as a surrogate for clinical outcome measures could be beneficial in reducing the durations required for clinical trials to demonstrate effectiveness. However, limited information is available on the effects of such multi-modal interventions on cerebrovascular function, and their relationship to longer term clinical outcomes. Future trials of lifestyle interventions would benefit from the addition of cerebrovascular outcomes to understand the effects on vascular structure and function, which could contribute to the identification of novel therapeutic targets.

AUTHOR CONTRIBUTIONS

LB and RP jointly drafted the manuscript. All authors contributed to the article and approved the submitted version.

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Prostaglandin E₂ Dilates Intracerebral Arterioles When Applied to Capillaries: Implications for Small Vessel Diseases

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Prostaglandin E₂ (PGE₂) has been widely proposed to mediate neurovascular coupling by dilating brain parenchymal arterioles through activation of prostanoid EP4 receptors. However, our previous report that direct application of PGE₂ induces an EP1-mediated constriction strongly argues against its direct action on arterioles during neurovascular coupling, the mechanisms sustaining functional hyperemia. Recent advances have highlighted the role of capillaries in sensing neuronal activity and propagating vasodilatory signals to the upstream penetrating parenchymal arteriole. Here, we examined the effect of capillary stimulation with PGE₂ on upstream arteriolar diameter using an *ex vivo* capillary-parenchymal arteriole preparation and *in vivo* cerebral blood flow measurements with two-photon laser-scanning microscopy. We found that PGE₂ caused upstream arteriolar dilation when applied onto capillaries with an EC₅₀ of 70 nM. The response was inhibited by EP1 receptor antagonist and was greatly reduced, but not abolished, by blocking the strong inward-rectifier K⁺ channel. We further observed a blunted dilatory response to capillary stimulation with PGE₂ in a genetic mouse model of cerebral small vessel disease with impaired functional hyperemia. This evidence casts previous findings in a different light, indicating that capillaries are the locus of PGE₂ action to induce upstream arteriolar dilation in the control of brain blood flow, thereby providing a paradigm-shifting view that nonetheless remains coherent with the broad contours of a substantial body of existing literature.

Keywords: functional hyperemia, cerebral small vessel diseases, CADASIL, microcirculation, neurovascular coupling, potassium channel, prostaglandin E₂, epidermal growth factor receptor

INTRODUCTION

As a leading cause of stroke and dementia, cerebral small vessel diseases (SVDs) pose a horrendous threat to the elderly population (Pantoni, 2010; Wardlaw et al., 2019). Despite their major contribution to age-related vascular cognitive impairment and disability (Iadecola et al., 2019), the disease processes and key biological mechanisms underlying these disorders remain largely

unknown. Moreover, there are no specific treatments outside the management of vascular risk factors and use of anti-platelets after ischemic stroke. However, accumulating experimental evidence suggests that functional alterations in the cerebral microcirculation have early and deleterious consequences on functional hyperemia—the ability of the brain to increase local blood flow in response to local increases in neuronal activity (Huneau et al., 2018; Iadecola et al., 2019).

Cerebral Autosomal Dominant Arteriopathy with Subcortical Infarcts and Leukoencephalopathy (CADASIL) is the most common genetic form of cerebral SVDs. Remarkably, CADASIL includes all clinical and MRI manifestations of sporadic SVDs, hence offering a lens through which to view more common forms of sporadic SVDs (Chabriat et al., 2009). Both CADASIL patients and the *TgNotch3^{R169C}* mouse model, hereafter referred to as SVD mice, exhibit deficits in functional hyperemia at an early stage of the disease progression (Joutel et al., 2010; Huneau et al., 2018). We have recently demonstrated that activation of the epidermal growth factor receptor (EGFR) by its ligand heparin-binding EGF-like growth factor (HB-EGF) ameliorates the cerebral vascular deficits of the SVD mouse—including impaired functional hyperemia (Capone et al., 2016; Dabertrand et al., 2021). Part of this effect is mediated by reenabling capillary-to-arteriole signaling during the neurovascular coupling that underpins functional hyperemia (Dabertrand et al., 2021). In physiological conditions, action potentials increase extracellular K⁺ concentration which activates the strong inward-rectifier K⁺ (Kir2.1) channel in capillary endothelial cells (cECs). This creates a hyperpolarizing signal that rapidly propagates to upstream arterioles, driving vasodilation and local hyperemia (Longden et al., 2017; Harraz et al., 2018a; Moshkforoush et al., 2020). This mechanism is vulnerable to pathology, and we recently demonstrated that SVD lowers the synthesis of the phospholipid PIP₂, which prevents Kir2.1 channels from acting as sensors of increases in external K⁺ (Dabertrand et al., 2021). Strikingly, we showed that both HB-EGF and systemic injection of exogenous PIP₂ improved functional hyperemia deficits in SVD mice by restoring capillary-to-arteriole signaling.

Functional hyperemia is a vital process controlled by multiple mechanisms that provide layers of redundancy, and thus neurovascular coupling agents other than K⁺ ions have been postulated (Attwell et al., 2010; Kaplan et al., 2020). Among these, prostaglandin E₂ (PGE₂), produced by cyclooxygenase-2 (COX-2) from arachidonic acid, has been widely proposed to be released from excitatory neurons to relax arteriolar smooth muscle cells (SMCs) by activation of the G_S protein-coupled prostanoid EP₄ receptor and subsequent cyclic adenosine monophosphate-dependent pathway (Zonta et al., 2002; Takano et al., 2005; Gordon et al., 2008; Attwell et al., 2010; Lacroix et al., 2015). Yet, this interpretation appears incompatible with our previous demonstration that PGE₂ causes constriction, rather than dilation, when applied directly to isolated cortical arterioles—an effect that occurs via activation of the G_q protein-coupled prostanoid EP₁ receptor (Dabertrand et al., 2013). Consistent with these observations, the EP₁ receptor is robustly expressed not only in the arteriolar SMCs, but also in cECs of the brain microcirculation (Vanlandewijck et al., 2018). We therefore

hypothesized that PGE₂ contributes to neurovascular coupling by initiating a vasodilatory signal from the capillaries to the upstream parenchymal arterioles, rather than by targeting SMCs directly. Using a combination of *ex vivo* and *in vivo* approaches, we report that capillary stimulation with PGE₂ induces upstream arteriolar dilation and increases blood flow *in vivo*. Consistent with defective functional hyperemia, we further show that PGE₂-induced capillary-to-arteriole signaling is blunted in the SVD mouse model and can be rescued by HB-EGF.

MATERIALS AND METHODS

Animals

All experimental protocols used in this study were approved by the Institutional Animal Care and Use Committee (IACUC) of the University of Colorado, Anschutz Medical Campus. Adult (2–3 months old) male C57/BL6J mice (Jackson Laboratories, United States), were housed on a 12-h light:dark cycle with environmental enrichment and free access to food and water. *TgNotch3^{WT}* (WT) and *TgNotch3^{R169C}* (SVD) lines have been previously described (Joutel et al., 2010) and were used at 6 months of age in order for the *TgNotch3^{R169C}* mice to develop the SVD phenotype, and for consistency with our previous studies (Dabertrand et al., 2015, 2021; Capone et al., 2016; Fontaine et al., 2020). All mice were euthanized by i.p. injection of sodium pentobarbital (100 mg/kg) followed by rapid decapitation.

Ex vivo Capillary-Parenchymal Arteriole Preparation

The CaPA preparation was obtained as previously described (Longden et al., 2017; Rosehart et al., 2019) by dissecting intracerebral arterioles arising from the M1 region of the middle cerebral artery, leaving the attached capillary bed intact. Arteriolar segments were cannulated on glass micropipettes with one end occluded by a tie and pressurized using a Living Systems Instrumentation (United States) pressure servo controller with a mini peristaltic pump. The ends of the capillaries were then sealed by the downward pressure of an overlying glass micropipette. CaPA preparations were superfused (4 mL/min) with prewarmed (36°C ± 1°C), gassed (5% CO₂, 20% O₂, and 75% N₂) artificial cerebrospinal fluid (aCSF) for at least 30 min. The composition of aCSF was 125 mM NaCl, 3 mM KCl, 26 mM NaHCO₃, 1.25 mM NaH₂PO₄, 1 mM MgCl₂, 4 mM glucose, 2 mM CaCl₂, pH 7.3 (after aeration with 5% CO₂). Application of pressure (40 mmHg) to the cannulated parenchymal arteriole segment in this preparation pressurized the entire tree and induced myogenic tone in the arteriolar segment. Only viable CaPA preparations, defined as those that developed pressure-induced myogenic tone greater than 15%, were used in subsequent experiments. Arteriolar viability was validated by bath-applying NS309 (1 μM), which causes an endothelial-dependent vasodilation through activation of small- and intermediate-conductance, Ca²⁺-sensitive K⁺ (SK and IK, respectively) channels, or the thromboxane receptor agonist U46619 (1 μM), which causes robust vasoconstriction.

Dilatory responses of the attached arteriole segment to K⁺ and PGE₂ were obtained by applying aCSF containing 10 mM K⁺ or PGE₂ onto capillary extremities by pressure ejection from a glass micropipette (tip diameter, ~5 μm) attached to a Picospritzer III (Parker, United States) at ~5 psi for 20 s. The dose-responses were performed using pipettes filled with the different concentrations of PGE₂ and testing them sequentially starting with the lowest concentration. Contrary to arteriolar endothelial cells, cECs do not express functional IK and SK channels and spatial restriction of the drugs applied onto the capillary ends was validated by the lack of response to local stimulation with NS309 (1 μM), as previously described (Rosehart et al., 2019). In some control experiments, K⁺ and PGE₂ were applied directly to the arteriole segment and the other drugs were applied via the bath perfusion. The luminal diameter of the parenchymal arteriole was acquired at 15 Hz using a CCD camera and IonWizard 6.2 edge-detection software (IonOptix, United States). Two regions were simultaneously recorded, zone 1 where the capillary tree sprouted from of the arteriole and zone 2 located 250 μm upstream of this, and diameter from both of these sites was averaged unless noted otherwise. Changes in arteriolar diameter were calculated from the average luminal diameter measured over the last 10 s of stimulation and were normalized to the maximum dilatory responses in 0 mM Ca²⁺ bath solution at the end of each experiment using the following equation: [(change in diameter)/(maximal diameter-initial diameter)] × 100.

In vivo Imaging of Cerebral Hemodynamics

As previously described (Longden et al., 2017), mice were anesthetized with isoflurane (5% induction and 2% maintenance) during the surgical procedure. The skull was exposed, cleaned, and a stainless-steel head plate was attached with a mixture of dental cement and superglue. Isoflurane was replaced with α-chloralose (50 mg/kg) and urethane (750 mg/kg) during recording. FITC-dextran (2000 kDa) was injected via the retro-orbital sinus to visualize the cerebral vasculature and for contrast imaging of RBCs. A penetrating arteriole, identified by the direction of the RBCs flowing into the brain, was followed and a downstream capillary was selected for study in cortical layers 2 or 3. A pipette was then maneuvered into the brain and positioned adjacent to the capillary under study, and aCSF containing, or not, 1 μM PGE₂ was ejected (200–300 ms, 8 ± 1 psi, ~4 picoliters). The ejected volume was monitored by including 0.2 mg/mL tetramethylrhodamine isothiocyanate (TRITC; 150 kDa)-labeled dextran in the pipette (Figure 1E). RBC flux data were collected by line scanning at 5 kHz. Images were acquired using a Zeiss LSM-7 multiphoton microscope (Zeiss, United States) equipped with a 20x Plan Apochromat 1.0 N.A. DIC VIS-IR water-immersion objective and coupled to a Coherent Chameleon Vision II Titanium-Sapphire pulsed infrared laser (Coherent, United States). FITC and TRITC were excited at 820 nm, and emitted fluorescence was separated through 500–550 and 570–610 nm bandpass filters, respectively.

Reagents

HC030031, HC067047, PGE₂, NS309, and SC51322 were purchased from Tocris Bioscience (United States); All other chemicals and reagents were obtained from Sigma-Aldrich (United States). The vehicle for HB-EGF solutions was 0.2-μm-filtered PBS containing 0.1% BSA.

Statistical Analysis

Data in figures and text are presented as means ± standard error of the mean (SEM). Statistical testing was performed using GraphPad Prism 8 software. All data passed the Kolmogorov-Smirnov test for normality. Statistical significance was determined using one-way analysis of variance (ANOVA) followed by Tukey's *post hoc* test, unless otherwise stated.

RESULTS

To investigate the effect of capillary stimulation with PGE₂ on upstream arteriolar diameter, we used our previously developed *ex vivo* capillary-parenchymal arteriole (CaPA) preparation that allows to apply vasoactive substances at specific points along the arteriole-capillary continuum by pressure ejection (Longden et al., 2017; Rosehart et al., 2019). Consistent with our previous report that PGE₂ acts as a vasoconstrictor (Dabertrand et al., 2013), local application of 1 μM PGE₂ directly on the arteriolar segment caused a decrease in diameter (Figures 1A–C). In contrast, and as expected (Longden et al., 2017; Dabertrand et al., 2021), 10 mM K⁺ also applied on the arteriole caused a robust dilation. Interestingly, either 10 mM K⁺ or 1 μM PGE₂ caused upstream dilations when applied onto capillary extremities and the amplitudes of these were virtually identical (56.2% ± 3.9% and 47.8% ± 4.2%, respectively) (Figures 1A–C and Supplementary Movie 1). However, the kinetics of the responses appear significantly different with slower onset (5.27 ± 1.35 s) and time to peak (9.65 ± 1.5 s) for PGE₂ compared to K⁺ responses, (1.7 ± 0.4 s and 3.87 ± 1.13 s, respectively).

We next tested the effect of increasing concentrations of PGE₂ locally applied to the arteriolar segment and observed a concentration-dependent constriction with a calculated EC₅₀ of 145 nM (Figure 1D). The arteriolar dilation in response to capillary stimulations with PGE₂ was also concentration-dependent with a calculated EC₅₀ of 70 nM and a maximal response at 1 μM, and thus we chose to use the latter concentration throughout the study for capillary stimulation (Figure 1E and Supplementary Movie 1). Finally, we tested the effect of a modest concentration of PGE₂ (500 nM) applied via the bath perfusion, hence stimulating both the capillary ends and the arteriolar segment, and measured a small but clear constriction (Figures 1F,G).

In our initial report of capillary-to-arteriole electrical signaling (Longden et al., 2017), we identified the cEC strong inward-rectifier K⁺ channel, Kir2.1, as the molecular cornerstone of the capillary-to-arteriole electrical mechanism elicited by capillary stimulation with 10 mM K⁺. Increasing extracellular K⁺ concentration from 3 to 10 mM activates Kir2.1 channels in cECs, which induces a regenerative hyperpolarization that

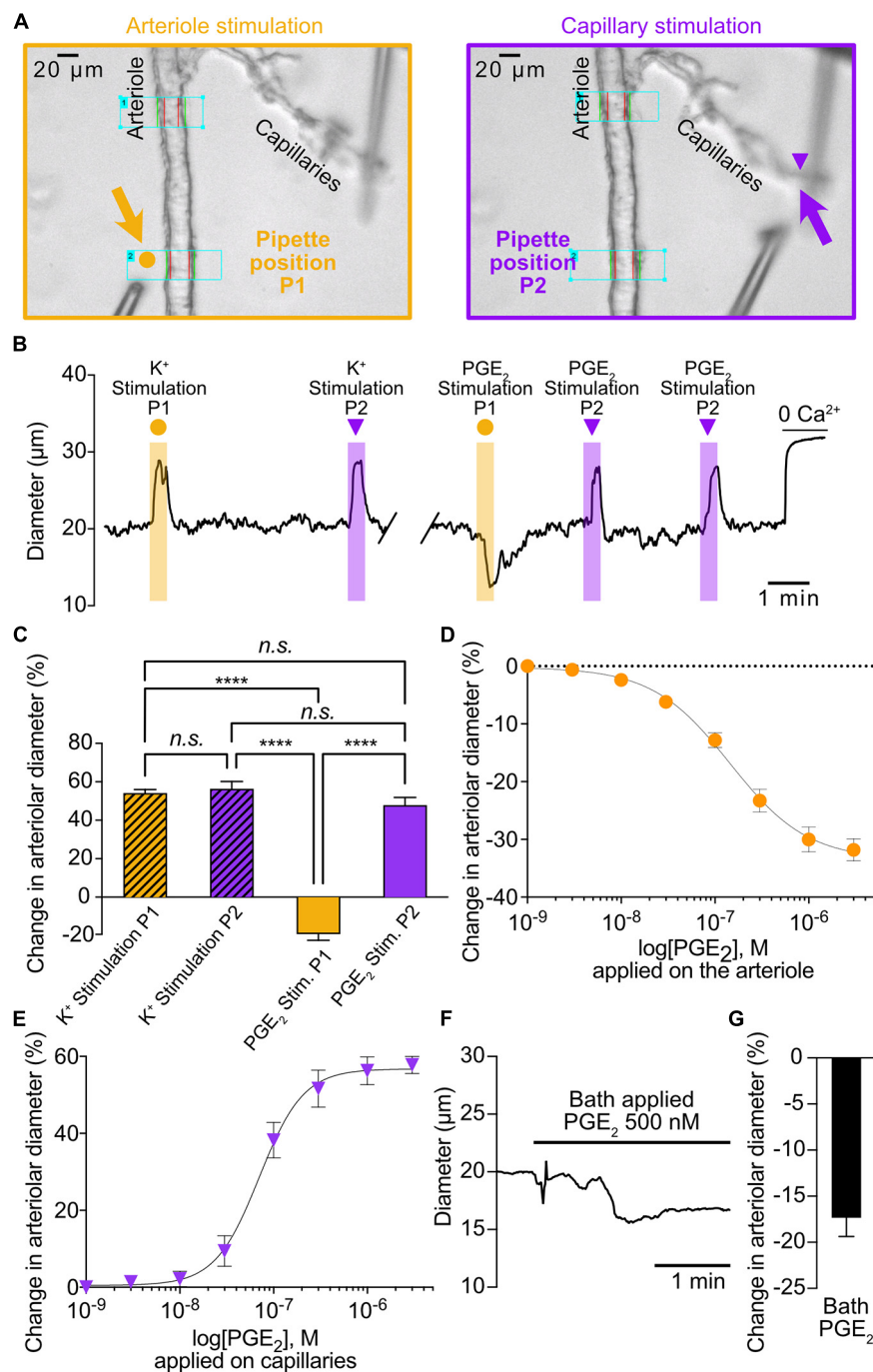
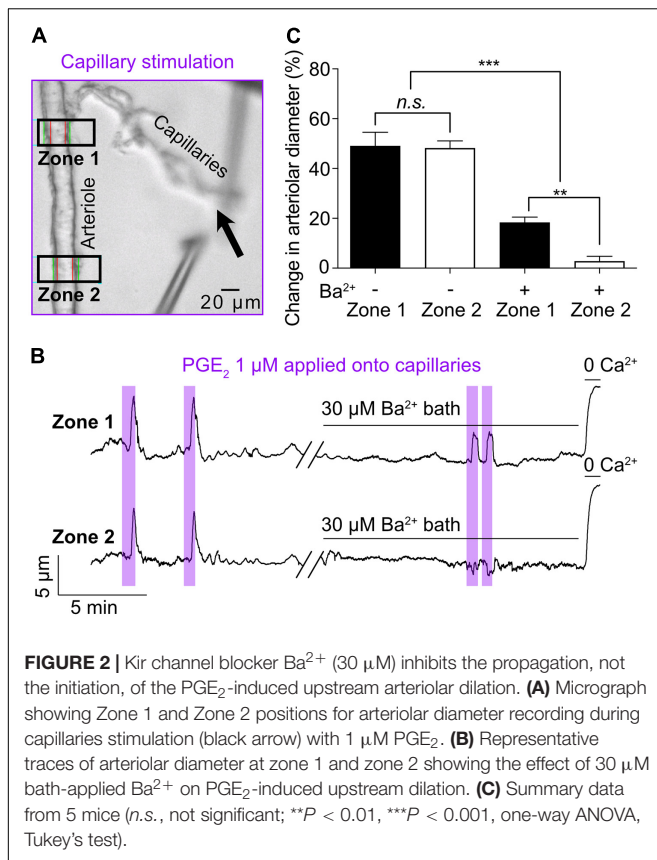
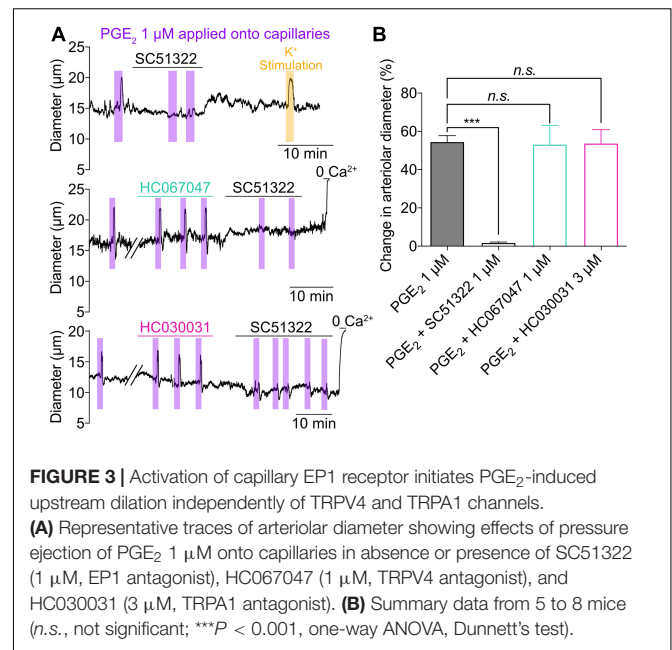


FIGURE 1 | PGE₂ causes upstream arteriolar dilation when applied onto capillaries. **(A)** Pipette positions for arteriole stimulation (left, orange arrow) and capillary stimulation (right, purple arrow) in CaPA preparations. **(B)** Representative trace of arteriolar diameter showing effects of pressure ejection of 10 mM K⁺ or 1 μM PGE₂ onto arteriole (P1, orange dot) and capillaries (P2, purple triangle). **(C)** Summary data from 6 mice (n.s., not significant; **** $P < 0.0001$, one-way ANOVA, Tukey's test). **(D)** Concentration-response curve produced by locally applying PGE₂ to the arteriolar segment over a concentration range of 1 nM to 3 μM (5 mice). An EC₅₀ of 145 nM was calculated from the non-linear regression curve. **(E)** Concentration-response curve produced by locally applying PGE₂ to capillary extremities over a concentration range of 1 nM to 3 μM (8 mice). An EC₅₀ of 70 nM was calculated from the non-linear regression curve. **(F)** Representative trace of arteriolar diameter showing effects of bath application of 500 nM PGE₂. **(G)** Summary data from 5 mice.



travels retrogradely to dilate the upstream arteriole. Consistent with this model, 30 μM Ba²⁺—which, among the inward rectifier K⁺ channels expressed in the microcirculation, preferentially blocks Kir2 channels (Longden and Nelson, 2015)—completely abolished the arteriolar dilation in response to capillary-applied 10 mM K⁺. Therefore, we tested the effect of Ba²⁺ on arteriolar dilation induced by capillary stimulation with PGE₂. Arteriolar diameter was recorded at two distinct zones: zone 1 was placed at the branching of the transitional segment from the arteriole and zone 2 was located 250 μm upstream of this (Figure 2A). PGE₂ locally applied onto capillaries led to similar dilatory responses in zone 1 (49.1% ± 5.5%) and zone 2 (48.2% ± 2.9%) (Figures 2B,C). Remarkably, bath application of 30 μM Ba²⁺ decreased upstream arteriolar dilation at zone 1 in response to capillary stimulation with PGE₂ by 63% and virtually eliminated dilation at zone 2. The onset of the remnant dilation observed at zone 1 was unchanged by Ba²⁺ (5.27 ± 1.35 s 6.49 ± 2.81 s). Taken together, these data suggest that the Kir2.1 channel is not required to initiate PGE₂-induced upstream arteriolar dilation, but rather participates in the amplification and propagation of the vasodilatory signal, as evidenced by the difference between zone 1 and zone 2.

To investigate the mechanistic underpinnings of the capillary-to-arteriole signaling induced by PGE₂, we first superfused the *ex vivo* CaPA preparation with 1 μM SC51322, a prostanoid receptor antagonist specific for the EP1 receptor. SC51322 abolished the response to PGE₂, while the dilation induced by



10 mM K⁺ remained intact (Figures 3A,B). This observation suggests that PGE₂ acts through activation of G protein-coupled receptors of the G_{q/11} subtype (G_qPCR) and subsequent Ca²⁺ signaling. Depletion of plasma membrane phosphatidylinositol 4,5-bisphosphate (PIP₂) following G_qPCR activation is known to stimulate transient receptor potential (TRP) channels, a major pathway for extracellular Ca²⁺ influx (Kim et al., 2008; Harraz et al., 2018b). Moreover, recent work by Thakore et al. (2021) has revealed that activation of TRPA1 channel in cECs can initiate a biphasic, propagating retrograde signal that dilates upstream parenchymal arterioles during functional hyperemia. However, the TRPA1 antagonist HC030031 at 3 μM had no effect on the dilation induced by capillary stimulation with PGE₂ (Figures 3A,B). Inhibition of TRPV4, another Ca²⁺-permeable TRP channel expressed by cECs (Harraz et al., 2018b), with 1 μM HC067047 also did not impact the effect of PGE₂ (Figures 3A,B). These results suggest that PGE₂ induces upstream arteriole dilation via activation of the EP1 receptor but independently of TRPV4 and TRPA1 channels.

We then tested the ability of PGE₂ to dilate upstream arterioles *in vivo* by measuring capillary red blood cell (RBC) flux with two-photon laser-scanning microscopy. Fluorescein isothiocyanate (FITC)-labeled dextran was used to visualize the cortical microcirculation and a pipette containing 1 μM PGE₂ was maneuvered into the brain through a cranial window and positioned next to a capillary of interest (Figure 4A). Pressure ejection (8 ± 1 psi) for up to 300 ms of PGE₂ evoked a significant increase in capillary RBC flux (Δ = 12 ± 4 RBCs/s) consistent with the notion that the mechanisms we observed in CaPA preparations are also at play in the intact system (Figures 4B–D). In contrast, ejection of aCSF vehicle had no effect on capillary RBC flux (Figure 4E). To determine whether PGE₂ also causes upstream dilation we next performed experiments in which we imaged at the parenchymal arteriole-transitional zone junction

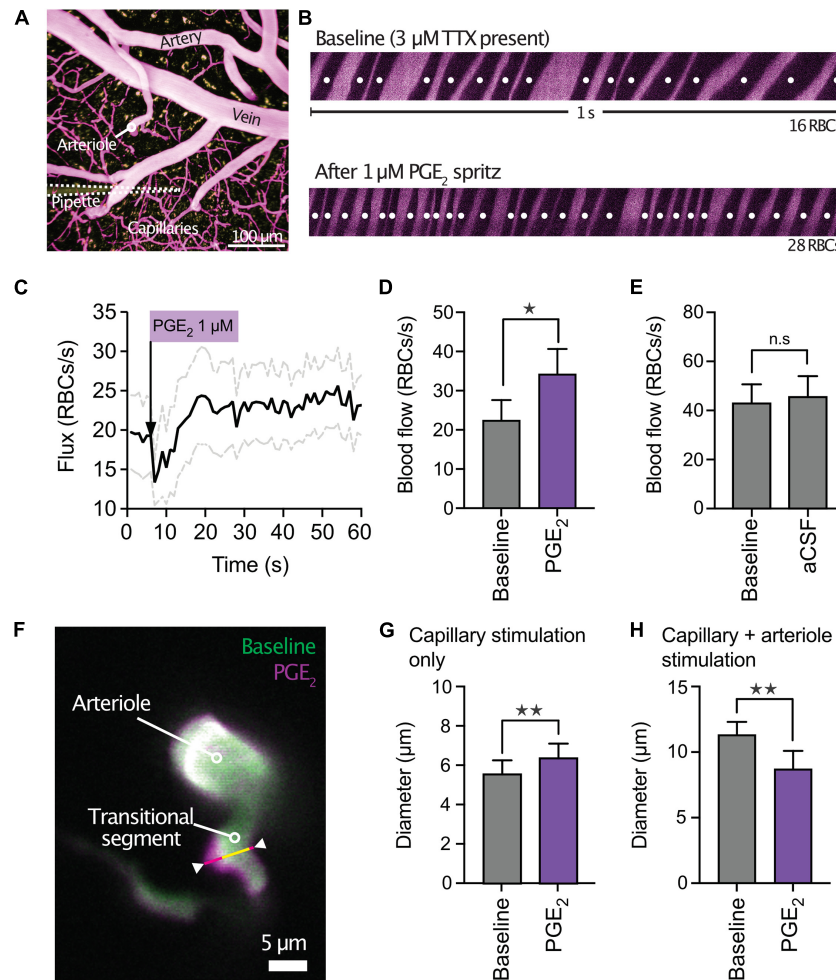
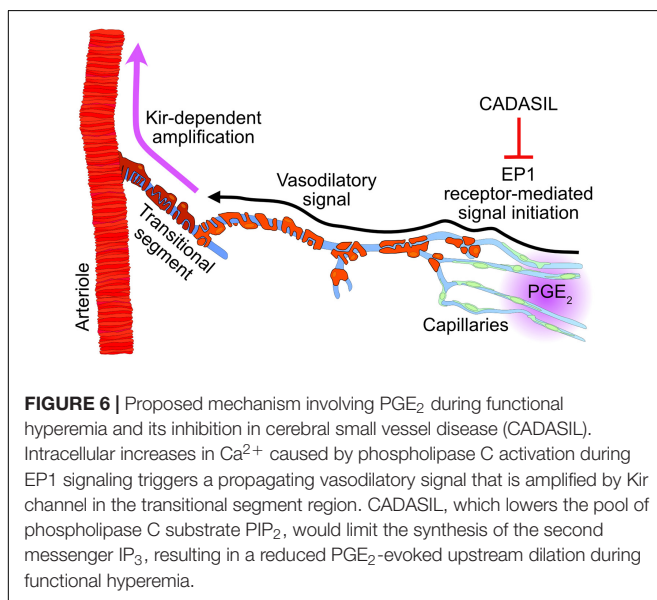
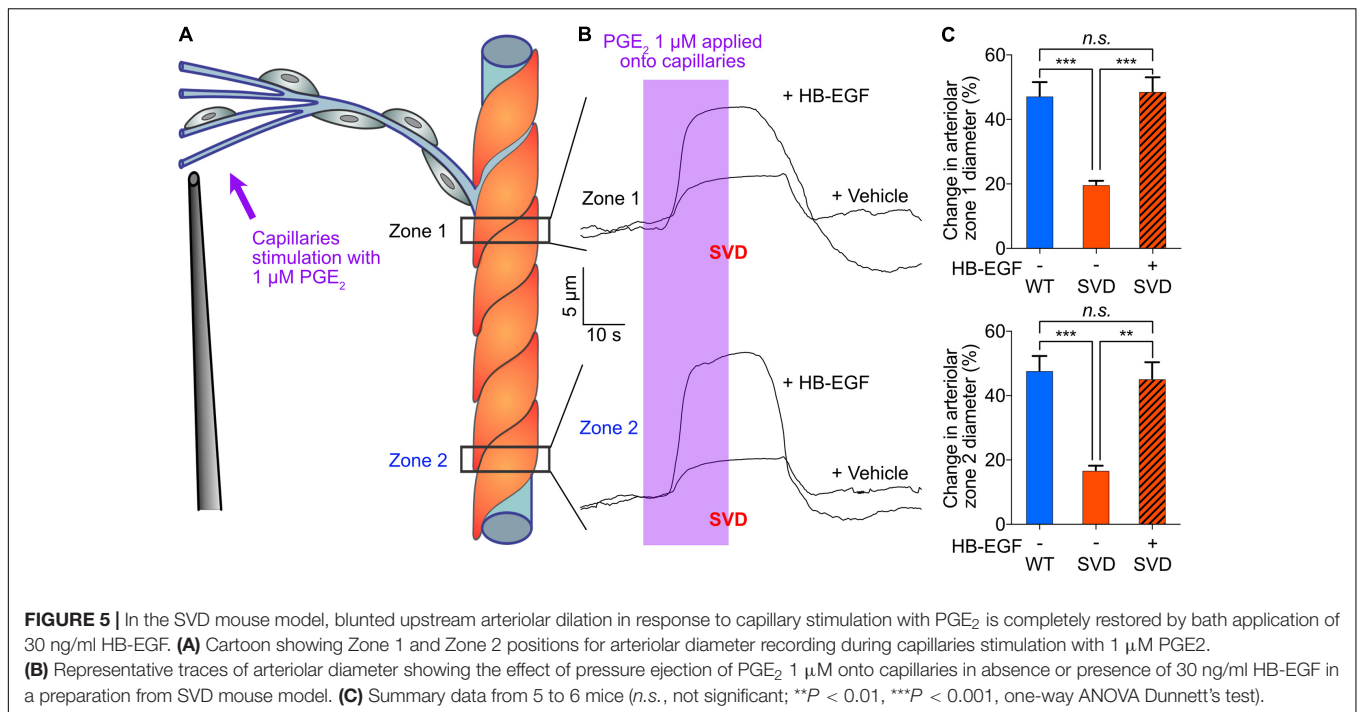


FIGURE 4 | PGE₂ causes capillary hyperemia *in vivo*. **(A)** Micrograph of mouse cortical vasculature showing a micropipette containing PGE₂ and TRITC-dextran (yellow) in close apposition to a capillary (FITC-dextran, purple). **(B)** Red blood cell (RBC) flux was measured by high-frequency line scanning over a period of 1 min at baseline (top panel) and after application of 1 μ M PGE₂ (lower panel) onto a capillary. RBCs appear as black streaks in plasma (purple). **(C)** Average traces (black line) plus SEM (gray lines) showing the increase in RBC flux to PGE₂. The dip immediately following the ejection of PGE₂ is caused by momentary pressure on the capillary wall. **(D)** Summary data of RBC flux showing significant hyperemia following capillary stimulation with 1 μ M PGE₂ ($n = 14$ experiments, 6 mice, $^*P < 0.05$, paired Student's t -test). **(E)** In contrast, ejection of aCSF vehicle onto capillaries had no effect on RBC flux ($n = 9$ experiments, 5 mice, $P > 0.05$, paired Student's t -test). **(F)** Overlay showing the diameter of a penetrating arteriole and the transitional segment to the capillary bed at baseline (green) and after PGE₂ ejection onto a downstream capillary (magenta). The upstream dilation was most prominent in the region highlighted by the white arrowheads. **(G)** Summary data showing upstream dilation to capillary ejection of PGE₂ ($n = 5$ experiments, 4 mice, $^{**}P < 0.01$, paired Student's t -test). **(H)** In contrast, simultaneous stimulation of both capillaries and arterioles with PGE₂ *in vivo* led to constriction ($n = 7$ experiments, 3 mice, $^{**}P < 0.01$, paired Student's t -test).

(Figure 4F). Here, we found that ejection of 1 μ M PGE₂ selectively onto capillaries routinely produced a small upstream dilation (Figures 4F,G and Supplementary Movie 2). In contrast, if we applied PGE₂ onto both capillaries and arterioles *in vivo* by increasing the duration of ejection, causing spread of the ejected solution to the upstream arteriole along a paravascular route, we observed constriction (Figure 4H), consistent with our *ex vivo* data. As expected, ejection of aCSF vehicle alone had no effect on upstream arteriolar diameter ($12.95 \pm 0.57 \mu\text{m}$ baseline diameter vs. $12.96 \pm 0.8 \mu\text{m}$ after aCSF ejection, $n = 6$ experiments, 3 mice, $P = 0.97$, paired Student's t -test).

Finally, we investigated the effect of SVD on PGE₂-initiated capillary-to-arteriole signaling. *Ex vivo* stimulation of capillaries from SVD mice with PGE₂ induced an upstream arteriolar

dilation that was 58% smaller at Zone 1 and 64.8% smaller at Zone 2 compared with that in *TgNotch3*^{WT} control (WT) mice, revealing attenuated PGE₂-mediated signaling (Figure 5). The difference between zone 1 and 2 dilations were not significant. We previously reported that activation of the epidermal growth factor receptor (EGFR) by its ligand heparin-binding EGF-like growth factor (HB-EGF) ameliorates the cerebral vascular deficits of the SVD mouse—including neurovascular coupling and functional hyperemia deficits (Capone et al., 2016; Dabertrand et al., 2021). We then directly tested the effect of HB-EGF on PGE₂-induced upstream vasodilation in *ex vivo* preparations from SVD mice. Bath-applied 30 ng/mL HB-EGF dramatically enhanced the upstream arteriolar dilation induced by capillary stimulation with PGE₂, increasing the average dilation from $19.8\% \pm 1\%$ to



48.7% \pm 4.3% at Zone 1, and from 16.8% \pm 1.4% to 45.3% \pm 5.1% at Zone 2 (**Figure 5** and **Supplementary Movie 3**). HB-EGF completely restored PGE₂-induced upstream dilation, abolishing the differences measured between WT and SVD animals.

DISCUSSION

Our progress in understanding functional hyperemia in health and disease has been hampered by large gaps in our comprehension of the mechanism underlying this

basic physiological response—and persistent controversies surrounding it (Kaplan et al., 2020). Our recent work identified Kir2.1 channels in cECs as the molecular cornerstone initiating and propagating a retrograde hyperpolarizing vasodilatory signal from capillaries to arterioles (Longden et al., 2017; Harraz et al., 2018a; Moshkforoush et al., 2020; Dabertrand et al., 2021). The present study extends this capillary-based paradigm, providing support for a new signaling modality that posits a central role for the G_qPCR EP1 receptor in mediating PGE₂-induced vasodilatory signal that propagates upstream to cause dilation of feeding parenchymal arterioles (**Figure 6**). Importantly, these findings hold the promise of resolving controversies surrounding how PGE₂, a widely proposed mediator of neurovascular coupling (Zonta et al., 2002; Takano et al., 2005; Gordon et al., 2008; Attwell et al., 2010; Watkins et al., 2014), can promote functional hyperemia despite evidence that it directly constricts arterioles (Dabertrand et al., 2013).

Our observations raise the immediate question of which effect of PGE₂, capillary-mediated vasodilation or direct arteriolar constriction, predominates during neurovascular coupling. We calculated an EC₅₀ of 145 nM for the PGE₂-induced constriction of the parenchymal arteriole, while the vasodilation induced by capillary stimulation displayed an EC₅₀ of 70 nM. In our experimental conditions, a simultaneous stimulation of capillaries and the arteriole with 500 nM PGE₂ led to a constriction, suggesting that the arteriolar effect prevails when both vascular segments are exposed to PGE₂. A possible explanation for these observations is that the microvascular response to PGE₂ has a kinetic component in which the fast constriction predominates over the slower capillary-mediated response. However, this stimulation via the bath perfusion likely does not reflect *in vivo* conditions. Accordingly, we tested the

effect of increasing our ejection duration *in vivo* such that PGE₂ not only stimulated the targeted capillaries but also spread upstream to the arteriole. Here too this maneuver produced arteriolar constriction, again suggesting that this response will predominate when the arteriole is exposed directly to PGE₂. Since the dense capillary network within the brain lies in close proximity to all neurons (Nishimura et al., 2007; Blinder et al., 2013), it is expected that cECs are the primary sensors of neuronal activity and any neurally derived PGE₂. Therefore, the vasodilatory effect would be expected to predominate during neurovascular coupling in physiological conditions and our data suggest that this occurs through local exposure of the capillaries to PGE₂. However, a variety of brain conditions, including ischemia and neurodegeneration (Minghetti, 2004), are known to up-regulate COX-2 expression in excitatory neurons which results in an EP1-dependent neurotoxicity of PGE₂ (Kawano et al., 2006). In this pathological situation, higher PGE₂ concentrations could lead to arteriolar constriction and then contribute to PGE₂ neurotoxicity by limiting local blood supply.

Neuronal activation leads to rapid increases in blood flow within, and on the surface, of the brain. Hillman and colleagues elegantly provided evidence for involvement of the endothelium in stimulus-evoked, conducted vasodilation from the brain parenchyma to arterioles and pial arteries *in vivo* (Chen et al., 2014). Our previous study on neurovascular coupling demonstrated how capillary endothelium is capable of transmitting an electrical signal to cause upstream vasodilation in support of functional hyperemia (Longden et al., 2017). We showed that activation of the Kir2.1 channel in cECs by extracellular K⁺ ions propagates a regenerative hyperpolarization from cell-to-cell up to the feeding arteriole to cause vasodilation (Longden et al., 2017; Harraz et al., 2018a; MacMillan and Evans, 2018). Interestingly, recent work from Thakore et al. (2021) showed that 4-hydroxynonenal (4-HNE), an endogenous product of lipid peroxidation, activates transient receptor potential ankyrin 1 (TRPA1) channel in cECs to cause upstream arteriolar dilation during functional hyperemia. These recent findings introduce the concept that a slowly propagating short-range Ca²⁺ signal is initiated in the capillary endothelium and converted into the fast-propagating hyperpolarization that causes dilation of upstream arterioles. The conversion is proposed to occur in the transitional region between the capillaries and the arteriole (Ratelade et al., 2020) by activation of the small- and intermediate-conductance Ca²⁺-activated K⁺ channels (SK and IK, respectively) and amplification of the hyperpolarization by Kir channel (Thakore et al., 2021). The transitional region refers to the first segment sprouting out of the arteriole, visible on **Figure 1** micrographs, while local stimulations are applied onto the 3rd and 4th order capillary branches after the transitional region. A vast body of literature reports (de Wit et al., 1999; Domeier and Segal, 2007; Bagher and Segal, 2011) supports the concept that a propagating Ca²⁺ signal is capable of acting through IK/SK channels, which are not present in cECs (Longden et al., 2017) but are expressed by transitional and arteriolar ECs (Hannah et al., 2011; Thakore et al., 2021). The generated hyperpolarizing signal is further amplified via Kir channel

activation (Sonkusare et al., 2016) and conveyed through myoendothelial junctions to adjacent SMCs. In the dilation induced by capillary stimulation with PGE₂, inhibition of the Kir2.1 channel with Ba²⁺ had a profound effect, particularly measurable on the propagation of the dilation, which is consistent with the biphasic propagative model proposed by Thakore et al. (2021). Interestingly, inhibition of TRPA1 channels did not prevent PGE₂ from causing upstream dilation, suggesting a different initiation mechanism, likely involving G_q protein activation and inositol trisphosphate (IP₃)-mediated Ca²⁺ release, as opposed to direct Ca²⁺ entry across the plasma membrane. Our previous work also highlighted the role of G_qPCR activation in breaking down phosphatidylinositol 4,5-bisphosphate (PIP₂), resulting in decreased Kir2.1 channel activity and increased open probability of the Ca²⁺/Na⁺-permeable TRPV4 channel (Harraz et al., 2018a,b, 2020). However, blocking TRPV4 channels had no effect on the dilation induced by PGE₂ either. Given these data, we propose that EP1-initiated IP₃-dependent Ca²⁺ signals arriving in the transitional segment activate endothelial IK/SK channels, and the ensuing membrane potential hyperpolarization activates Kir channels, converting the incoming Ca²⁺ signal into a Kir-dependent hyperpolarizing signal. The characterization of such a Ca²⁺ signal will require more extensive investigation, but the vasodilation induced by capillary stimulation with PGE₂ is clearly central to the postulated role of this molecule as a neurovascular coupling agent.

Cerebral SVDs have emerged as a central link between two major co-morbidities. They account for more than 30% of strokes worldwide and at least 40% of dementia cases (Pantoni, 2010; Iadecola, 2013). CADASIL is caused by dominant mutations in the NOTCH3 receptor, expressed by SMCs and pericytes, that stereotypically lead to the extracellular deposition of the NOTCH3 ectodomain (NOTCH3^{ECD}), which recruits and aggregates other proteins on vessels, ultimately forming deposits termed granular osmiophilic material (GOM) (Joutel et al., 2000, 2016; Chabriat et al., 2009). One of these proteins is the tissue inhibitor of metalloproteinases 3 (TIMP3), which directly complexes with NOTCH^{ECD} and abnormally accumulates in the extra cellular matrix of brain vessels in patients and mice with CADASIL (Monet-Leprêtre et al., 2013). A deficit in CBF hemodynamics, including functional hyperemia, is an early disease manifestation in patients (Chabriat et al., 2000; Pfefferkorn et al., 2001; Liem et al., 2009; Huneau et al., 2018) and a prominent feature of the well-established *TgNotch3*^{R169C} CADASIL mouse model used in the present study (Joutel et al., 2010; Capone et al., 2016; Dabertrand et al., 2021). Our recent work indicates that TIMP3 effects on cerebrovascular reactivity are attributable to inhibition of ADAM17 and subsequent suppression of EGFR signaling by inhibition of ectodomain shedding of its ligand HB-EGF (Dabertrand et al., 2015, 2021; Capone et al., 2016).

Consistent with this model, we previously found that EGFR activation with exogenous soluble HB-EGF restores cerebral arterial tone and functional hyperemia (Dabertrand et al., 2015, 2021; Capone et al., 2016). Here, we

found that PGE₂-induced dilation was impaired in the CADASIL mouse model and fully restored by HB-EGF.

We previously identified two downstream consequences of the suppressed TIMP3-ADAM17-EGFR signaling module: (i) the upregulation of voltage gated K⁺ (K_V1.5) channels in the arteriolar SMCs (Dabertrand et al., 2015; Capone et al., 2016); and (ii) the partial inhibition of Kir2.1 channels in cECs, but not in arteriolar ECs and SMCs (Dabertrand et al., 2021). Here, we report a third consequence: the disruption of PGE₂-induced capillary-to-arteriole signaling, reinforcing the concept that extracellular matrix alterations have profound impacts on cerebrovascular dynamics in SVDs. Using computational modeling, we previously investigated the impact of K_V channel upregulation on membrane potential dynamics in the context of concurrent activation of myocyte Kir channels (Koide et al., 2018). Interestingly, while these analyses showed that a higher K_V channel current density would reduce the membrane potential range over which Kir channels can be activated to cause and propagate dilation, the more hyperpolarized resting membrane potential (9 mV) actually facilitates Kir channel activation. Thus, arterioles from the CADASIL mouse model would still hyperpolarize and dilate in response to Kir channel opening, as observed experimentally (Dabertrand et al., 2015; Koide et al., 2018), but at the cost of a smaller vasodilatory reserve. At the capillary level, we previously described, and modeled, how a 50% reduction in cECs Kir2.1 current is sufficient to completely abolish the capillary-to-arteriole electrical signaling in response to 10 mM K⁺ (Harraz et al., 2018a; Moshkforoush et al., 2020), and then strongly reduces functional hyperemia in the CADASIL mouse model (Dabertrand et al., 2021). We attributed this endothelial dysfunction to a reduced cEC metabolism caused by inhibition of the EGFR pathway (Dabertrand et al., 2021). The resulting lower ATP/ADP ratio in CADASIL compared to WT cECs decreases the synthesis of PIP₂ and its availability to act as an essential cofactor for Kir2.1 channel—hence reducing the channel activity (Huang et al., 1998; Hansen et al., 2011; Harraz et al., 2018a, 2020). Of particular note, PIP₂ is also a substrate for phospholipase C during EP1 signaling, and a reduced pool of PIP₂ would certainly limit the synthesis of the second messenger IP₃, which mobilizes Ca²⁺ from endoplasmic reticulum stores through its action on its cognate receptor. The observation that HB-EGF restores both Kir2.1- and PGE₂-initiated signaling is consistent with this concept. The impact of CADASIL on neurovascular coupling is thus multifaceted, involving disruption of two different propagating signals sharing connections to the EGFR pathway in cECs, and Kir channel activation in the transitional and arteriolar segments.

We have made major progress in establishing potential contributing mechanisms to cerebral hemodynamics impairment observed at an early stage of CADASIL, a Mendelian paradigm of SVDs (Chabriat et al., 2009) and the most common hereditary cause of stroke (Pantoni, 2010) and dementia (Schmidt et al., 2012). Here we demonstrate that functional hyperemia deficits in this SVD involve a PGE₂-initiated capillary-to-arteriole signal that is normally regulated by the TIMP3-ADAM17-EGFR signaling module. Furthermore, the evidence that PGE₂ induces arteriolar dilation via capillary stimulation

has the potential to reconcile disparate findings in neurovascular studies. Our mechanistic studies thus lay the groundwork for novel targeted strategies for treating CADASIL and other cerebrovascular diseases.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author.

ETHICS STATEMENT

The animal study was reviewed and approved by the Institutional Animal Care and Use Committee (IACUC) of the University of Colorado, Anschutz Medical Campus.

AUTHOR CONTRIBUTIONS

AR, JF, and FD performed *ex vivo* experiments, data collection, and analysis. TL and NW performed *in vivo* experiments, data collection, and analysis. AJ contributed to the study design. FD designed and directed the research and wrote the manuscript. All authors edited the manuscript and approved its submission.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnagi.2021.695965/full#supplementary-material>

Supplementary Movie 1 | Capillary stimulation with PGE₂ 1 μM evokes upstream arterial dilation in *ex vivo* CaPA preparation.

Supplementary Movie 2 | Capillary stimulation with PGE₂ causes upstream dilation of the arteriole and transitional segment *in vivo*.

Supplementary Movie 3 | Exogenous application of 30 ng/ml HB-EGF fully restores upstream arterial dilation in response to capillary stimulation with 1 μM PGE₂ in CaPA preparation from a SVD mouse.

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Neurovascular Alterations in Vascular Dementia: Emphasis on Risk Factors

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Vascular dementia (VaD) constitutes the second most prevalent cause of dementia in the world after Alzheimer's disease (AD). VaD regroups heterogeneous neurological conditions in which the decline of cognitive functions, including executive functions, is associated with structural and functional alterations in the cerebral vasculature. Among these cerebrovascular disorders, major stroke, and cerebral small vessel disease (cSVD) constitute the major risk factors for VaD. These conditions alter neurovascular functions leading to blood-brain barrier (BBB) deregulation, neurovascular coupling dysfunction, and inflammation. Accumulation of neurovascular impairments over time underlies the cognitive function decline associated with VaD. Furthermore, several vascular risk factors, such as hypertension, obesity, and diabetes have been shown to exacerbate neurovascular impairments and thus increase VaD prevalence. Importantly, air pollution constitutes an underestimated risk factor that triggers vascular dysfunction via inflammation and oxidative stress. The review summarizes the current knowledge related to the pathological mechanisms linking neurovascular impairments associated with stroke, cSVD, and vascular risk factors with a particular emphasis on air pollution, to VaD etiology and progression. Furthermore, the review discusses the major challenges to fully elucidate the pathobiology of VaD, as well as research directions to outline new therapeutic interventions.

Keywords: vascular dementia (VaD), stroke, cerebral small vessel disease (cSVD), neurovascular abnormalities, blood-brain barrier, neuroinflammation, air pollution

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INTRODUCTION

Dementia affects nearly 50 million people worldwide, and the World Health Organization (WHO) estimates that this number will triple by 2050 (Patterson, 2018). Dementia is a heterogeneous neurodegenerative pathology that encompasses Alzheimer's disease (AD), vascular dementia (VaD), Lewy body dementia (LBD), frontotemporal dementia (FTD), and Parkinson's disease (PD). Although the overall prevalence of dementia is higher in aging men, its severity is more pronounced in aging females, a disparity that might implicate sex hormones (Appelros et al., 2009; Podcasy and Epperson, 2016; Poorthuis et al., 2017). VaD comes just after AD as a main cause of dementia, accounting for approximately 15–20% of dementia cases in the Western countries and could reach up to 30% in Asia and developing countries (Rizzi et al., 2014). Vascular deficiencies are now considered relevant contributors to mixed dementia (MxD), which accounts for 25–35% of all dementia cases (Jellinger, 2007; Rosa et al., 2020).

The Vascular Impairment of Cognition Classification Consensus Study (VICCCS) defines VaD as “clinically significant deficits in at least one cognitive domain comprising sensation, perception, motor skills and construction, attention and concentration, memory, executive functioning, processing speed and language/verbal speed, that are of sufficient severity to cause severe disruption of activities of daily living” (Sachdev et al., 2006; Andrianopoulos et al., 2017; Skrobot et al., 2018). The cognitive functions are assessed through the Montreal Cognitive Assessment Test which evaluates five cognitive domains; executive function, attention, memory, language, and visuospatial function (Pendlebury et al., 2012; Skrobot et al., 2018; Iadecola et al., 2019). Diagnosis of VaD is divided into two major research fields; cognitive tests and neuroimaging. Indeed, the diagnosis does not rely only on memory impairments, but it is supported by the presence of diverse cognitive deficits accompanied by diagnostic imaging evidencing cerebrovascular abnormalities such as brain atrophy, white matter hyperintensities, infarcts, and hemorrhages. Accordingly, four different subtypes arise: post-stroke dementia (PSD) in which dementia appears 6 months after stroke, subcortical ischemic vascular dementia (SIVaD), multi-infarct dementia, and MxD (Skrobot et al., 2018). The cognitive deficits associated with VaD are caused by structural and functional vascular abnormalities that are exacerbated with age. These abnormalities promote the emergence of chronic alterations in the neurovascular functions that underlie the etiology of cognitive decline observed in VaD (**Figure 1**).

VaD is tightly associated with several risk factors that can be categorized into four groups, which comprise: (i) cerebrovascular disease-related factors; (ii) atherosclerotic factors, such as smoking, myocardial infarction, diabetes mellitus, and hyperlipidemia; (iii) demographic factors, such as age, biological sex and education; and (iv) genetic factors, such as the emergence of mutations leading to vascular encephalopathies (Ritchie and Lovestone, 2002; Gorelick, 2004). Noteworthy, these vascular risk factors are now being recognized as clinical risk factors for AD pathology (O'Brien and Markus, 2014; **Figure 1**).

The cerebrovascular disease-related factors include cerebral tissue loss volume, bilateral cerebral infarction, strategic infarction, and white matter disease (WMD; Ritchie and Lovestone, 2002; Gorelick, 2004). Furthermore, hypertension was shown to be associated with larger white matter and smaller brain volumes, silent or strategic subcortical or cortical infarcts, and loss of volume in the thalamus or temporal lobe that are critical for cognitive functions (Ritchie and Lovestone, 2002; Gorelick, 2004). While age remains a principal risk factor for dementia, the presence of familial dementia history, and the epsilon 4 allele of apolipoprotein E (ApoE)4 susceptibility gene were recognized as an important risk factor for VaD (Ritchie and Lovestone, 2002; Gorelick, 2004). In addition, variables such as the female sex, various types of infection of lipid concentrations, history of head injury, head circumference, hormone replacement therapy (HRT), as well as thyroid dysfunction and preceding history of depression could interact with ApoE genotype and hence increase the risk of dementia (Ritchie and Lovestone, 2002; Gorelick, 2004). The

genetic factors include vascular encephalopathies such as cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy (CADASIL), autosomal recessive cerebral arteriopathy with subcortical infarcts and leukoencephalopathy (CARASIL), and potentially ApoE4 (Ritchie and Lovestone, 2002; Gorelick, 2004). Recently, environmental factors, namely air pollution, have been shown to constitute an important, yet underestimated, risk factor for dementia, inducing VaD and AD (Azarpazhooh and Hachinski, 2018; Béjot et al., 2018). Indeed, numerous studies have demonstrated that the elevated levels of air pollutants are directly linked to brain chronic inflammation and neurodegenerative diseases (Campbell et al., 2005; Schwartz et al., 2005; Calderón-Garcidueñas et al., 2007; Hartz et al., 2008; Block and Calderón-Garcidueñas, 2009; Mills et al., 2009; Rozemuller et al., 2012; Paul et al., 2019). Among air pollutants, ultrafine particles (UFPs) are particularly deleterious due to their ability to reach the brain where they act as inflammatory triggers and neurotoxins (Block and Calderón-Garcidueñas, 2009; Hameed et al., 2020).

As mentioned, VaD prevalence is strongly linked to cerebrovascular diseases, which essentially include stroke and cerebral small vessel disease (cSVD; Grinberg and Thal, 2010; Gorelick et al., 2011; Jellinger, 2013). Indeed, one patient in 10 has a stroke before developing a form of dementia that is not related to AD. In turn, cSVD was found in up to 62% of patients diagnosed with VaD, outlining a strong correlation between these pathologies (Gorelick et al., 2011; Venkat et al., 2015; van Veluw et al., 2017; Shih et al., 2018; Iadecola et al., 2019). Stroke and cSVD are characterized by the dysfunction of the neurovascular unit, which anatomically comprises sealed endothelial cells forming the blood-brain barrier (BBB), perivascular cells that include pericytes, and vascular smooth muscle cells (VSMCs), astrocytes, microglia, and neurons (Hermann and ElAli, 2012). The neurovascular unit integrates signals from the different neighboring cells to generate critical functions that include BBB maintenance, neurovascular coupling, vascular stability, and immunomodulation (Zlokovic, 2011; Hermann and ElAli, 2012). Neurovascular functions are impaired after stroke and cSVD leading to BBB dysfunction, neurovascular uncoupling, hypoperfusion, inflammation, and loss of neurons (Zlokovic, 2008, 2011; Guo and Lo, 2009; Moskowitz et al., 2010). These pathological events are at the origin of ischemic and hemorrhagic lesions, which strongly correlate with the cognitive deficits observed in VaD.

Despite being the second most common form of dementia after AD, little is known about the molecular and cellular mechanisms underlying the pathobiology of VaD. This gap in the literature is mainly due to disease heterogeneity in the clinical setup and the lack of an optimal experimental model that can accurately replicate most of the pathological events underlying the etiology and progression of the different forms of VaD. It is now established that accumulation of brain lesions over time mediated by neurovascular impairments constitutes a major contributor to the pathobiology of VaD in the elderly (Venkat et al., 2015; Corrada et al., 2016; Ince et al., 2017; Summers et al., 2017; van Veluw et al., 2017; Shih et al., 2018). The review summarizes the current knowledge related to the

pathological mechanisms underlying the pathobiology of VaD with an emphasis on stroke, cSVD and risk factors with a focus on air pollution. We will discuss the challenges and research directions that might help in a better understanding of VaD pathobiology, thereby outlining new therapeutic interventions.

MAJOR RISK FACTOR-MEDIATED MECHANISMS IMPLICATED IN VAD PATHOBIOLOGY

Stroke-Related Dementia

Ischemic or hemorrhagic strokes trigger major pathophysiological mechanisms that underlay VaD (Mijajlović et al., 2017). Indeed, epidemiological studies indicate that stroke history doubles the risk of dementia in the elderly (<65 years) and increases the incidence of early mortality (Savva and Stephan, 2010). Approximately 10% of patients exhibit signs of dementia before their first cerebrovascular accident, and another 10% manifest cognitive deficits soon after their first event (Desmond et al., 2002; Pendlebury and Rothwell, 2009). Particularly, recurrent stroke events raise the prevalence of dementia to 30%, constituting the most prominent causal factor of the disease (Pendlebury and Rothwell, 2009). Noteworthy, an examination of the association between stroke rates and dementia in the frame of the National Long-Term Care Survey (NLTC) between 1984–2001 reported that the elevated incidence of post-stroke dementia is related to increased patient survival, due to clinical improvements in stroke management (Ukrainitseva et al., 2006). Assessment of the neuropsychological parameters revealed that deterioration of the executive functions, abstraction, visual memory, and visuoconstruction constitute some of the most critical long-term cognitive disabilities observed in stroke patients (Sachdev et al., 2004). In contrast, praxis-gnosis, working memory, and language have been shown to be impacted to a lesser extent (Sachdev et al., 2004). The Sydney Stroke Study disclosed that in 50–85 years old patients diagnosed with VaD, the stroke volume and premorbid function were the most significant determinants of cognitive deterioration following the initial insult (Sachdev et al., 2006). It is remarkable to notice that in the last decade, stroke incidence increased by 23% in young adults aged between 35 and 50 years, especially because of the unhealthy lifestyle that meaningfully increased the rate of risk factors in this population, including smoking, hypertension, and obesity (Ekker et al., 2019; George, 2020). Due to the advances in acute stroke care, young stroke patients live longer and thus are at high risk of developing dementia at later stages (Pinter et al., 2019).

BBB Dysfunction

Hallmark of stroke pathophysiology (Yang et al., 2019), BBB dysfunction constitutes a pivotal factor implicated in the initiation and exacerbation of the cascade of events leading to dementia (Zlokovic, 2011; Sachdev et al., 2014; Noe et al., 2020). Following primary injury, BBB breakdown allows the uncontrolled infiltration into the brain of blood-borne molecules, including plasma proteins, metabolites, neurotoxic compounds, and peripheral immune cells that

contribute to secondary injury progression *via* edema formation, neuroinflammation, and glial reactivity (Halder and Milner, 2019; Koizumi et al., 2019) that aggravate the initial neurological deficits (Khanna et al., 2014; Jiang et al., 2018). The experimental findings indicate that acute BBB impairment is widely mediated by early inflammatory mediators, such as cytokines and chemokines, as well as oxidative stress, including reactive oxygen species (ROS) and reactive nitrosative species (RNS; Yang et al., 2019). The action of these substrates is further potentiated by matricellular proteins, proteoglycans, and metalloproteinases (MMPs) secreted in the extracellular space (Jones and Bouvier, 2014). Experimental and clinical studies revealed that MMP-9 significantly contributes to long-term BBB breakdown in several brain disorders, namely stroke and neurodegenerative diseases (Barr et al., 2010; Montagne et al., 2017; Underly et al., 2017; **Figure 2**).

Upon ischemic stroke, MMP-9 is secreted by the cells forming the neurovascular unit *via* regulation of the extracellular signal-regulated kinase (ERK)-1/2 and the signal transducer and activator of transcription (STAT)-3 pathways, leading to the degradation of basal lamina/extracellular matrix (ECM) proteins, and the recruitment and extravasation of peripheral immune cells (Nishikawa et al., 2018; Jäkel et al., 2020). Interestingly, human brain studies showed that MMP-9 is implicated in the degradation of type IV collagen at the basal lamina, resulting in hemorrhagic transformations (Rosell et al., 2006, 2008), enhanced leukocyte infiltration, and poor neurological outcomes (Kim et al., 2016). In parallel, studies employing MMP-9^{-/-} mice showed that leukocyte-derived MMP-9 plays an essential role in mediating BBB dysfunction and is associated with elevated leukocyte transmigration that exacerbates the inflammatory signaling in the acute phase of stroke (Gidday et al., 2005). Furthermore, photothrombotic mouse models of cerebral ischemia have reported that BBB permeability at the level of the capillary is governed by pericytes exhibiting MMP-9 activation, which was later neutralized by specific MMP-9 inhibition (Underly et al., 2017). Elevated MMP-9 activity has been also reported to be associated with increased brain edema and IgG extravasation after ischemia in hyperlipidemic mice (ElAli et al., 2011). These observations indicate that hyperlipidemia exacerbates stroke-mediated BBB dysfunction, which could eventually aggravate dementia (**Figure 2**).

On an important note, stroke-induced MMP-9 expression has risen as an informative prognostic marker for a poor neurological outcome, increased mortality, and the emergence of typical signs of dementia (Zhong et al., 2017). Clinical investigations disclosed that high MMPs expression correlates with increased levels of albumin cerebrospinal fluid (CSF) in patients with vascular cognitive impairment (VCI) derived from SIVD, multiple strokes, and leukoaraiosis (Candelario-Jalil et al., 2011). In parallel, independently of the presence of vascular risk factors, elevated serum MMP-9 levels are associated with mild (25.6% of patients) and severe (27.4%) cognitive impairment 3 months following stroke according to the Mini-Mental State Examination and Montreal Cognitive Assessment (Zhong et al., 2018). Moreover, high MMP-9 levels in patients with cardioembolic

stroke involving the middle cerebral artery (MCA) territory are related to large infarct volumes and poor behavioral scores based on the National Institutes of Health Stroke Scale (NIHSS; Montaner et al., 2001). Otherwise, the strong correlation between MMP-9 expression and the hyperintense acute reperfusion injury marker (HARM) has led to consider this protein as a revealing marker for BBB disruption (Barr et al., 2010). Furthermore, increased MMP-9 activity has been detected in the frontal and parietal cortex of postmortem human brains diagnosed with AD and patients exhibiting cognitive deficits (Bruno et al., 2009). MMP-9 enhanced expression is also notable in the CSF of AD patients, directly correlated with T-tau and P-tau levels (Stomrud et al., 2010). Importantly, a longitudinal study (4–10 years) enrolling AD and VaD patients revealed higher MMP-9 levels in the CSF of VaD patients compared to AD or controls (Adair et al., 2004). These findings imply that assessment of MMP-9 levels might constitute a potential strategy to distinguish between the different types of dementia.

Cell-based assays have demonstrated that exposure of pericytes to amyloid- β (A β)₄₂ induced MMP-9 activity, which in turn ameliorated protein aggregation (Schultz et al., 2014). This outlines the presence of a direct pathological link between the molecular markers of dementia and MMP-9 activity, even though MMP-9 is an active substrate for A β degradation (Hernandez-Guillamon et al., 2015). Overall, current evidence suggests that MMP-9-mediated BBB dysfunction following stroke may constitute an early pathological mechanism that initiates the neurodegenerative cascades leading to cognitive deficits over time. Although MMP-9 inhibition has been proposed as a therapeutic strategy to attenuate BBB breakdown after cerebral ischemia (Dong et al., 2009; Chaturvedi and Kaczmarek, 2014), animal studies showed that MMP-9 is required for neurovascular remodeling and adaptation in the chronic phase after stroke (Zhao et al., 2006). Furthermore, MMP-9 is implicated in the clearance of various misfolded proteins involved in several neurodegenerative diseases, such as A β (Hernandez-Guillamon et al., 2015). The current knowledge implies that MMP-9 impact on BBB breakdown and neurodegeneration is time and context-dependent, which entails a contextualized modulation of protein expression/activity to preserve BBB integrity and attenuate cognitive deficits associated with stroke.

Post-stroke Neuroinflammation

Evidence obtained from AD studies outlined an important pathological link among chronic neuroinflammation, vascular damage, and cognitive decline in aged patients (Rhodin and Thomas, 2001; Kinney et al., 2018). Importantly, the neuroinflammatory responses associated with AD could be observed as well in other forms of dementia, including frontotemporal dementia (FTD; Bevan-Jones et al., 2020), PD (Caggiu et al., 2019), and VaD (Iadecola, 2013). Neuroinflammation plays a critical role in modulating tissue injury and repair after stroke. This process integrates various molecular and cellular mechanisms that comprise the release of inflammatory and oxidative stress mediators, glial reactivity, and peripheral immune cell activation and extravasation (Guruswamy and ElAli, 2017; Dzyubenko et al., 2018; Jayaraj

et al., 2019). In this regard, uncontrolled microglial activation and the subsequent release of proinflammatory cytokines after stroke (Zhao et al., 2017) are strongly associated with demyelination and axonal loss (Sachdev et al., 2004).

It has been reported that in a rodent model of cerebral hypoperfusion, microglial activation *via* the complement (C)3-C3aR pathway, which is implicated in myelin phagocytosis, resulted in learning and memory deficits (Zhang et al., 2020). Interestingly, the cognitive impairments were attenuated by the genetic deletion of *c3ar1* or *via* the administration of SB290157, a potent C3aR antagonist (Zhang et al., 2020). Similarly, experimental investigations using cerebral ischemia have shown that fingolimod (FTY720), a potent agonist of sphingosine 1 phosphate (S1P), induced a microglial anti-inflammatory phenotype (M2 phenotype) through the activation of STAT-3 signaling pathway (Qin et al., 2017). Modulating microglial activation to adopt a protective M2 phenotype resulted in enhanced oligodendrocytogenesis and white matter integrity, and reduced cognitive deficiencies associated with working memory (Qin et al., 2017). Induction of severe chronic cerebral hypoperfusion (SCCH) in APP/PS1 mice that overproduced A β accelerated spatial learning and memory decline in 4-month adult animals. This was correlated to the accumulation of parenchymal A β plaques in the hippocampus and diminished activity of the ERK-1/2 pathway. APP/PS1 mice subjected to SCCH had higher levels of patrolling monocytes in peripheral blood. Interestingly, this model revealed that SCCH reduces microglial interaction with A β plaques in the hippocampus, denoting a reduced capacity for A β clearing in the brain parenchyma (Bordeleau et al., 2016). Alternatively, the release of prostaglandins, which act as inflammatory mediators upon stroke, has been shown to be associated with exacerbated A β -mediated cognitive decline and impaired synaptic plasticity (Kotilinek et al., 2008; **Figure 2**).

Loss of white matter integrity by hyper-reactive ramified and amoeboid microglia was also found in the posterior cingulate cortex of post-mortem brains of patients diagnosed with Down syndrome who are at higher risk of developing AD neuropathology (Martini et al., 2020). Furthermore, microglia in post-mortem AD brains exhibit accelerated aging and transcriptional alterations associated with the isoforms of ApoE, a protein broadly related to both dementia and cardiovascular disease (Srinivasan et al., 2020). In this regard, findings from subarachnoid hemorrhage in mice indicate that ApoE mediates protective effects following injury by inducing M1 microglial quiescence (Pang et al., 2018), suggesting that adequate lipid metabolism modulates neuroinflammation. Functional human brain investigations using positron emission tomography (PET) coupled to 11C-PK11195, which is an *in vivo* marker of activated microglia, have unraveled a progressive microglial activation and neuroinflammation, which were correlated with long-term (14 to 16 months) cognitive decline in AD patients (Malpetti et al., 2020). In this regard, activated microglia exhibiting a pro-inflammatory neurotoxic phenotype (M1 phenotype) trigger the activation of pro-inflammatory astrocytes (A1 astrocytes) *via* tumor necrosis factor (TNF)- α , interleukin (IL)-1 α , and C1q cytokines (Liddelow et al., 2017). In turn, A1 reactive

astrocytes exacerbate oligodendrocyte and neuronal death (Liddel et al., 2017).

In line with these findings, an exaggerated astrocyte reactivity has been related to dementia and cognitive decline (Jo et al., 2014; Csipo et al., 2020). The evidence suggests that morbid neuroinflammatory responses maintained by A1 reactive astrocytes could result in mediating brain injury or age-related neurodegeneration and cognitive deficits. For instance, hippocampal astrocytes are susceptible to the upregulation of inflammatory-related genes and pathways, such as C3 and C4b, C-X-C motif chemokine ligand (CXCL)-10, and the peptidase inhibitor serine protease inhibitor A3N (SERPINA3N; Clarke et al., 2018). Interestingly, in a mouse model of familial Danish dementia (FDD), abundant A1 reactive astrocytes were detected in the brain, which correlated with the appearance of cerebral amyloid angiopathy (CAA), a disease characterized by the deposition of A β within the cerebral vasculature and a major risk of VaD. In this context, an increased number of astrocytes was observed in perivascular zones, accompanied by numerous cell branches and enhanced glial fibrillary acid protein (GFAP) expression (Taylor et al., 2020; **Figure 2**).

Several approaches have demonstrated that the usage of anti-inflammatory strategies could attenuate cognitive deficits associated with dementia. For instance, activation of the cannabinoid receptor 2 (CB₂R) using different agonists promoted memory restitution through the reduction of oxidative stress and mitochondrial dysfunction (Jayant and Sharma, 2016). In line with these findings, cell-based assays have shown that CB₂R activation stimulated the microglial release of IL-10, a key anti-inflammatory cytokine, *via* activation of ERK1/2, c-Jun N-terminal kinase (JNK), and mitogen-activated protein kinases (MAPKs) pathways, accompanied by the inhibition of the nuclear factor- κ B (NF- κ B) pathway (Correa et al., 2010). Likewise, administration of the CB₂R agonist paeoniflorin (PF) ameliorated memory and learning deficits in mice, accompanied by induction of M2 cells and the release of anti-inflammatory mediators, such as transforming growth factor (TGF)- β 1, and IL-10, instead of pro-inflammatory ones, such as TNF- α , IL-1 β , and IL-6. This phenotypic switch is driven by the enhanced activity of phosphoinositide-3-kinase (PI3K/AKT) anti-inflammatory pathway and the inhibition of the mammalian target of rapamycin (mTOR)/NF- κ B pro-inflammatory signaling (Luo et al., 2018). Experimental findings indicate that the inhibition of mTOR attenuated cognitive deficits, which were accompanied by a restoration of M1/M2 microglia phenotypic switch following cerebral hypoperfusion (Chen et al., 2016).

Remarkably, in a rodent model of VaD, it has been shown that acupuncture could attenuate inflammation by reducing TNF- α and Toll-like receptor (TLR)4 expression in microglia, and suppressing the myeloid differentiation factor (MyD88)/NF- κ B pathway (Wang et al., 2020). Furthermore, pharmacological administration of PLX5622, a potent inhibitor of colony stimulating factor-1 receptor (CSF-1R) that is required for microglial cell survival, improved short-term memory in a rodent model of induced hypertension. This effect was accompanied by controlled microglia reactivity and preservation of BBB integrity (Kerkhofs et al., 2020). It has also been demonstrated that

depletion of microglia *via* CSF1R inhibition prevented A β plaque development in the hippocampus of a mouse model of AD (Spangenberg et al., 2019). Attenuation of microglial reactivity *via* blockage of CCL-5 signaling also preserved BBB integrity in the context of systemic inflammation (Haruwaka et al., 2019). Taken together, there is strong evidence associating chronic uncontrolled neuroinflammatory responses after stroke to the emergence of long-term cognitive disabilities and dementia, mediated essentially by microglial reactivity. Therefore, strategies aiming to modulate microglial response by stimulating a protective phenotype might constitute a potential approach to attenuate VaD occurrence after stroke.

Stroke-Mediated Proteinopathies

Accumulation of A β in the brain constitutes a hallmark of AD pathogenesis (Chen et al., 2017). Early studies using human post-mortem brains revealed that amyloid precursor protein (APP) is not implicated exclusively in AD pathology, and its expression is as well induced in the brain after stroke (Cochran et al., 1991; Jendroska et al., 1997). For instance, mutant mice overexpressing APP exhibited a substantial reduction of cerebral blood flow (CBF) accompanied by larger infarcts after stroke, suggesting that APP exacerbated ischemic injury by impairing structural and functional vascular integrity (Zhang et al., 1997). Moreover, it has been shown that cerebral ischemia promotes APP deposition in the lesion core, the perilesional regions, as well as in the white matter areas exhibiting myelin loss (Nihashi et al., 2001; Zhan et al., 2015).

Endothelin (ET)-1 is a powerful vasoconstrictor synthesized by endothelial cells and reactive astrocytes, which have been shown to be implicated in ischemic stroke pathobiology as well as A β deposition. Examination of post-mortem human brains showed strong ET-1 expression in reactive astrocytes surrounding A β plaques (Hung et al., 2015). Furthermore, ET-1 overexpression in the acute phase after stroke has been involved in BBB disruption, glial reactivity, and neuronal death. Indeed, mutant mice exhibiting astrocytic ET-1 overexpression (GET-1 mice) experience severe memory and spatial learning deficits, associated with the upregulation of cleaved caspase-3, TNF- α , and IL-1 β (Thiel et al., 2014; Hung et al., 2015). Using cell-based assays, ET-1 overexpression in reactive astrocytes has been shown to amplify A β production (Hung et al., 2015). A β accretion contributes to the development of cognitive deficits by impairing the receptor for advanced glycation endproducts (RAGE)-mediated A β clearance, which exacerbates inflammation, oxidative stress, and neurodegeneration (Min et al., 2020). These findings indicate that ET-1 upregulation after ischemic stroke is tightly associated with A β production and deposition and has considerable effects on excitotoxicity and BBB integrity. Furthermore, comorbid models of A β toxicity and cerebral ischemia have reported that A β deposition exacerbates ischemic damage. This condition leads to ventricular enlargement and striatal atrophy, morphological alterations in microglia, increased production of inflammatory mediators and enhanced glial communication *via* the gap junction proteins connexin (CX)-43. These observations are particularly important as ventricular enlargement was associated with deposition of

neurofibrillary tangles and A β plaques, directly implicated in the pathogenesis of various forms of dementia (Amtul et al., 2015). Furthermore, evidence indicates that the interactions among lipoprotein-associated triggering receptor expressed on myeloid cells (TREM) 2 and apolipoproteins are involved in modulating microglia-mediated A β phagocytosis (Yeh et al., 2016), thus suggesting that A β clearance is associated with lipid metabolism (**Figure 2**). Likewise, it has been shown that exogenous administration of A β triggers tau phosphorylation and magnifies learning and memory deficits in animals subjected to cerebral ischemia (Song et al., 2013).

Finally, the generation of RNS, including peroxynitrite, revealed by the formation of 3-nitrotyrosine (3-NT), is increased in perivascular astrocytes and microglial cells after A β ₄₂ injection, strongly correlating with BBB leakage (Ryu and McLarnon, 2006). These findings suggest that A β pathology could trigger the release of reactive nitrogen species in astrocytes, directly undermining cerebrovascular integrity. Moreover, it has been demonstrated that A β binding to RAGE induces ROS production leading to loss of the tight junction proteins claudin-5, occludin, and zonula occludens (ZO)-1, as well as deficient endothelial cell function (Carrano et al., 2011). Overall, this bidirectional pathological crosstalk implies that exacerbated cognitive decline strongly emerges when cerebral injury and A β toxicity occur comorbidly.

cSVD

cSVD comprises numerous pathologies impacting cerebral arteries, arterioles, venules, and capillaries, which are associated with diverse pathological and etiological processes (Østergaard et al., 2016; Staszewski et al., 2017; Li et al., 2018; Parkes et al., 2018). Six different types of cSVD are classified according to their etiology (Pantoni, 2010; Li et al., 2018). Atherosclerosis and sporadic and hereditary CAA are the most frequent forms. Recent reports outlined a significant increase in the number of genetic microangiopathies distinct from CAA such as CADASIL or Fabry's disease (Razvi and Bone, 2006; Ballabio et al., 2007; Dichgans, 2007; Hara et al., 2009). Microangiopathies caused by inflammation or mediated by immunity are rare and characterized by the presence of inflammatory cells within the vasculature (Jennette and Falk, 1997), generally caused by mechanisms associated with systemic pathologies. Venous collagenosis is a pathologic thickening of the wall of veins and venules that are located near the lateral ventricles, thus leading to a smaller lumen and sometimes to an occlusion (**Figure 2**). Finally, post-radiation angiopathies are a side effect of cerebral irradiation that appears a few months to years after treatment. These angiopathies mainly affect small vessels of the white matter associated with fibrinoid necrosis, resulting in an increased thickness of the walls accompanied by a reduced diameter, which jointly could lead to a thrombotic occlusion (Dropcho, 1991).

cSVD Associated Parenchymal Pathology

cSVD refers to various and complex pathological and etiological processes. Therefore, the clinical manifestations depend on both the cause of the pathology and the affected brain territory. Among the most common symptoms are stroke-related

manifestations, progressive cognitive deterioration, VaD, gait disturbance, sphincter dysfunction, and psychiatric disorders (van der Flier et al., 2005; Pantoni, 2010; Del Bene et al., 2013; Li et al., 2018; de Laat et al., 2011). cSVD is thought to constitute the major cause of vascular cognitive deficits and are responsible for up to 45% of dementia cases (Shi and Wardlaw, 2016; Li et al., 2018). Cognitive deficits are associated with impaired executive functions, decline in memory and attention, regression in verbal fluency, and delayed recall. These symptoms are accompanied by others that are not specific, including dizziness, trouble sleeping, tinnitus, and hearing loss. Moreover, neuropsychiatric symptoms can be observed, including hallucinations, agitation, depression, anxiety, disinhibition, apathy, irritability, and changes in appetite. Most of these manifestations are often accompanied by brain microbleeds. Cerebral microangiopathies are accountable for up to 20–30% of ischemic stroke as well as a considerable proportion of hemorrhage and encephalopathies caused by emboli, thrombosis, or stenosis of the vessel (Cai et al., 2015; Shi and Wardlaw, 2016; Regenhardt et al., 2019).

In addition to the cerebrovascular pathologies, cSVD exhibits a unique parenchymal pathology characterized by small subcortical infarcts, lacunar stroke, microbleeds, white matter hyperintensities (WMH), enlarged perivascular spaces, and brain atrophy detectable in imaging (van der Flier et al., 2005; de Laat et al., 2011; Del Bene et al., 2013; Wardlaw et al., 2013; Li et al., 2018; Regenhardt et al., 2018; Das et al., 2019). Small subcortical ischemic stroke is the result of severe tissue ischemia caused by the occlusion of a perforating arteriole. Patients either have typical stroke symptoms or a lesion visible only using neuroimaging approaches (Wardlaw et al., 2013; Li et al., 2018). Lesions can be anywhere in the brain and are round or ovoid and less than 20 mm in diameter (Smith et al., 2012; Brundel et al., 2012; van Veluw et al., 2017; Hartmann et al., 2018). They appear hyperintense in a diffusion-weighted image (DWI), hypointense on the map of apparent diffusion coefficients, and normal to hyperintense in fluid-attenuated inversion recovery (FLAIR)/T2 imaging (Okazaki et al., 2015; Potter et al., 2015; Li et al., 2018). DWI is the most sensitive technique currently used to detect ischemia a few hours after stroke onset. Recent infarcts will form a cavity characterized by morphological changes that include a reduction in volume and diameter within 90 days of the onset of the infarction (Moreau et al., 2012; Potter et al., 2015; Li et al., 2018). These infarcts can evolve in three different ways, namely lacuna, WMH without cavitation T2-weighted sequence, and finally, they could disappear without visible consequences in conventional magnetic resonance imaging (MRI). When a recent small subcortical ischemic stroke resolves into lacuna, it actually forms a fluid-filled cavity called a lacunar stroke and represents 40% of acute ischemic strokes. Lacunar insults are divided in two different categories; cavitated old infarcts and incomplete infarcts (Fisher, 1965; Lammie et al., 1998; Regenhardt et al., 2018). Old infarcts are pan-necrotic cavitation with scattered macrophages whereas incomplete infarcts are described as exhibiting loss of neurons and oligodendrocytes associated with invading CD68⁺ macrophages and reactive microglia in addition to reactive astrocytes that are found inside and around the lesion site

(Merino and Hachinski, 2000; Brundel et al., 2012; Regenhardt et al., 2018).

The vascular damage can also develop into BBB leakage or cerebral microbleeds, which appear as small, round, and homogeneous hypointense foci on T2-weighted MRI and are mostly asymptomatic. They originate essentially from the rupture of a precapillary arteriole and are usually associated with vascular risk factor exposition or vascular A β deposition (Cordonnier et al., 2007; De Silva and Faraci, 2016; Shi and Wardlaw, 2016; Toth et al., 2017). The rupture is caused by various factors such as age, hypertension, cerebral ischemia, dementia, and cerebral amyloid angiopathy (CAA), and participates in cognitive deficits, dementia, and transient neurological deficits (Martinez-Ramirez et al., 2014; Shi and Wardlaw, 2016; Li et al., 2018). Microbleeds trigger proliferation and migration of microglia and astrocytes as well as monocyte recruitment (Liddel et al., 2017). The immune cells release various inflammatory factors that impair neuronal function, as well as neurotransmitters that may be neurotoxic and interfere with neuronal circuitry to promote cognitive decline (Tancredi et al., 2000; Beattie et al., 2002; Rosidi et al., 2011; Donzis and Tronson, 2014).

The cerebral white matter is composed of myelinated axons, myelinating oligodendrocytes, oligodendrocyte precursor cells (OPCs), astrocytes, and microglia (Hase et al., 2018). WMH is common in older people and is a typical feature of cerebral microangiopathies, which are associated with BBB disruption, small white matter infarcts, glial activation, loss of oligodendrocytes, and demyelination caused by chronic diffuse hypoperfusion associated with a reduced CBF (Prins and Scheltens, 2015; Li et al., 2018). WMH is generally located within the white matter including the pons and brainstem but also in the deep gray matter. It is distributed symmetrically and bilaterally and appears hyperintense on FLAIR or T2 MRI. Importantly, WMH triples the risk of stroke, doubles the risk of dementia, and substantially increases the risk of death (Debetto and Markus, 2010; Pantoni, 2010; Shi and Wardlaw, 2016). Symptoms develop insidiously and are associated essentially with cognitive impairments, dementia, and depression (Debetto and Markus, 2010; Pantoni, 2010; Shi and Wardlaw, 2016).

The perivascular space is an extension of the subarachnoid space that surrounds the brain microvasculature. It is a liquid-filled space that cannot be detected by conventional imaging in a physiological context. When this space is widened, it often appears hyperintense on T2 MRI, hypointensity on T1 weighting, and sometimes hypointense on FLAIR (Aribisala et al., 2013; Shi and Wardlaw, 2016; Li et al., 2018). Finally, brain atrophy refers to a diminished brain volume on neuroimaging characterized by symmetrical or asymmetrical decreased total volume, increased ventricular volumes, enlarged superficial sulci, and decreased specific gray or white matter volumes (Mok et al., 2011). One main region affected is the hippocampus and is associated with cognitive decline (Muller et al., 2011; Jokinen et al., 2012).

Atherosclerosis

Atherosclerosis is an age-related condition that constitutes a major risk factor for cerebral microangiopathies. As its severity

is increased by diabetes and hypertension, it is also called hypertensive microangiopathy (Tan et al., 2017; Li et al., 2018; Ter Telgte et al., 2018). The risk factors for atherosclerosis are also hyperlipidemia, smoking, and moderate to severe sleep apnea (Østergaard et al., 2016; Cannistraro et al., 2019). Atherosclerosis is characterized by a chronic inflammation associated with the deposition of low-density lipoproteins (LDL) within the vasculature, leading to its internalization by endothelial cells (Tabas et al., 2015), and resulting in the thickening and hardening of the arterial walls (Lusis, 2000; Shabir et al., 2018). Upon deposition, LDL undergoes oxidation by ROS to form oxidized (ox)-LDL, which further exacerbates the inflammatory response within the vasculature (Tabas et al., 2007). Indeed, by binding to vascular cell adhesion molecule (VCAM)-1 and P-selectin, monocytes can infiltrate the intima and differentiate into macrophages to engulf ox-LDL (Chistiakov et al., 2016). Macrophages, which are now called foam-cells due to the intracellular accumulation of lipids (Spann et al., 2012), accumulate and form stable fatty-streaks into the intima, and cells can calcify over time to slowly occlude the vessel (Alexander and Owens, 2012; Chistiakov et al., 2017). These pathological events occur in large to medium size arteries and lead to microbleeds, microinfarcts, as well as lipohyalinosis, characterized by the deposition of hyaline into the walls of connective tissue (Gorelick et al., 2011). This aspect is specific to the brain due to inflammation caused by ROS, ox-LDL, and gliosis involving astrocytes, OPCs, and microglia (Caplan, 2015). Lipohyalinosis fosters the infiltration of monocytes and T helper (TH)-1 lymphocytes that amplify the production of inflammatory mediators, such as TNF- α and interferon (INF)- γ (Stemme et al., 1995; Frostegård et al., 1999). Moreover, the assembly of inflammasomes can be promoted through the activation of nucleotide-binding oligomerization domain (NOD)-like receptor protein (NLRP)-3, stimulated by the formation of cholesterol crystals, caspase-1 and apoptosis-associated speck-like protein containing (ASC), caspase activation and recruitment domain (CARD; Weber and Noels, 2011). This results in IL-1 β release, which in turn stimulates the release of IL-6 and C-reactive protein (CRP), implicated in the pathogenesis of atherosclerosis and thrombosis (Ridker et al., 2017; **Figure 2**).

Besides, the elevated levels of LDL combined with the low levels of high-density lipoprotein (HDL) play an important role in the pathogenesis of atherosclerosis and constitute as well a major risk factor for VaD (Hao and Friedman, 2014; Georgakis et al., 2020). Indeed, HDL exerts a protective role through its antioxidant properties and mediates beneficial effects on platelets and endothelial function, thus on coagulation and inflammation (Bandeali and Farmer, 2012). Furthermore, HDL interacts with triglyceride-rich lipoproteins, attenuating their deferential effects (Bandeali and Farmer, 2012). Moreover, it could contribute to the removal of cholesterol excess from the brain microvasculature through ApoE and heparin sulfate proteoglycans (Mulder and Terwel, 1998). HDL counteracts the inhibition of vessel relaxation caused by ox-LDL and decreases LDL peroxidation which affects cellular function and impairs membrane-bound receptors and enzymes (Braucher

and Hall, 1992; Matsuda et al., 1993). Importantly, the impact of hyperlipidemia seems to be differentially modulated depending upon biological sex. Indeed, a recent study shows that females exhibit greater expression of genes related to neuroprotection in response to lipid stress compared to age-matching males (Nuthikattu et al., 2020). Finally, lipid derivatives are now being under the scope of researchers who are trying to unravel novel biomarkers to better understand and diagnose VaD pathology. Indeed, in an interesting recent study that aimed to discover lipid biomarkers in the context of VaD, it has been reported that patients with dementia exhibit low levels for ceramides, cholesterol esters, and phospholipids, and high levels of glycerides compared to controls (Liu et al., 2020). These observations indicate that lipid derivatives could indeed be used as novel diagnostic and prognostic biomarkers in VaD. However, more research is needed in this direction to validate the use of lipid derivatives as diagnostic and prognostic biomarkers in VaD.

Vascular damage caused by atherosclerosis can lead to microatheromas, microaneurysms, and even stenosis or obstruction of the vessel, impairing the mechanisms of blood flow autoregulation and leading to chronic cerebral hypoperfusion (Pantoni, 2010; Li et al., 2018). Importantly, occlusion of the cerebral arteries results in local ischemia or lacunar infarction (Kraft et al., 2017; Ter Telgte et al., 2018), while stenosis and hypoperfusion in the white matter cause incomplete ischemia lesions evidenced by neuroimaging as white matter hyperintensity (Rigsby et al., 2007). When the pathology affects the cerebral arterioles $<50\ \mu\text{m}$ in diameter, it is called small cerebrovascular atherosclerosis (Li et al., 2018).

Sporadic and Hereditary CAA

CAA is a chronic degenerative disease characterized by the loss of VSMCs and the accumulation into the vessel wall of eosinophilic hyaline material composed of soluble $\text{A}\beta_{40}$ (Attems et al., 2011; Charidimou et al., 2017). CAA affects 50–60% of the elderly population affected by dementia, including 85–95% of AD cases (Sacco, 2000; Jellinger, 2002; Thal et al., 2003; Keage et al., 2009; Charidimou et al., 2017; Zhang et al., 2017; Li et al., 2018). The initial cognitive deficits associated with VaD could be explained by the particular sensitivity of the hippocampus and the cortex to CAA (Arvanitakis et al., 2011; Li et al., 2018). CAA is associated with changes in basal membrane (BM) composition and morphology that could predispose $\text{A}\beta$ accumulation in the vessel even though the mechanism is not yet fully understood (Perlmutter et al., 1990, 1991; Su et al., 1992; Morris et al., 2014; Howe et al., 2020). Among the reported changes are BM thickening and degeneration, abnormal heparan sulfate proteoglycans (HSPGs) deposits, and irregular vasculature accompanied by increased collagen IV, fibronectin, agrin, and perlecan expression (Berzin et al., 2000; Farkas et al., 2000; Bourasset et al., 2009; Gama Sosa et al., 2010; Keable et al., 2016; Lepelletier et al., 2017; Magaki et al., 2018; Singh-Bains et al., 2019). Furthermore, vascular functional impairments are featured by BBB dysfunction caused by loss of endothelial cells, deregulation of mural cells mediated by oligomeric $\text{A}\beta$ accumulation, as well as induction of astrogliosis with dystrophic endfeet surrounding BM $\text{A}\beta$ deposits (Shimizu et al., 2009; de

Jager et al., 2013; Giannoni et al., 2016; Yang et al., 2017; Magaki et al., 2018; Nortley et al., 2019). Two possible mechanisms for $\text{A}\beta$ deposition have been proposed: (i) release of vascular $\text{A}\beta$ from VSMCs directly into the vessel wall; and (ii) release of parenchymal $\text{A}\beta$ by neurons which deposits afterward into the vessel wall (Davis et al., 2004; Herzig et al., 2004; Vidal et al., 2009; ElAli et al., 2013). In both cases, the protein accumulates due to a poor clearance towards the periphery (Davis et al., 2004; Herzig et al., 2004; Vidal et al., 2009; ElAli et al., 2013). Insufficient $\text{A}\beta$ clearance can impair perivascular drainage pathways or diminish the ATP binding cassette subfamily B member-1 (ABCB1) and low-density lipoprotein receptor-related protein (LRP)1, a specialized endothelial-mediated active transport system implicated in $\text{A}\beta$ mobilization from the brain into the blood circulation, namely (Deane et al., 2004; Herzig et al., 2006; Weller et al., 2008; Hawkes et al., 2011). Interestingly, LRP1 plays an important role in protecting against neurodegeneration. Indeed, LRP1 downregulation doesn't only affect $\text{A}\beta$ clearance but causes as well BBB breakdown through activation of MMP-9, thus leading to loss of neurons and cognitive deficits (Nikolakopoulou et al., 2021). Furthermore, recent evidence reveals that vascular $\text{A}\beta$ could be engulfed and eliminated by circulating patrolling monocytes, which act as the housekeeper vascular homeostasis by surveying endothelial cells (Auffray et al., 2007; Carlin et al., 2013; Michaud et al., 2013; Thériault et al., 2015). In this regard, it has been demonstrated that patrolling monocytes located at the luminal wall internalize $\text{A}\beta$ microaggregates that are diffusing from the parenchyma into the blood. Unfortunately, patrolling monocyte ability to phagocyte vascular $\text{A}\beta$ in AD is defective, resulting in an overall increase of highly toxic $\text{A}\beta_{40}$ and $\text{A}\beta_{42}$ oligomers (Hallé et al., 2015; Gu et al., 2016). Moreover, chronic mild cerebral hypoperfusion impairs BBB functional properties and promotes the accumulation of circulating $\text{A}\beta$ into the vessel wall, which initiates the cascade of parenchymal $\text{A}\beta$ deposition (ElAli et al., 2013). $\text{A}\beta$ accumulation and BM rearrangement trigger BBB breakdown, endorsing the formation of perivascular edema and the infiltration of toxic blood-derived substrates into the brain, which in turn contribute to the exacerbation of localized injuries and enlargement of the perivascular space (Holland et al., 2008; Hartz et al., 2012; Wardlaw et al., 2013; Li et al., 2018; **Figure 2**).

The overwhelming evidence is suggesting that the here mentioned vascular abnormalities leading to dementia reported in CAA occur as well in different forms of dementia, including AD and LBD (Salat et al., 2006; Okamoto et al., 2010; Soontornniyomkij et al., 2010; Arvanitakis et al., 2011; Love et al., 2014; Martinez-Ramirez et al., 2014; Reijmer et al., 2016; Li et al., 2018). This form of cSVD can be sporadic or of a genetic origin. For instance, a syndrome called hereditary brain hemorrhage with amyloidosis (HBHA) is associated with a mutation in the *APP* gene. This syndrome results in the deposition of misfolded amyloid fibrils in the walls of cerebral arterioles, which in turn activates a cascade of events leading to the development of CAA. The clinical phenotype develops between the ages of 45–65 years and is associated with intracerebral hemorrhages, WMH, multifocal lesions of a hemorrhagic and ischemic nature (Kamp et al., 2014; Marini et al., 2020). The presence of *ApoE4*

allele, which constitutes the main risk factor for AD, has also been demonstrated to constitute an important risk factor for this form of cSVD (Hermann and ElAli, 2012). ApoE4 is a lipid-binding protein which plays an important role in lipoprotein metabolism as well as transport of triglycerides and cholesterol (Hirsch-Reinshagen et al., 2009). It binds to LDL, very-low-density lipoprotein (VLDL) debris, and some HDL *via* LRP (Bu, 2009; Leduc et al., 2010). ApoE4 can form complexes with A β and impairs A β through LRP thus attenuating its clearance and subsequently leading to its accumulation in the brain (Cho et al., 2001; Verghese et al., 2013). Moreover, ApoE4 increases the formation of A β oligomers, which are now well established to constitute the most neurotoxic form of A β (Hashimoto et al., 2012; Youmans et al., 2012). Interestingly, human pericytes of the prefrontal cortex and hippocampus of ApoE4 carriers exhibit increased activation of nuclear factor of activated T-cells (NFAT), which might account for CAA occurrence (Blanchard et al., 2020; **Figure 2**).

Moreover, non-APP sources of CAA exist and are caused by mutations of the BRI2 [i.e., integral membrane protein (ITM)2B] gene, essentially a codon stop mutation. Indeed, processing of the mutated form of BRI2 protein leads to the generation of 34-mer amyloid Bri (ABri) and amyloid Dan (ADan) peptides that accumulate in the brain, either to the ABri amyloid subunit or the AD amyloidogenic fragment. ABri and ADan are responsible for the Familial British (FBD) and Danish (FDD) dementia characterized among other pathological features by severe CAA (Vidal et al., 1999, 2000; Yamada and Naiki, 2012).

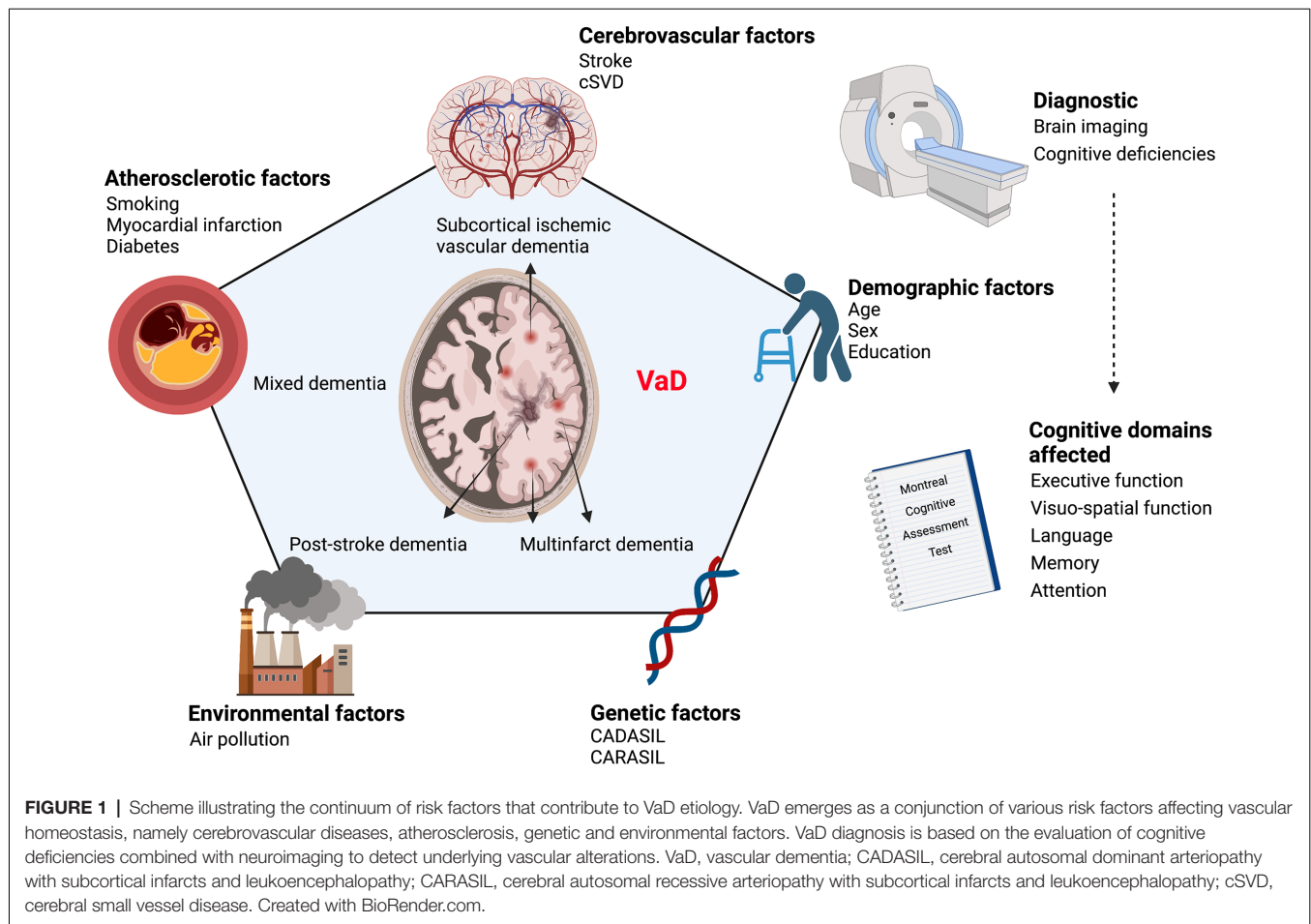
Genetic cSVD

Mutations in specific genes constitute the third most common cause of cSVD, among which the mutation of the *neurogenic locus notch homolog protein (NOTCH)3* gene is the better characterized (Cannistraro et al., 2019; Marini et al., 2020). NOTCH3 is a member of the transmembrane receptor NOTCH family, which is critically involved in developmental patterning, cell fate decisions, regulation of cell survival, and proliferation (Kopan and Ilagan, 2009; Bray, 2016; Baron, 2017; Hosseini-Alghaderi and Baron, 2020). During adulthood, NOTCH3 regulates stem cells and their lineages to promote tissue maintenance and repair. NOTCH3 is expressed by VSMCs and pericytes and plays a key role in regulating the crosstalk between the mural and endothelial cells. It controls the vascular tone and flow-mediated dilation *via* the modulation of the Ras homolog family member A (RHOA)/ Rho-associated protein kinase (ROCK) pathway in cerebral arteries (Joutel et al., 2000; Belin de Chantemèle et al., 2008; Li et al., 2009; Marini et al., 2020). However, the role of NOTCH3 is not restricted to the vasculature, since it is expressed in neural stem cells and is implicated in neuronal differentiation (Alunni et al., 2013; Kawai et al., 2017).

A mutation in the *NOTCH 3* gene is responsible for CADASIL, the most common autosomal dominant inherited cSVD (Louvi et al., 2006; Di Donato et al., 2017; Hosseini-Alghaderi and Baron, 2020; Marini et al., 2020). *NOTCH3* gene is affected essentially by missense mutations that lead to an odd number of cysteine residues located in the extracellular domain of the encoded receptor, and is associated with an early

accumulation of the receptor's extracellular domain containing aggregates in small vessels (Joutel et al., 2000, 2001; Monet-Leprêtre et al., 2013; Yamamoto et al., 2013). The function and activity of the NOTCH3 receptor are differently impacted by the mutations. However, the accumulation of extracellular domain containing aggregates in small vessels leads to mural cell degeneration *via* apoptosis or impaired proliferation (Joutel et al., 2000, 2001; Monet-Leprêtre et al., 2013; Yamamoto et al., 2013). Furthermore, the mutation itself causes profound morphological changes in pericytes, associated with dysfunctional mitochondria that could lead to oxidative and phosphorylation deficiencies, secondary lysosomes, and large cytoplasmic vesicles that result in cellular injury and autophagic apoptosis (de la Peña et al., 2001; Gu et al., 2012). This cascade of events cause neurovascular unit dysfunction characterized by detachment of astrocytic endfeet, destabilization of the vasculature, deregulation of vascular contractility, leakage of the BBB, and infiltration of toxic blood-born components into the brain parenchyma due to the decreased endothelial adherens junction protein, thus jointly resulting in diminished reactivity to CO₂ (Ghosh et al., 2015; **Figure 2**).

Pericytes and endothelial cells are intimately interconnected through peg-and-socket junctions, which degenerate upon *NOTCH3* mutations. For instance, endothelial cells exhibit degenerative features, such as selective death or swelling, causing vessel stenosis or occlusion (Dziewulska and Lewandowska, 2012). Deregulation of pericyte-endothelial cells crosstalk causes cerebrovascular dysfunction that comprises reduced vascular density and impaired CBF (Tuominen et al., 2004; Lacombe et al., 2005; Miao et al., 2006; Gu et al., 2012; De Guio et al., 2014; Liu X.-Y. et al., 2015; Ihara and Yamamoto, 2016; Ping et al., 2019). Moreover, chronic cerebral hypoperfusion resulting from cerebrovascular dysfunction exacerbates pericyte degeneration, reduces pericyte coverage for the capillaries, and subsequently increases BBB permeability leading to white matter impairments and neuronal loss (Ueno et al., 2002; Bell et al., 2010; Montagne et al., 2018; Liu et al., 2019; Nikolakopoulou et al., 2019). BBB breakdown allows the infiltration into the parenchyma of toxic blood-born metabolites that accumulate around the vasculature, thus inducing macrophage, microglia, and T-cells activation and recruitment, which jointly promote axonal degeneration (Davalos et al., 2012; Ryu et al., 2015). Neuronal loss is mainly provoked by the secretion of pro-inflammatory mediators and the generation of ROS and RNS by pericytes within the perivascular space, which further exacerbates leukocyte adhesion and infiltration as well as microglial cell activation (Matsumoto et al., 2018; Erdener and Dalkara, 2019). Finally, the structural lesions within the white matter are worsened by the release of pericyte-derived bone morphogenetic protein (BMP)-4, which promotes astrogliosis (Uemura et al., 2018, 2020). Evidence of these pathological events could be detected using imaging approaches that indicate WMH, ischemic manifestations, subcortical hemorrhages, and microbleeds. A new sensitive assay was recently developed allowing pericyte injury detection in the CSF, a new technology that could serve as a diagnostic tool for WMD (Sweeney et al., 2020).



CADASIL develops gradually over time and the earliest symptoms appear on average around 30 years of age, usually 10 years earlier in women than men, and are manifested as migraines with aura (Guey et al., 2016; Di Donato et al., 2017). The migraine could also manifest with atypical attacks with basilar, hemiplegic, or prolonged aura and a few patients can even develop very severe attacks leading to confusion, fever, meningitis, or even coma that can mimic encephalopathy (Schon et al., 2003; Vahedi et al., 2004; Ragno et al., 2013; Tan and Markus, 2016; Drazzyk et al., 2019). Adults between the age of 20 and 65 years are subject to transient ischemic attacks and stroke (Lesnik Oberstein et al., 2001). CADASIL is associated as well with some psychiatric manifestations, which include mood disturbances, severe depression, and schizophrenia (Lågas and Juvonen, 2001; Valenti et al., 2008, 2011; Noh et al., 2014; Ho and Mondry, 2015; Di Donato et al., 2017). Finally, 40% of symptomatic cases report apathy which drastically impacts the quality of life of CADASIL patients (Reyes et al., 2009). The final stage of CADASIL progression is dementia, but cognitive decline starts years before (Brookes et al., 2016).

A recessive form of CADASIL exists under the name of CARASIL. This form of hereditary cSVD is caused by the mutation of the *high-temperature requirement A serine peptidase 1 (HTRA1)* gene which has two major functions, degrading

various substrates and inhibiting TGF- β 1 signaling pathway that is involved in various processes namely angiogenesis and BBB formation *via* pericyte-endothelial cell crosstalk (Oka et al., 2004; Hara et al., 2009; Shiga et al., 2011; Akhtar-Schaefer et al., 2019; Kandasamy et al., 2020). HTRA1 is expressed in various brain cells comprising endothelial cells and VSMCs (De Luca et al., 2003; Oka et al., 2004; Campioni et al., 2010; Tennstaedt et al., 2012; Tiaden and Richards, 2013). Loss of HTRA1 function results in increased TGF- β 1 availability and thereby signaling, leading to vascular fibrosis and extracellular matrix synthesis, which jointly cause microvascular degeneration, CBF reduction, and neurogenesis alterations (Wyss-Coray et al., 2000; Tarkowski et al., 2002; Gaertner et al., 2005; Yamamoto et al., 2011; Zhang et al., 2012; Martinez-Canabal et al., 2013; Beaufort et al., 2014; Friedrich et al., 2015). Moreover, the mutation has been shown to be associated with impaired pericyte proliferation, accumulation of protein within the vessel walls, MMPs activity, and BBB permeability (Joutel et al., 2016; Baron-Menguy et al., 2017; Ikawati et al., 2018; Marini et al., 2020). Cognitive decline begins early compared with CADASIL, as well as gait disturbances, lower back, pain and alopecia (Shiga et al., 2011; Marini et al., 2020).

Fabry's disease is an X-inherited rare disorder that belongs to the family of lysosomal storage diseases and is caused by a

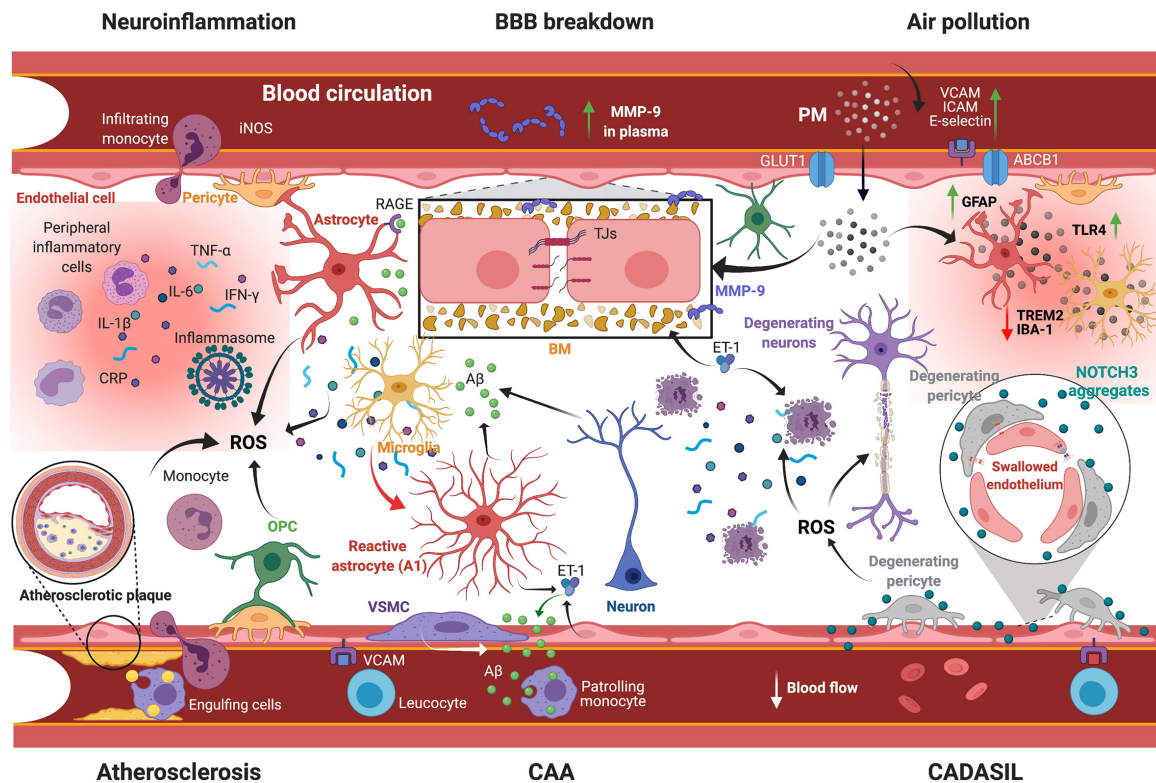


FIGURE 2 | Scheme illustrating the mechanisms underlying VaD pathobiology. Several vascular risk factors are implicated in orchestrating pathological responses leading to VaD: (i) BBB breakdown involves the impairment of TJs and degradation of BM formed by ECM proteins via MMP-9 activity. Plasma MMP-9 levels constitute an effective prognostic marker for a poor neurological outcome; (ii) Post-stroke neuroinflammation comprises extravasation of peripheral immune cells and secretion of inflammatory mediators (e.g., CRP, TNF- α , IL-1 β , IL-6, IFN- γ), as well as ROS generation and glial activation, accompanied by cerebral A β aggregation; (iii) Atherosclerosis comprises the accumulation of lipids and the calcification of immune cells into the intima, leading to vessel occlusion and hypoperfusion. This condition is associated with the generation of ROS that causes chronic inflammation; (iv) CAA is associated with the degeneration of VSMCs and vascular A β aggregation due to impaired clearance; (v) CADASIL is associated with NOTCH3 aggregation, causing endothelial cell swelling and pericyte degeneration and subsequently CBF impairment; and (vi) Exposure to air pollution, which implies PM infiltration into the brain, exacerbates BBB breakdown and neuroinflammation. VaD, vascular dementia; BBB, blood-brain barrier; TJs, tight junctions; BM, basement membrane; ECM, extracellular matrix proteins; MMP-9, matrix metalloproteinase-9; CRP, C-reactive protein; TNF- α , tumor necrosis factor- α ; IL-1 β /6, interleukin-1 β /6; IFN- γ , interferon- γ ; ROS, reactive oxygen species; A β , amyloid- β ; CAA, cerebral amyloid angiopathy; VSMCs, vascular smooth muscle cell; CADASIL, cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy; NOTCH3, neurogenic locus notch homolog protein-3; CBF, cerebral blood flow; PM, particulate matter; GLUT1, glucose transporter-1; ABCB1, ATP binding cassette subfamily B member-1; VCAM, vascular cell adhesion protein; ICAM, intercellular adhesion molecule; TLR4, Toll like receptor-4; TREM2, triggering receptor expressed on myeloid cells-2; GFAP, glial fibrillary acidic protein; RAGE, receptor for advanced glycation endproducts; ET-1, endothelin-1; iNOS, inducible nitric oxide synthase; OPC, oligodendrocyte progenitor cell. Created with BioRender.com.

mutation in the α -galactosidase (*GAL*)A gene that encodes for α -GAL enzyme that plays a key role in sphingolipid metabolism (El-Abassi et al., 2014). The mutation causes deficiencies in α -GAL activity that results in the accumulation of sphingolipids in various organs and tissues including the vessels (Rolfs et al., 2013). The cerebrovascular complications in Fabry's disease arise from peripheral neuropathy and are associated with mild to severe headache, vertigo, transient ischemic attacks, ischemic stroke, intracerebral hemorrhage, and VaD (Okeda and Nisihara, 2008). Furthermore, deposition of toxic metabolites within the vasculature and VSMCs lead to ischemia, vessel stenosis, occlusion, and dilation with local changes in CBF (Shimotori et al., 2008). The disease is more severe in men compared to women and is often present with infantile neuropathy, gastrointestinal symptoms, corneal opacity, hearing loss, and

angiokeratoma (Sims et al., 2009; Schiffmann, 2015; Marini et al., 2020). Brain structural damage and symptoms exacerbate with age.

Collagen IV, which exists as a heterodimer derived from the transcription of *COL4A1* and *COL4A2* genes, is an essential component of the vascular BM. Mutations in these two genes are associated with microangiopathies in several organs (Germain et al., 2019; Marini et al., 2020). More precisely, *COL4A1* mutation is responsible for ocular, renal, muscular, and cerebral deficits (Vahedi and Alamowitch, 2011; Marini et al., 2020). Furthermore, cerebral microangiopathies have been shown to affect half of the carriers of this mutation. WMH, dilation of the perivascular spaces, lacunar infarctions, and microbleeds have been reported as well (Vahedi and Alamowitch, 2011). Pontine autosomal dominant microangiopathy and

leukoencephalopathy (PADMAL) syndrome is a specific form of cerebral microangiopathies associated with the *COL4A1* mutation (Verdura et al., 2016), and is associated with an overexpression of the gene with the absence of protein misfolding. Patients with this syndrome have dysarthria, ataxia, and stroke as well as mood disorders and dementia. On the other hand, *COL4A2* mutation is associated with an increased prevalence of lacunar ischemic stroke and deep intracerebral hemorrhages (Verdura et al., 2016). In contrast to *COL4A1* mutation, *COL4A2* mutation impairs the trimerization of collagen IV due to defects in the α -helix structure, which cause BM instability, loss of vascular wall integrity, and increased BBB permeability, mediated by the intra- and extracellular accumulation of deficient collagen IV (Kuo et al., 2012; Meuwissen et al., 2015; Verdura et al., 2016; Zhang et al., 2017; Malik et al., 2018; Germain et al., 2019).

Forkhead Box C1 (FOXC1) is highly expressed in pericytes. Its expression plays an important role in controlling endothelial cell proliferation and vascular stability. *FOXC1* mutation reproduces some of the events reported upon *COL4A1* mutation, especially the ischemic infarctions, and cerebral microangiopathies that lead to WMH, which are visible in neuroimaging (French et al., 2014). The retinal vasculopathy with cerebral leukodystrophy (RVCL) syndrome includes three pathological conditions: (i) cerebral retinal vasculopathy (CRV); (ii) hereditary vascular retinopathy (HRV); and (iii) hereditary endotheliopathy with retinopathy, nephropathy, and stroke (HERNS). Patients with these syndromes possess a mutation in three prime repair exonuclease (TREX)-1 that encodes for a DNA exonuclease (Stam et al., 2016). This mutation causes a defect in apoptosis and INF signaling (Rice et al., 2015; Marini et al., 2020). All characteristics of the cerebral microangiopathies are found in RVCL patients who report the following symptoms: neurological deficits, migraines, cognitive deficits, psychiatric disorders, and seizures (Stam et al., 2016; Marini et al., 2020).

Environmental Risk Factors: Air Pollution

The increasing interaction with the environmental risk factors associated with human activities has a significant impact on health due to the exposure to various hazardous pollutants. Currently, environmental factors are directly implicated in the etiology and progression of diverse pathologies, including brain diseases. Indoor and outdoor air pollution is among the environmental factors that play a particularly important role in the deterioration of vascular health (Block and Calderón-Garcidueñas, 2009). Indeed, air pollution, which is defined as the release of an amalgam of pollutants into the atmosphere, has been reported to increase the prevalence of cardiovascular, cerebrovascular, and respiratory diseases, as well as cancer (Campbell et al., 2005; Schwartz et al., 2005; Calderón-Garcidueñas et al., 2007; Hartz et al., 2008; Block and Calderón-Garcidueñas, 2009; Mills et al., 2009; Rozemuller et al., 2012; Cho et al., 2018; Paul et al., 2019). Epidemiological studies have indicated that nearly one-third of the global stroke burden and about one-fifth of the global dementia burden, including VaD, are attributable to air pollution (Feigin et al., 2016; Azarpazhooh and Hachinski, 2018; Béjot et al., 2018).

Furthermore, numerous studies have outlined a link between high levels of air pollutants, chronic brain inflammation, and neurodegeneration (Campbell et al., 2005; Schwartz et al., 2005; Calderón-Garcidueñas et al., 2007; Hartz et al., 2008; Block and Calderón-Garcidueñas, 2009; Mills et al., 2009; Rozemuller et al., 2012; Paul et al., 2019). These effects are mainly attributable to the exposure to fine particulate matter (PM), and more precisely to PM of 2.5 microns or less in diameter (PM_{2.5}). Indeed, the experimental findings have indicated that UFPs could reach the brain through different routes, including the intranasal cavity, where they act as an inflammatory mediators, thus deregulating the function of cells forming the neurovascular unit (Oberdörster et al., 2004; Peters et al., 2006). In particular, exposure to PM leads to impaired olfactory function, one of the initial atypical symptoms that emerge in individuals affected by different forms of dementia (Campbell et al., 2005; Schwartz et al., 2005; Calderón-Garcidueñas et al., 2007; Hartz et al., 2008; Block and Calderón-Garcidueñas, 2009; Mills et al., 2009; Rozemuller et al., 2012; Paul et al., 2019). The correlation between air pollution and dementia, including VaD and AD, was highlighted in various epidemiological studies relying mostly on cohort studies in polluted regions (Åström et al., 2021). The Betula cohort revealed an association of dementia incidence, AD in particular, with traffic-related air pollution (TRAP; Oudin et al., 2019). Moreover, studies have found that exposure of the elderly to air pollution, notably PM₁₀ and PM_{2.5} was associated with cognitive decline (Wu et al., 2015). In an interesting case-control study, it was reported that elevated long-term PM₁₀ levels were associated with a significantly increased risk of AD and VaD prevalence in the elderly. A dose-response relationship between PM₁₀ exposure and the risk of AD and VaD was reported (Wu et al., 2015).

Impact on Endothelial Functions

Various studies have investigated the impact of air pollutants, and more specifically PM, on endothelial functions. For instance, exposure of isolated rat brain capillaries to diesel exhaust particles (DEP) altered BBB function through oxidative stress generation and proinflammatory cytokine production, which jointly induced the expression of adhesion molecules that exacerbate infiltration of immune cells into the brain (Hartz et al., 2008). Exposure to DEP deregulated the expression of several transporters and receptors that are critically involved in BBB functionality namely ABCB1, multidrug-resistance associated proteins (MRP)1, MRP2, MRP4, breast cancer resistance protein (BCRP), glucose transporter (GLUT)1, and the metabolizing enzyme glutathione S-transferase (GST) π (Hartz et al., 2008). Several reports have demonstrated an increased BBB permeability following exposure to a mixed vehicular emission, translated essentially by deregulation of the TJs (Rojas et al., 2011; Oppenheim et al., 2013; Bernardi et al., 2021). For instance, human brain microvascular endothelial cells in culture exposed to nanoparticles of aluminum oxide exhibit reduced cell viability, altered mitochondrial function, increased oxidative stress, and diminished expression of the TJs proteins claudin-5 and occludin (Chen et al., 2008; Block and Calderón-Garcidueñas, 2009). Interestingly, epidemiological studies have outlined a strong correlation between air pollution

and neuroinflammation in highly exposed residents (Calderón-Garcidueñas et al., 2008). Indeed, these studies reported an upregulation of some inflammatory markers, such as expression of cyclooxygenase (COX)-2 and IL-1 β , as well as infiltration of immune cells into the olfactory bulb (OB), frontal cortex, substantia nigrae, and vagus nerves (Calderón-Garcidueñas et al., 2008). Importantly, most of the inflammatory responses were concentrated at the vasculature translated by activation of NF- κ B pathway in brain endothelial cells, accompanied by oxidative stress, A β ₄₂ immunoreactivity, trafficking of inflammatory cells into the perivascular space, and an altered BBB (Calderón-Garcidueñas et al., 2008). In line with these observations, direct exposure of brain endothelial cells in culture to PM_{2.5} deregulated TJs and increased permeability and monocyte transmigration across the endothelial monolayer (Liu F. et al., 2015). In addition, exposure to both PM_{2.5} and PM₁₀ induced the activation of endothelial cells accompanied by an enhanced adhesion of U937 monocytic cells to the endothelial monolayer (Montiel-Dávalos et al., 2007). Interestingly, exposure to PM_{2.5} also induced ICAM-1 expression, whereas exposure to PM₁₀ induced expression of E-selectin and P-selectin (Montiel-Dávalos et al., 2007; **Figure 2**).

In vivo experiments in which mice were exposed to a mixed vehicular emission, a combination of gasoline and diesel engine exhausts, the animals exhibited altered BBB integrity through the deregulation of the TJs protein, namely claudin-5 and occludin (Oppenheim et al., 2013). This was accompanied by an augmentation of inducible nitric oxide synthase (iNOS) levels, an increase in the production of IL-1 β in the parenchyma, and deregulation of ABCB1 transport activity (Oppenheim et al., 2013). Moreover, exposure to PM *in vitro* and *in vivo* has been shown to stimulate the re-localization of the TJs protein ZO-1 from the cell membrane and reduce its protein level (Wang et al., 2012). Importantly, PM mediated the intracellular mobilization of calcium (Ca²⁺) dependently upon ROS, activating calpain that is implicated in ZO-1 degradation and disruption of the endothelial barrier (Wang et al., 2012). Using a 3D human *in vitro* BBB model, indoor nanoscale particulate matter (INPM) was shown to translocate across the BBB and accentuate inflammation by inducing ROS (Li et al., 2020). This induction was followed by abnormal nuclear reactor factor (NRF)-2 expression and a disruption of the kelch ECH associating protein (KEAP)-1/antioxidant response elements (ARE) pathway which is involved in supporting cells to overcome stress (Li et al., 2020). Interestingly, exposure of rodents to urban PM increased the levels of ET-1 mRNA and reduced TNF- α mRNA levels in the cerebral hemisphere and the pituitary gland. These results suggest that the cerebrovascular effects of urban pollutants are associated with the modulation of gene expression involved in the regulation of vasoconstriction in the brain and pituitary gland (Benatti et al., 1993; Thomson et al., 2007).

Exposure to PM_{2.5} has been shown to increase the prevalence of carotid artery stenosis (CAS), a well-established risk factor for ischemic stroke, correlating with an increased BBB permeability (Newman et al., 2015; Szarmach et al., 2017). In line with these observations, a strong association between air pollution with systemic brain inflammation was revealed in children

living in polluted areas, associated with short-term memory deficits, prefrontal WMH, and BBB disruption (Calderón-Garcidueñas et al., 2016). The same study reported a leaking vascular network, degeneration of pericytes, VSMCs, and endothelial cells, thickening of the BM, and reduced perivascular astroglial coverage in the prefrontal white matter of dog brains (Calderón-Garcidueñas et al., 2016). Exposure to PM_{2.5} has been demonstrated as well to accelerate atherosclerosis development through induction of vascular dysfunction as well as promotion of coagulopathies, which were accompanied by a strong inflammatory response and lipid abnormalities (Liang et al., 2020). Finally, exposure of ApoE-deficient mice to TRAP, mixed vehicle emissions, induced the cerebral expression of ICAM-1 and the release of pro-inflammatory mediators, such as TNF- α and IL-1 β (Adivi et al., 2021).

Impact on the Dynamics of Astrocytes and Oligodendrocytes

Evidence indicates that astrocytes respond to PM in a context-dependent manner (Allen et al., 2014). For instance, early postnatal exposure to ambient UFPs decreased GFAP immunoreactivity in male subjects and increased GFAP expression as well as other neuroinflammation markers in females (Allen et al., 2014). Exposure of male and female rodents during gestation and early postnatal development to TRAP attenuated astrogliosis specifically in the dentate gyrus (DG) associated with reduced GFAP immunoreactivity, which remained unchanged in CA1 and CA3 regions (Patten et al., 2020). Previous findings have shown that maternal exposure to carbon black nanoparticles (CB-NP) induced astrogliosis in the cortex of rodents, affecting the interaction of the astrocyte endfeet with the endothelium and perivascular macrophages (Onoda et al., 2017). Interestingly, intranasal delivery of PM_{2.5} to male rodents subjected to ischemic stroke exacerbated astrocytic reactivity *via* GFAP activation and iNOS induction, aggravating post-stroke neurobehavioral impairments (Chen et al., 2016). In line with these results, rodents exposed to PM for a prolonged period exhibited altered neuronal and astrocytic functions *via* impairment of mitochondrial activity (Araújo et al., 2019). Likewise, exposure of rodents to natural air pollution sources, such as volcanic-derived particles, increased GFAP immunoreactivity in astrocytes (Camarinho et al., 2019; Navarro et al., 2021). Moreover, exposure of rodents to low doses of CB-NP induced endoplasmic reticulum (ER) stress in perivascular macrophages and reactive astrocytes, specifically around the vasculature of offspring animals, associated with the accumulation of β -sheet rich misfolded proteins (Onoda et al., 2020). Cell-based assays have shown that exposure of astrocytes to PM activated janus kinase (JAK)-2/STAT-3 and p38/JNK/ERK pathways in reactive astrocytes triggering iNOS induction and IL-1 β production (Li et al., 2016). In this regard, PM has been reported to increase the expression and release of proinflammatory mediators through activation of the NF- κ B signaling pathway (Li et al., 1999; Gómez-Budia et al., 2020).

Furthermore, exposure to UFPs altered adult OPCs turnover and survival of mature oligodendrocytes (OLs), accompanied by increased oxidative stress and decreased total antioxidant

capacities (TAC) that impair the remyelination capacity of the brain (Kim J. Y. et al., 2020). Furthermore, prenatal exposure to concentrated ambient particles (CAPs), which were defined as UFPs, promoted a premature maturational shift in OLs in the corpus callosum (CC), followed by hypermyelination (Klocke et al., 2018). Interestingly, females showed significant alterations in oligodendrocytogenesis in the CC (Klocke et al., 2018). Finally, in a mouse model of lysolecithin-induced demyelination of the subcortical white matter, exposure to PM_{2.5} hampered remyelination and disrupted oligodendroglia differentiation (Parolisi et al., 2021).

Impact on Microglial Reactivity and Consequences on Neurons

In response to air pollution, microglia are activated through the upregulation of several proinflammatory mediators that affect neuronal function (Gómez-Budia et al., 2020). Microglia respond to air pollution by adopting an amoeboid shape *in vitro* (Cole et al., 2016; Roqué et al., 2016; Mumaw et al., 2017), as well as *in vivo* (Araújo et al., 2019). Indeed, mice exposed to DEP exhibited reduced adult neurogenesis through activation of microglia (Coburn et al., 2018). In contrast, attenuating microglial reactivity hampered neuroinflammation and oxidative stress (Coburn et al., 2018). Cell-based assays showed that exposure of BV-2 immortalized microglial cell to PM_{2.5} increased mRNA expression of various proinflammatory markers, namely IL-6, IL-1 β , TNF- α , iNOS, COX-2, TREM2, and TLR2/4, while reducing mRNA expression of key anti-inflammatory markers, such as IL-10 and Arginase (ARG)-1 (Kim R.-E. et al., 2020). In parallel, exposure of rodents to DEP for a long period triggered the activation of microglia in the nuclei of the solitary tract (NTS), accompanied by ER dilation and mitochondrial vacuolization in the medulla, hence outlining structural alterations of the neuronal network (Chen et al., 2021). Interestingly, rodents exposed to DEP for a long period showed an induced ionized calcium-binding adaptor molecule (IBA)-1 expression in the brain (Levesque et al., 2011). Interestingly, microglial reactivity was accompanied by an increased mRNA expression of TNF- α , IL-6, and macrophage inflammatory protein (MCP)-1 α in the midbrain, cortex, and OB, while IL-1 β was expressed particularly in the midbrain (Levesque et al., 2011). Mice exposed to PM also triggered the induction of mRNA expression of TLR4, MyD88, TNF- α , and tumor necrosis factor receptor (TNFR)-2 in microglia (Woodward et al., 2017a). Activation of the latter was further confirmed through an increased intracellular expression of inflammatory mediators, such as COX-2, NF- κ B, prostaglandin E2 (PGE2), and iNOS both *in vitro* and *in vivo* (Babadjouni et al., 2018; Chen et al., 2018). Using cell-based assays, TNF- α induction in microglia upon PM exposure inhibited neurite outgrowth (Cheng et al., 2016). Inhibition of NRF-2 activity prior to exposure of BV-2 microglial cells to PM_{2.5} attenuated cell viability, induced ROS generation, and stimulated NF- κ B pathway, outlining NRF-2 role in mitigating PM deleterious effects (Chen et al., 2018).

Importantly, *in vivo* experiments showed that exposure to DEP in TREM2^{-/-} mice accentuated IL-1 β expression (Greve et al., 2020). PM has been demonstrated to impact

neuron-glia crosstalk. Acute exposure of adult mice to PM_{2.5} increased the levels of lipoperoxidation and proinflammatory cytokines in the brain and activated microglia, accompanied by reduced neurogenesis in the subgranular zone (SGZ) and subventricular zone (SVZ; Bernardi et al., 2021). Neuronal cell cultures exhibited reduced viability upon exposure to PM_{2.5} associated with increased release of glutamate (Liu F. et al., 2015). Prior treatment of cells with the *N*-methyl-D-aspartate (NMDA) receptor mitigated PM_{2.5}-mediated neuronal loss (Liu F. et al., 2015). Moreover, neuronal cultures displayed dopaminergic neurotoxicity upon exposure to DEP only in presence of microglia, which was associated with elevated levels of ROS (Block et al., 2004). Co-culture of neurons and microglia exposed to PM_{2.5} in presence of oligomeric oA β exacerbated IL-1 β and ROS release aggravating oA β -induced neuronal injury and inflammation (Wang et al., 2018). PM have been shown to directly impact neuronal function. For instance, exposure of rodents to nano PM caused hippocampal neurite atrophy and decreased expression of myelin basic protein (MBP), accompanied by increased TNF- α mRNA expression (Woodward et al., 2017b). In this regard, epidemiological studies have outlined a strong correlation between the levels of PM in air and neuronal chromatolysis and satellitosis in exposed dogs, associated with cortical neurons degeneration, and neurofibrillary tangle formation (Calderón-Garcidueñas et al., 2002).

Interestingly, epidemiological studies comprising elderly women revealed that residence in places contaminated with high levels of fine PM increases the risks for global cognitive decline and all-cause of dementia by 81 and 92%, respectively, with stronger adverse effects in ApoE4 carriers (Cacciottolo et al., 2017). Experimental findings obtained from female AD mouse models (5xFAD; Familial AD) expressing either ApoE3 or ApoE4 mice that were exposed to urban nano PM for 15 weeks showed increased A β plaques and soluble A β oligomers, which were associated with neuronal changes in the hippocampus (Cacciottolo et al., 2017). These findings were confirmed *in vitro* upon exposure of neuroblastoma cells (N2a-APP/swe) to nano PM translated by an enhanced pro-amyloidogenic processing of the APP, explaining the elevated cerebral A β production (Cacciottolo et al., 2017). Furthermore, long-term exposure to ambient air pollution was found to be associated with rapid cognitive decline in aged adults, where ApoE4 carriers exhibited the fastest cognitive decline (Kulick et al., 2020; **Figure 2**).

Further *in vivo* experimental investigations implicating exposure to TRAP nano PM showed an increased production of A β peptides, associated with oxidative damage (Cacciottolo et al., 2020). Indeed, exposure of J20-APPswe mice, an AD mouse model, to nano PM for 150 h revealed exacerbated lipid oxidation and pro-amyloidogenic processing of APP in lipid raft fractions compared to controls (Cacciottolo et al., 2020). These observations were further confirmed *in vitro* using N2a-APPswe cells exposed to nano PM (Cacciottolo et al., 2020). Importantly, the link between air pollution and HDL was highlighted in the Multi-Ethnic Study of Atherosclerosis Air Pollution (MESA Air) study showing that exposure to air pollution was significantly associated with low levels of

HDL (Bell et al., 2017). In this regard, ApoE-deficient mice exposed for 2 weeks to DEP, exhibited high systemic pro-oxidant effects associated with dysfunctional HDL (Yin et al., 2013). These observations would indicate that PM exposure could be responsible for the attenuated HDL protective effects against atherosclerosis (Yin et al., 2013).

CONCLUSION AND PERSPECTIVES

Through this review, we aimed to highlight the structural and functional neurovascular alterations underlying VaD pathobiology. The epidemiological, clinical, and experimental investigations are indicating that the long-term outcomes of early pathophysiological events impacting neurovascular functions upon cerebrovascular disorders have major consequences on imitating a cascade of events that lead to VaD. Moreover, the recent findings are outlining air pollution as a major vascular risk factor that is directly implicated in promoting neurovascular impairments associated with VaD. Given the heterogeneity of cerebrovascular disorders, which include stroke, genetic and sporadic microangiopathies, combined with the diverse effects of environmental factors, a better understating of the short- and long-term remodeling processes at the neurovascular unit is urgently required to allow getting new insights into VaD pathobiology. Such knowledge will allow the identification of key “targetable” mechanisms for therapeutic purposes.

Recent findings are suggesting that BBB disruption occurs in some cases for a prolonged period beyond the acute and subacute phases after stroke (Bernardo-Castro et al., 2020). A prolonged, yet subtle, disruption of the BBB could trigger a cascade of events that lead to VaD. For instance, high expression levels of MMPs, the main mediator of BBB disruption, are significantly associated with cognitive deficits (Yang and Rosenberg, 2011). However, targeting MMPs is a double-edged sword as while preserving the BBB in the acute and sub-acute phase, its inhibition impairs neurovascular adaptation in the chronic phase, thus impeding brain plasticity (Yang and Rosenberg, 2011). More promising appears to be the inhibition of beta-site APP cleaving enzyme (BACE)1 that seems to protect endothelial cell integrity in the context of CADASIL and AD (Chacón-Quintero et al., 2021). In the same way, rodent models of aging suggest that the mitochondrial overexpression of catalase improves diminish vascular impairment and benefits neurovascular coupling (Csiszar et al., 2019).

On the other hand, uncontrolled post-stroke neuroinflammation, especially excessive microglial activation, is associated with cognitive deficits (Guruswamy and ElAli, 2017; Gefen et al., 2019; Jayaraj et al., 2019; Zhang et al., 2021). In this regard, strategies aiming to mitigate neuroinflammation *via* modulation of microglia could attenuate cognitive decline related to stroke (Guruswamy and ElAli, 2017; Dzubyenko et al., 2018; Jayaraj et al., 2019). In this regard, evidence suggests that the prominent microglial regulator, insulin-like growth factor (IGF)-1, reduces gliosis while preserving brain volume and myelination, as well as motor performance and memory when administered intranasally in aged mice (Farias Quipildor

et al., 2019). Efficient immunomodulatory approaches capable of fine-tuning microglial activation are still to be developed.

Importantly, the emergence and turnover of stroke-mediated proteinopathies are associated with BBB disruption and neuroinflammation. Although the role of cSVD is critically important in the etiology of VaD, there is still a huge gap in the literature as to our understanding of the underlying mechanisms. This is due to the fact that the current knowledge is obtained either from genetic or non-clinically relevant sporadic animal models. Indeed, the majority of cSVD cases are sporadic and associated with diverse risk factors, as such, it is critically important to develop animal models that replicate some of the pathological features associated with VaD. However, the overwhelming findings indicate the loss-of-function on the local cerebrovascular network caused a central role in initiating a pathological cascade of events that lead to altered neuro-glial functions, and thus subsequently dementia (Rouhl et al., 2012; Shoamanesh et al., 2015; Fu and Yan, 2018; Li et al., 2018; Jian et al., 2020).

Air pollution has emerged as a significant risk factor for cerebrovascular and neurodegenerative disorders (Wu et al., 2015; Åström et al., 2021). It is becoming clear that exposure to PM, one of the most deleterious air pollutants, increases the risk of chronic neuroinflammation that leads to dementia (Wu et al., 2015; Åström et al., 2021). Cell-based assays have demonstrated that PM acts as a powerful inflammatory and oxidative stress mediator in various brain cell types. Interestingly, PM exposure seems to amplify the pathological responses underlying VaD pathobiology. Although the impact of PM on neurovascular functions is evidenced *in vitro*, little is known about its role in mediating neurovascular impairments *in vivo*. Future studies should consider investigating the consequences of PM exposure as a comorbid condition in the design of preclinical experiments.

As previously mentioned, besides PM, various modifiable vascular risk factors are recognized to impact VaD. For instance, stroke and cSVD share common risk factors such as hypertension, atherosclerosis, obesity, atrial fibrillation, diabetes, dyslipidemia, high homocysteine, metabolic syndrome, smoking, as well as cardiac and carotid arterial disease (Barnes and Yaffe, 2011; O'Brien and Thomas, 2015; Kalaria et al., 2016; Tariq and Barber, 2018). Given their central role, various preventive approaches have been developed and adopted to attenuate the impact of this triad of vascular on VaD. Such approaches are exemplified by the landmark multidomain Finnish Geriatric Intervention Study to Prevent Cognitive Impairment and Disability (FINGER), which has been shown to enhance all cognitive sub-domains through a multidomain lifestyle intervention that include dietary counseling, physical exercise, cognitive training, and vascular and metabolic risk monitoring for a period of 2 years (Ngandu et al., 2015; Kivipelto et al., 2020). Furthermore, knowing the detrimental role of high blood pressure in VaD prevalence, the Systolic Blood Pressure Intervention Trial (SPRINT) and its sub-study the Memory and Cognition in Decreased Hypertension (MIND) were established with an emphasis on investigating the consequences of lowering systolic blood pressure. Although the incidence of dementia was not

improved, the trial reported a reduction of mild cognitive impairment (MCI) and MCI composite in subjects with lower blood pressure. Interestingly, these observations were associated with a reduction of WM lesion volumes (Kjeldsen et al., 2018; Peters et al., 2019). Despite that, each trial discloses important limitations, yet the impact of modifying main VaD risk factors was shown to allow the development of efficient interventions. Further research is needed for longer periods with a more representative population to lessen the limitations and improve the efficiency of interventions aiming to attenuate dementia prevalence.

Finally, a deep analysis of the current scientific knowledge outlines brain pericytes as a major effector in the pathobiology of VaD. Through their spatial localization, pericytes play a central role in integrating and processing signals from their milieu to generate critical neurovascular functions, which include BBB maintenance, CBF modulation, vascular stabilization, and immunomodulation (Zlokovic, 2011; Hermann and ElAli, 2012). Following a stroke, pericytes undergo cell death at the ischemic core and get activated in the peri-lesion site where they detach from the vasculature (Zlokovic, 2011; Hermann and ElAli, 2012). Importantly, it has been reported that even after successful recanalization, pericytes located at the peri-lesion site remain contracted impeding the capillary microcirculation, which leads to vascular constriction and chronic hypoperfusion (Dalkara, 2019). Moreover, microcirculation of the white matter was disrupted upon pericyte degeneration leading to the accumulation of toxic blood-derived fibrotic deposits within the vasculature, which promotes vascular fibrosis, accompanied by a reduction in the regional CBF (Montagne et al., 2018). These changes are supposed to be directly implicated in the pathogenesis of diffuse WMD associated with loss of

oligodendrocytes and subsequently myelinated axons (Montagne et al., 2018). The findings indicate that pericyte degeneration plays a major role in the pathogenesis, and thereby therapies, of WMD associated with cSVD (Montagne et al., 2018). Furthermore, the generation of pericytes is translated by elevated levels of the soluble-platelet derived growth factor receptor (PDGFR) β (sPDGFR β) in the CSF, strongly correlating with BBB breakdown, CBF reduction, and cognitive decline (Sweeney et al., 2020). Therefore, decoding the pericyte reactivity to stressors related to the vascular risk factors constitutes a promising avenue that might lead to achieving major breakthroughs in getting new mechanistic insights in the pathobiology of VaD and in developing novel therapeutic interventions.

AUTHOR CONTRIBUTIONS

The authors contributed to the review as follows: SL wrote, edited, and finalized the draft as well as prepared the figures. DM-C wrote, edited, and finalized the draft as well as prepared the figures. YE wrote the section related to air pollution. AEA conceptualized, wrote, and finalized the draft. All authors contributed to the article and approved the submitted version.

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Neurovascular Coupling Is Impaired in Hypertensive and Diabetic Subjects Without Symptomatic Cerebrovascular Disease

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The mechanistic link between hypertension, diabetes and cerebral small vessel disease (CSVD) is still poorly understood. We hypothesized that hypertension and diabetes could impair cerebrovascular regulation prior to irreversibly established cerebrovascular disease. In this study, 52 hypertensive patients [54% males; age 64 ± 11 years; 58% with comorbid diabetes mellitus (DM)] without symptomatic cerebrovascular disease underwent transcranial Doppler (TCD) monitoring in the middle (MCA) and posterior (PCA) cerebral arteries, to assess vasoreactivity to carbon dioxide (VRCO₂) and neurovascular coupling (NVC). 1.5T magnetic resonance imaging was also performed and white matter hyperintensity volume was automatically segmented from FLAIR sequences. TCD data from 17 healthy controls were obtained for comparison (47% males; age 60 ± 16 years). Hypertensive patients showed significant impairment of NVC in the PCA, with reduced increment in cerebral blood flow velocity during visual stimulation (22.4 ± 9.2 vs. 31.6 ± 5.7 , $p < 0.001$), as well as disturbed NVC time-varying properties, with slower response (lower rate time: 0.00 ± 0.02 vs. 0.03 ± 6.81 , $p = 0.001$), and reduced system oscillation (reduced natural frequency: 0.18 ± 0.08 vs. 0.22 ± 0.06 , $p < 0.001$), when compared to controls. VRCO₂ remained relatively preserved in MCA and PCA. These results were worse in hypertensive diabetic patients, with lower natural frequency ($p = 0.043$) than non-diabetic patients. White matter disease burden did not predict worse NVC. These findings suggest that hypertensive diabetic patients may have a precocious impairment of NVC, already occurring without symptomatic CSVD. Future research is warranted to evaluate whether NVC assessment could be useful as an early, non-invasive, surrogate marker for CSVD.

Keywords: hypertension, diabetes mellitus, neurovascular coupling (NVC), transcranial doppler (TCD), cerebral small vessel disease

INTRODUCTION

Cerebral small vessel disease (CSVD) has an enormous impact on public health worldwide (GBD 2017 Causes of Death Collaborators, 2018). It accounts for 25% of ischemic strokes and most hemorrhagic strokes and is the second leading cause for cognitive decline (Sudlow and Warlow, 1997; Iadecola et al., 2019).

Hypertension (HT) is the major vascular risk factor (VRF) for CSVD. Alongside HT, diabetes mellitus (DM) is a recognized VRF implicated in CSVD (Brundel et al., 2012; Liu et al., 2018).

While other etiologies for stroke are fairly well-studied, the pathophysiology and causality of CSVD are still poorly understood. Many of its manifestations are clinically silent until the development of clinical consequences, with stroke, cognitive decline, and gait impairment, limiting disease-specific preventive strategies (Pantoni, 2010). Also, CSVD radiological markers and clinical manifestations seem to be dissociated, for reasons not fully explained (Sorond et al., 2011; Jokumsen-Cabral et al., 2019). Biomarkers for the events predating irreversible damage could be key for better clinical management and pre-symptomatic preventive measures.

There is evidence of neurovascular dysfunction in CSVD and it may precede clinical and imaging manifestations (Wardlaw, 2010; Freeze et al., 2018; Castro et al., 2020). Very few studies have investigated neurovascular coupling (NVC) in HT, mostly using imaging modalities. Transcranial Doppler (TCD) is a non-invasive method that allows the monitoring of microvascular hemodynamic functional integrity (Claassen et al., 2016; Malojcic et al., 2017).

We aimed to study cerebrovascular regulation by TCD in hypertensive and diabetic patients without major CSVD-related impairment as a possible surrogate marker for the future development of symptomatic CSVD, to help guide therapies aimed at the cerebral microcirculation and neurovascular unit.

MATERIALS AND METHODS

Study Subjects

A cross-sectional observational study was conducted in a University Hospital. Hypertensive patients were recruited from the hospital's Hypertension Unit. Exclusion criteria were previous stroke or other significant brain pathology (dementia by clinical criteria, brain tumor, traumatic brain injury, previous cerebral infection or neurodegenerative disease), severe/unstable disease, contraindication for magnetic resonance imaging (MRI), inadequate acoustic temporal bone window, extra- or intracranial artery stenosis >50% and incapability to collaborate or to give informed consent. TCD data from healthy controls of similar age and gender were obtained from previous studies performed with the same protocol (Jokumsen-Cabral et al., 2019).

The local ethics committee approved the study protocol, which followed the tenets of the Declaration of Helsinki. Written informed consent was obtained.

Clinical Evaluation

Participant's clinical and demographic data were recorded. Vascular comorbidities were summarized into a vascular comorbidity score (VCS), including HT, DM, dyslipidemia, tobacco usage, chronic heart failure, coronary heart disease, arrhythmias, peripheral artery disease, and nephropathy. These conditions were scored as present (1 point) or absent (0 points), for a score ranging from 0 to 9 (Mossello et al., 2015). All participants underwent cervical and transcranial Doppler ultrasound (Philips iu22, The Netherlands) to exclude hemodynamically significant vessel pathology. The patients underwent routine 24-h ambulatory blood pressure monitoring (Spacelabs 90207, Redmond, Washington, USA). The mini-mental state examination (MMSE) and the Montreal cognitive assessment (MoCA) were used to screen for dementia. The patients were evaluated by an ophthalmologist, and all had normal binocular visual acuity, allowing for the TCD dynamic testing.

Monitoring Protocol

Evaluations were conducted in a dim-lighted, quiet room ($\approx 22^{\circ}\text{C}$), in a supine position. Cerebral blood flow velocity (CBFV) was continuously recorded in the M1 segment of the right middle cerebral artery (MCA) and the P2 segment of the left posterior cerebral artery (PCA), with 2-MHz TCD probes secured with a headframe (Doppler BoxX, DWL, Singen, Germany), in order to simultaneously obtain data from both arterial territories (Azevedo et al., 2012). Continuous non-invasive arterial blood pressure (BP) was measured with the Finometer (FMS, Amsterdam, The Netherlands). Heart rate was assessed with a three-lead electrocardiogram. End-tidal carbon dioxide (EtCO_2) was recorded by capnography (Respsense Nonin, Amsterdam, The Netherlands). Data was synchronized and digitally stored at 400 Hz with Powerlab (AD Instruments, Oxford, UK) for offline analysis. After resting for 20 min, the vasoreactivity to carbon dioxide (VRCO_2) and NVC protocols were performed, as described below. CBFV envelopes were continuously registered and analyzed offline.

VRCO_2

Participants were monitored through successive 2-min steps of resting, inhalation of a mixture of 5% CO_2 and 95% O_2 mixture (EtCO_2 7–10 mmHg above baseline), resting (room air, until normocapnia) and hyperventilation (EtCO_2 7–10 mmHg below baseline). VRCO_2 was calculated as the slope of the relationship between EtCO_2 average values plotted against those of relative CBFV achieved at the three stages, expressed as % mean CBFV per mmHg EtCO_2 (Madureira et al., 2017).

NVC

NVC was assessed in the PCA territory by a visual paradigm consisting of 10 cycles, each with a 20s resting phase (eyes closed) and 40s stimulating phase (flickering checkerboard) at 10 Hz (Rosengarten et al., 2001a). The 5s of stable measurement prior to stimulation were used as the baseline (Rosengarten et al., 2001a). All cycles were synchronized and averaged. Peak systolic data was used because it is less prone to artifacts (Rosengarten

et al., 2001b). Maximal systolic CBFV change was obtained to calculate the overshoot parameter as $\frac{\text{maximumCBFV} - \text{baselineCBFV}}{\text{baselineCBFV}} \times 100\%$. [12] The systolic CBFV curve was modeled into a second order linear system to describe the dynamics of NVC response in time according to the equation $G(s) = \frac{K \times (1 + T_{vs})}{\frac{s^2}{\omega^2} + 2\xi \frac{s}{\omega} + 1}$, where “K”

stands for gain, “ T_v ” for rate time, “ ω ” for natural frequency, and “ ξ ” for attenuation (Rosengarten et al., 2001a). All the parameters of the equation were determined by the least squares method. The sum of the squared residuals and the χ^2 were also calculated to ensure the goodness of fit into the real measured values, as provided by the lsqnonlin function. Gain describes the relative

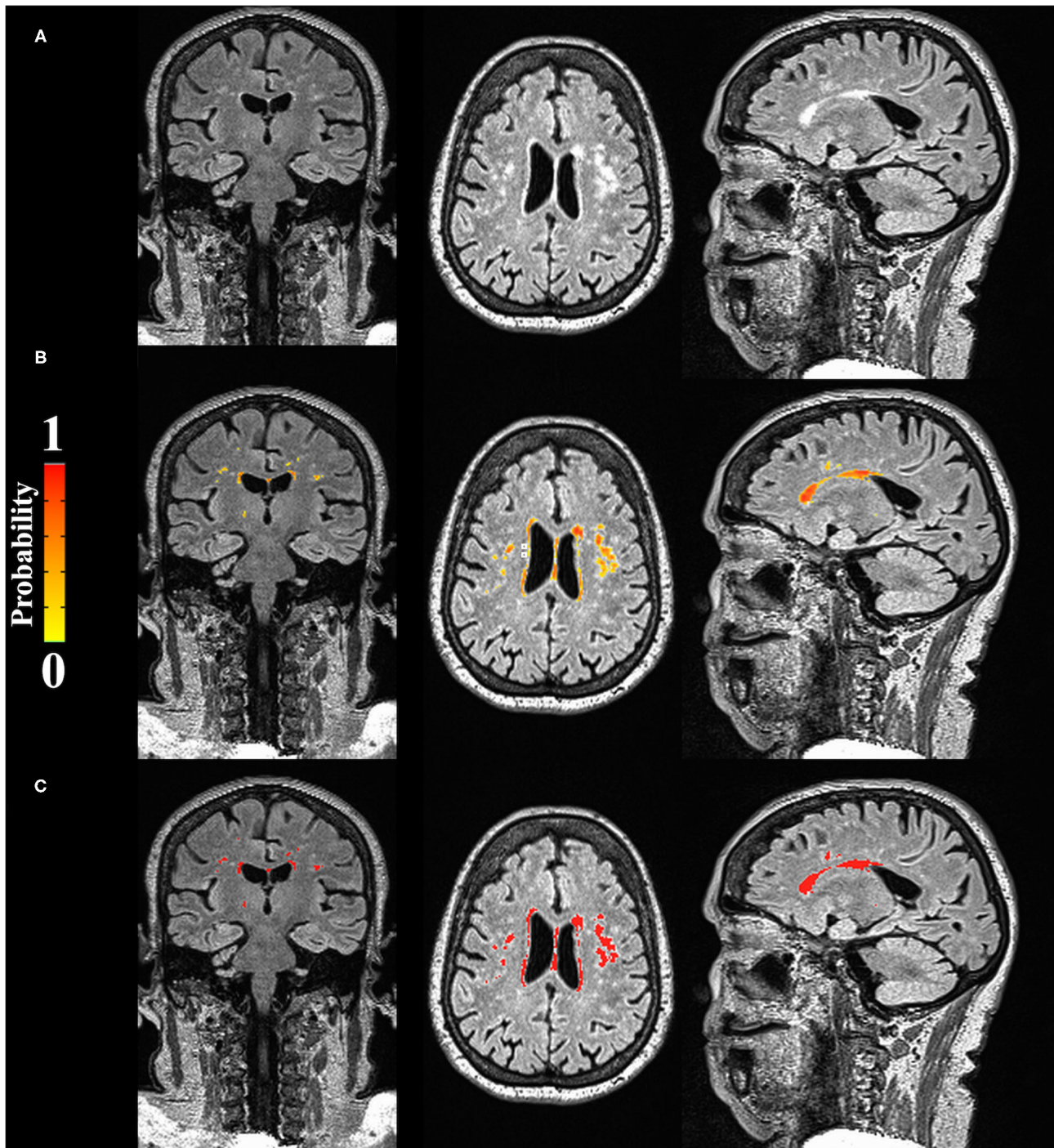


FIGURE 1 | Automatic segmentation of WMH (A) Raw data; (B) Probability map of WMH; (C) Binarized results of WMH (values above 0 were considered WMH).

CBFV difference between rest and steady-state level during visual stimulation; rate time indicates the initial steepness of the CBFV increase; natural frequency represents the oscillatory properties of the system; and attenuation describes dampening and tonus features, such as elastic properties of the vessel wall (Rosengarten et al., 2003). For this specific test, the MCA recordings were used as a control to detect non-specific changes in CBFV during the visual stimulation task.

MRI Imaging

Forty-six patients were eligible and agreed to undergo cerebral MRI (Siemens Aera 1.5T). Of these, data from 6 patients were excluded for lack of quality for the evaluations. White matter hyperintensity (WMH) volumes normalized by intracranial volume were derived from the T2-weighted fluid-attenuated inversion recovery sequences collected in the sagittal plane. Voxels resolution was $1 \times 1 \times 1$, slices = 256, FOV = 256 mm, TR = 5000 ms, TE = 336 ms, TI = 1800 ms (Figure 1). Briefly, WMH masks were created using the Lesion Segmentation Algorithm (LPA, 1) from the Lesion Segmentation Toolbox for SPM12 in MATLAB R2018a. Following an initial segmentation of the FLAIR image, probability maps were binarized using AFNI (2.3, v21.0.15) command 3dcalc. Resulting segmentations were quality-checked for sufficient accuracy and volumes were calculated using Freesurfer (v7.1) command mri_segstats.

Additional signs of CSVD were evaluated by an experienced vascular Neurologist to further characterize the CSVD in the patient cohort. Enlarged perivascular spaces (PVS) were defined as small (<0.3 mm) punctate or linear (if perpendicular or longitudinal to the plane of the scan, respectively) hyperintensities on T2 images in the basal ganglia (BG) and centrum semiovale (CS) (Potter et al., 2015). The PVS burden was then stratified into three groups: <11 , 11 – 20 and >20 (Lau et al., 2017). Lacunes were defined as rounded or ovoid lesions >3 mm and <20 mm in diameter in the BG, internal capsule, CS or brainstem, with CSF density on T2 images (Wardlaw et al., 2013). Cerebral microbleeds were defined as round, hypodense lesions <10 mm on susceptibility weighted imaging collected in the axial plane (slice thickness 2 mm, slices = 256, FOV = 230 mm, TR = 49 ms, TE = 40 ms), according to the guidelines (Greenberg et al., 2009).

Statistical Analysis

Normality was determined using the Shapiro–Wilk test and analysis of skewness. Data with high asymmetry was normalized using logarithmic transformation. Homogeneity of variances was tested for each analysis. Baseline characteristics were compared using the independent sample *t*-test and χ^2 -test. Mixed ANOVA or ANCOVA (for PCA) were used to compare hemodynamic data. Subgroup analyses were performed using univariate ANOVA, with Bonferroni *post-hoc* tests. The partial eta squared (η^2) was used as a measure of effect size: $\eta^2 > 0.14$ indicates a large effect, $\eta^2 > 0.06$ indicates a medium effect and $\eta^2 > 0.01$ indicates a small effect (Cohen, 1998). Age and gender were used as covariates in the ANOVA and ANCOVA comparisons. Supplemental analysis using body mass index (BMI) and VCS as covariates (plus age and gender) were

also performed. Patients were dichotomized by the median value of WMH volume to compare the results of VRCO₂ and NVC in the PCA between the two WMH burden groups. Subgroups were compared using the independent sample *t*-test.

Values of $p < 0.05$ were considered significant.

RESULTS

Fifty-two hypertensive patients were evaluated and TCD data from 17 healthy controls were used. Baseline characteristics are reported in Table 1. Supplementary File 1 depicts the burden of

TABLE 1 | Demographics and baseline characteristics.

Participant characteristics	Patients	Controls	<i>p</i> value*
Age, years (mean \pm SD)	64 \pm 11	60 \pm 16	0.376
Male, n (%)	28 (54)	8 (47)	0.627
BMI, kg/m ² (median \pm IQR)	29 \pm 5	25 \pm 4	<0.001
Diabetes Mellitus, n (%)	30 (58)	0 (0)	
VCS, n (%)			
0	0 (0)	17 (100)	
1	2 (4)	0 (0)	
2	5 (10)	0 (0)	
3	22 (42)	0 (0)	
4	13 (25)	0 (0)	
5	8 (15)	0 (0)	
6	2 (4)	0 (0)	
HT duration, years (median \pm IQR)	17 \pm 6	0 (0)	
Chronic medication			
No. antihypertensives (median \pm IQR)	3 \pm 2	0 (0)	
ACEI/ARB, n (%)	48 (92)	0 (0)	
Diuretics, n (%)	39 (75)	0 (0)	
CCB, n (%)	37 (71)	0 (0)	
BB, n (%)	14 (27)	0 (0)	
Alpha2-agonists, n (%)	4 (8)	0 (0)	
Antiplatelets, n (%)	14 (27)	0 (0)	
Statins, n (%)	40 (77)	0 (0)	
Cognitive parameters (median \pm IQR)			
Education, years	4 \pm 5		
MMSE	28 \pm 3		
MoCA	22 \pm 5		
ABPM, mmHg (median \pm IQR)			
24-h systolic BP	130 \pm 15	-	-
24-h diastolic BP	78 \pm 13	-	-
24-h pulse pressure	57 \pm 16	-	-
Finapres BP, 5 min (mean \pm SD)			
Systolic BP	136 \pm 24	109 \pm 25	<0.001
Diastolic BP	62 \pm 17	50 \pm 14	0.035
Mean BP SP, mmHg ₂ (median \pm IQR)	9 \pm 24	7 \pm 8	0.007
EtCO ₂ , mmHg (median \pm IQR)	36 \pm 4	38 \pm 3	0.553

*Values were obtained using the *t*-test or the χ^2 -test. ABPM, ambulatory blood pressure monitoring; BMI, body mass index; BP, blood pressure; EtCO₂, end-tidal carbon dioxide; IQR, interquartile range; SD, standard deviation; SP, spectral power; VCS, vascular comorbidity score. Bold values of statistically significant *p* values ($p < 0.05$).

PVS, microbleeds and lacunes in the patient group. Most patients had low burden of PVS in the BG (55.0%) and CS (65.0%). Nine patients (22.5%) presented lacunes and four patients (10%) presented microbleeds.

Cerebral Hemodynamics and VRCO₂

Baseline CBFV and CBFV variability (spectral power) were similar in patients and controls (Table 2). VRCO₂ in either territory (PCA or MCA) was similar between groups and

showed no change after adjusting for both VCS and BMI (Supplementary File 2).

There were no differences in VRCO₂ between hypertensive non-diabetics (HT-nDM), hypertensive diabetics (HT-DM) and controls (Table 3). Those results did not change significantly after controlling for BMI or VCS (Supplementary File 3).

NVC

NVC in the PCA territory was significantly altered in HT patients, with smaller increases in CBFV during visual

TABLE 2 | Cerebral hemodynamics, VRCO₂ and NVC: patients vs. controls, controlling for age and gender.

	Patients		Controls		Artery	Group	Interaction
	MCA	PCA	MCA	PCA	p value*	p value*	p value*
Cerebral hemodynamics							
Mean CBFV (cm/s)	47.6 ± 13.4 [†]	31.4 ± 8.2 [†]	52.4 ± 16.5 [†]	29.1 ± 10.1 [†]	0.044	0.925	0.161
MFV SP (cm/s ²)	3.3 ± 5.0 [†]	2.5 ± 4.7 [†]	3.8 ± 4.3 [†]	2.3 ± 3.1 [†]	0.976	0.770	0.119
VRCO ₂ (%/mmHg CO ₂)	1.4 ± 0.5 [†]	0.9 ± 0.4 [†]	1.7 ± 0.6 [†]	1.0 ± 0.7 [†]	0.107	0.588 [#]	0.437
Neurovascular coupling							
Overshoot [§] systolic CBFV (%)		22.4 ± 9.2 [†]		31.6 ± 5.7 [†]		<0.001	
Modeled parameters							
Gain (%)		14.0 ± 7.1 [†]		17.3 ± 4.8 [†]		0.118	
Natural frequency (Hz)		0.18 ± 0.08 [†]		0.22 ± 0.06 [†]		<0.001	
Attenuation (a.u.)		0.4 ± 0.4 [†]		0.4 ± 0.4 [†]		0.374	
Rate time (s)		0.00 ± 0.02 [†]		0.03 ± 6.81 [†]		0.001[#]	

*Two-factor mixed-design ANOVA for the interaction between group variable (patients vs. controls) and arterial territory (MCA vs. PCA), controlling for age and gender. For NVC, values were obtained using an ANCOVA. Effect size: rate time $\eta^2 = 0.174$, natural frequency $\eta^2 = 0.237$ and overshoot systolic CBFV $\eta^2 = 0.186$. [†]gender significantly interfered with the model; [#]age significantly interfered with the model; [†]Values are presented as mean ± standard deviation. [‡]Values are presented as median ± interquartile range. [§]Maximal CBFV increase during visual stimulation. a.u., arbitrary units; CBFV, cerebral blood flow velocity; VRCO₂, vasoreactivity to carbon dioxide; MFV SP, median flow velocity spectral power; MCA, middle cerebral artery; PCA, posterior cerebral artery. Bold values of statistically significant p values ($p < 0.05$).

TABLE 3 | Cerebral hemodynamics, VRCO₂ and NVC: controls vs. HT-nDM vs. HT-DM patients, controlled for age and gender.

	HT-nDM		HT-DM		Controls		Group	HT-nDM vs. Controls	HT-DM vs. Controls	HT-nDM vs. HT-DM
	MCA	PCA	MCA	PCA	MCA	PCA	p value*	p value*	p value*	p value*
Cerebral hemodynamics										
Mean CBFV (cm/s) [†]	47.0 ± 14.1	33.4 ± 8.4	48.0 ± 13.2	30.0 ± 7.8	52.4 ± 16.5	29.1 ± 10.1	0.993			
MFV SP (cm/s ²) [†]	3.1 ± 4.7	2.6 ± 5.0	3.7 ± 7.0	2.5 ± 5.0	3.8 ± 4.3	2.3 ± 3.1	0.921			
VRCO ₂ (%/mmHg CO ₂) [†]	1.4 ± 0.6	1.2 ± 0.7	1.4 ± 0.5	0.8 ± 0.3	1.7 ± 0.6	1.1 ± 0.4	0.676 [#]			
Neurovascular coupling										
Overshoot [§] systolic CBFV (%) [†]		25.1 ± 8.6		20.7 ± 9.3		31.6 ± 5.7	<0.001	0.080	<0.001	0.248
Modeled parameters										
Gain (%) [†]		13.9 ± 5.1		14.1 ± 8.2		17.3 ± 4.8	0.283			
Natural frequency (Hz) [†]		0.19 ± 0.04		0.16 ± 0.05		0.23 ± 0.06	<0.001[#]	0.052	<0.001	0.043
Attenuation (a.u.) [†]		0.4 ± 0.3		0.4 ± 0.3		0.5 ± 0.2	0.327 [#]			
Rate time (s) [†]		0.00 ± 0.00		0.00 ± 0.26		0.03 ± 6.81	0.005[#]	0.012	0.011	1.000

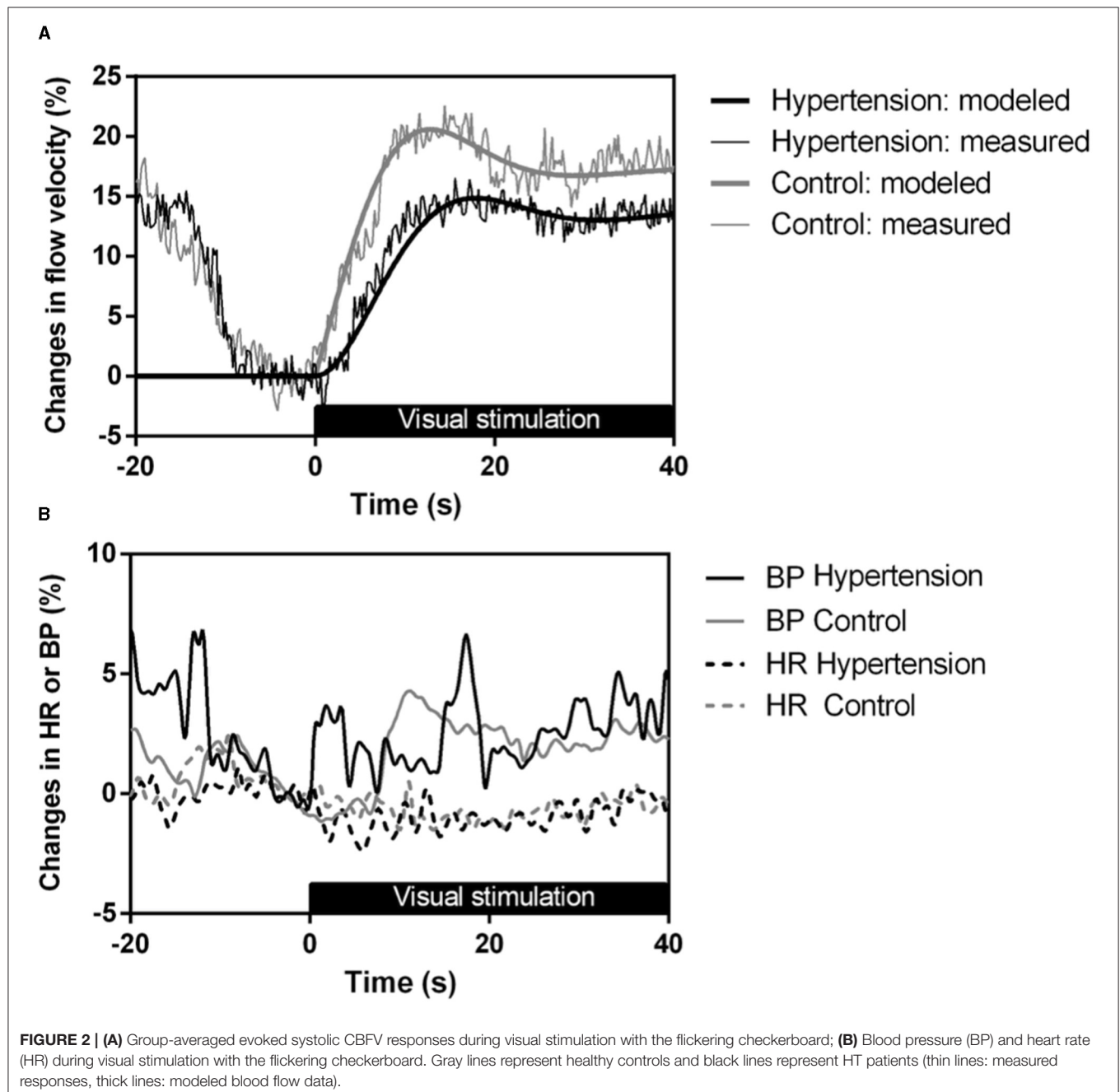
*Two-factor mixed-design ANOVA for the interaction between group variable (HT-DM vs. HT-nDM vs. controls) and arterial territory (MCA vs. PCA), controlling for age and gender, with Bonferroni post-hoc. For NVC, values were obtained using an ANCOVA. Effect size: rate time $\eta^2 = 0.174$, natural frequency $\eta^2 = 0.313$ and overshoot systolic CBFV $\eta^2 = 0.22$. [†]Values are presented as mean ± standard deviation. [‡]Values are presented as median ± interquartile range. [§]Maximal CBFV increase during visual stimulation. ^{||}gender significantly interfered with the model. [#]age significantly interfered with the model. a.u., arbitrary units; CBFV, cerebral blood flow velocity; HT-DM, hypertensive diabetic; HT-nDM, hypertensive non-diabetic; MCA, middle cerebral artery; MFV SP, median flow velocity spectral power; PCA, posterior cerebral artery; VRCO₂, vasoreactivity to carbon dioxide. Bold values of statistically significant p values ($p < 0.05$).

stimulation ($p < 0.001$) and disturbed NVC time-varying properties, with lower natural frequency ($p < 0.001$) and lower rate time ($p = 0.001$) (Table 2; Figure 2). These results remained similar when adjusting for BMI and VCS, although the overshoot systolic CBFV was over the limit of statistical significance when adjusting for the VCS ($p = 0.065$) (Supplementary File 2).

NVC results were worse in HT-DM than HT-nDM (Table 3). For the overshoot systolic CBFV and for natural frequency, only patients with both comorbidities showed

significant differences when comparing to controls. HT-DM showed lower natural frequency than non-diabetic patients ($p = 0.043$). When controlling for VCS, natural frequency was also worse in both HT-nDM ($p = 0.020$) and HT-DM ($p = 0.002$) when comparing to controls, with the worst results for HT-DM (HT-nDM vs. HT-DM, $p = 0.011$) (Supplementary File 3).

Figure 3 represents the evoked systolic CBFV responses in the PCA and MCA during the visual stimulation in one HT subject, to demonstrate the individual NVC responses.



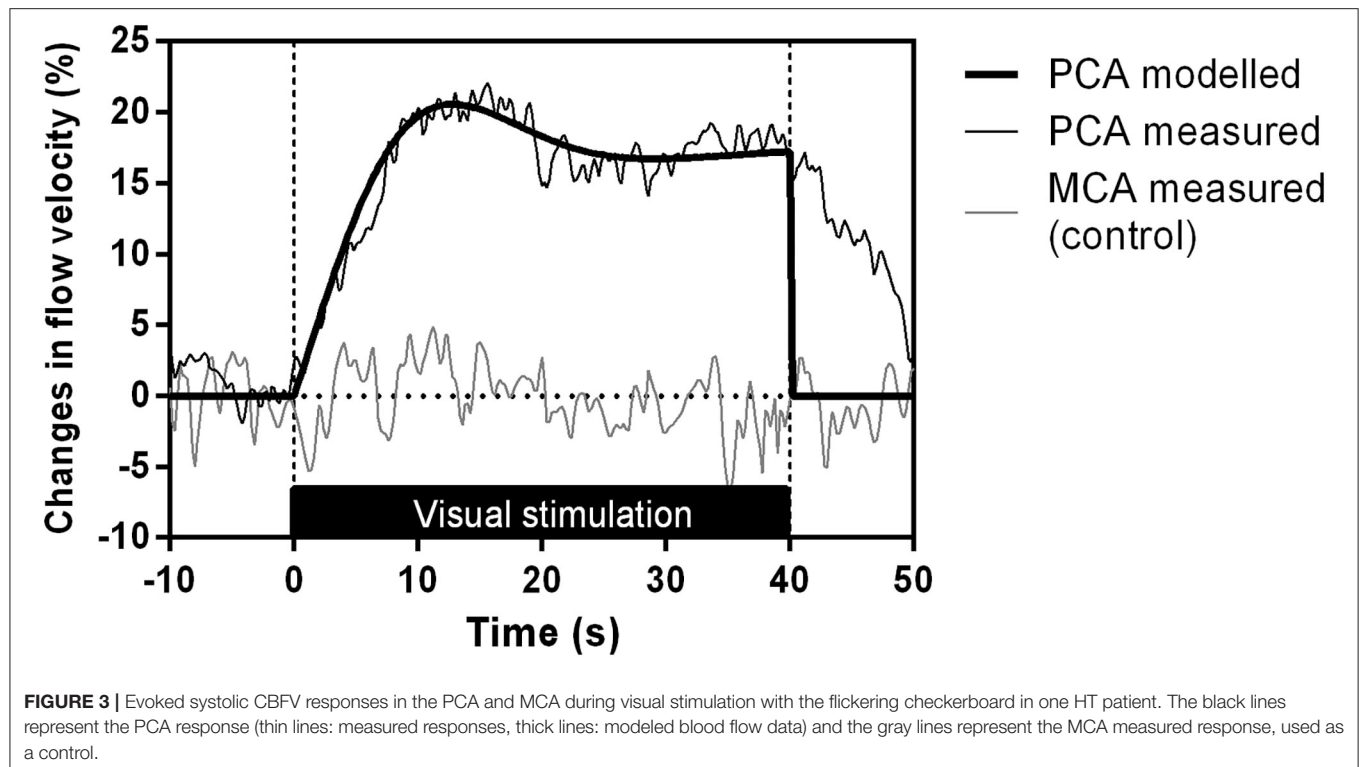


TABLE 4 | NVC and VRCO₂ in relation to the WMH volume in hypertensive patients.

	WMH volume*					
	MCA			PCA		
	≤ 0.14	> 0.14	<i>p</i> value	≤ 0.14	> 0.14	<i>p</i> value
VRCO₂ (%/mmHg CO₂)	1.6 ± 0.5	1.2 ± 0.5	0.004	1.0 ± 0.6 [‡]	0.8 ± 0.5 [‡]	0.006
Neurovascular coupling						
Overshoot [‡] systolic CBFV, (%)	-	-	-	23.4 ± 10.0 [†]	21.8 ± 8.8 [†]	0.573
Gain (%)	-	-	-	11.7 ± 11.1 [†]	14.6 ± 7.9 [‡]	0.297
Natural frequency (Hz)	-	-	-	0.2 ± 0.0 [†]	0.2 ± 0.1 [†]	0.432
Attenuation (a.u.)	-	-	-	0.4 ± 0.3 [†]	0.4 ± 0.2 [†]	0.891
Rate time (s)	-	-	-	0.0 ± 1.2 [‡]	0.0 ± 0.0 [‡]	0.169

*Dichotomized by median values. *p* values were obtained using the *t*-test. [†]Values are presented as mean ± standard deviation; [‡]Values are presented as median ± interquartile range. [‡]Maximal CBFV increase during visual stimulation. a.u., arbitrary units; CBFV, cerebral blood flow velocity; MCA, median cerebral artery; PCA, posterior cerebral artery; VRCO₂, vasoreactivity to carbon dioxide; WMH, white matter hyperintensities. Bold values of statistically significant *p* values (*p* < 0.05).

Association With White Matter Hyperintensities

As shown in **Table 4**, VRCO₂ was lower in the higher burden group in both the MCA (*p* = 0.004) and the PCA (*p* = 0.007). NVC parameters did not differ in both groups.

DISCUSSION

Our study shows that NVC was significantly impaired in hypertensive patients, with reduced CBFV increase and altered time behavior hemodynamic evoked response during visual stimulation. Moreover, NVC tended to be worse in the DM

subgroup. VRCO₂ remained relatively preserved. These results did not change when adjusting for other vascular risk factors. The novel finding is that the natural frequency seems to be the most sensitive parameter for discriminating abnormal NVC in these patients. Rosengarten and colleagues reported that this parameter had the lowest SD of the modeled parameters, thus having the potential for better differentiating between normal and abnormal NVC, but its capacity to identify vascular response dysfunction in disease settings needed to be studied (Rosengarten et al., 2001a). Our results seem to point toward natural frequency as a possible marker for NVC dysfunction in hypertension and diabetes.

Overall, the white matter disease (WMD) burden was low in the patient cohort. The NVC results were similar between

higher and lower WMD burden groups, while higher burden patients showed worse VRCO₂. It has been recently reported that reduced VRCO₂ precedes the development of WMH (Sam et al., 2016), which could help explain the difference between the disease burden groups. Similar NVC with higher and lower WMH volume has been previously reported (Sorond et al., 2013). Although WMH volume predicts an increased risk of stroke and cognitive decline, the clinical expression seems to vary (Sorond et al., 2011). Moreover, silent markers of CSVD are frequently detected on the MRI of older individuals without cognitive impairment (Vernooij et al., 2009; Debette and Markus, 2010). Thus, besides macrostructural changes, other modalities reflecting microstructural integrity and function, such as TCD dynamic studies, may provide additional information to further stratify patients at risk.

Neurovascular Unit at the Core of Target Brain Damage in Hypertension

Our results showed a significantly reduced PCA CBFV magnitude response and altered time behavior of reactive hyperemia during visual stimulation in HT patients, especially when diabetes was an added comorbidity. These findings indicate disturbed NVC in these patients' PCA cortical territory, independently of established WMD, as was observed by other groups (Birns et al., 2009; Purkayastha et al., 2014). The decreased CBFV during visual stimulation reflects less robust functional hyperemia, causing failure to meet the metabolic demands and neuronal damage (Iadecola et al., 2019).

The reduced overshoot systolic CBFV in the patient group was not accompanied by a reduction in the modeled parameter gain. This could reflect lack of statistical power to detect the differences between the groups, since the mean gain was lower in the patient group. In addition, it has been demonstrated that the overshoot of systolic CBFV is significantly influenced by not only gain, but also rate time and attenuation (Rosengarten et al., 2001a). Hence, the parameter gain and the overshoot systolic CBFV may not be perfectly matched.

These results are in line with studies on genetically determined CSVD. CADASIL patients demonstrated less robust functional hyperemia in the PCA during visual stimulation, with changes in time dynamics, very similar to our findings (Jokumsen-Cabral et al., 2019). Comparable NVC dysfunction in the posterior circulation was shown in Fabry patients (Azevedo et al., 2012; Castro et al., 2020). Both diseases are characterized by abnormal material deposition in the vessel walls (Baudrimont et al., 1993; Rombach et al., 2010). Interestingly, amyloid deposition also seems to cause NVC impairment (Iadecola and Davisson, 2008; Brickman et al., 2015), and HT appears to have a role in promoting amyloid deposition, thus working synergistically to worsen CSVD and cognitive decline (Iadecola and Davisson, 2008).

Chronic HT leads to structural (mal)adaptations in the cerebral circulation, with remodeling of the cerebral arteries and arterioles. This remodeling involves smooth muscle cell hypertrophy and hyperplasia, increased deposition of extracellular matrix components and degenerated smooth

muscle (lipohyalinosis) and fibrinoid necrosis, leading to arterial stiffening and loss of elasticity (Iadecola and Gottesman, 2019). These changes in the proximal resistance arteries cause substantial burden on the vulnerable downstream microcirculation, promoting pressure-induced oxidative stress to the endothelial cells and neuroinflammation (Ungvari et al., 2021). In experimental models, HT results in impairment of endothelium mediated neurovascular coupling responses, in part resulting from this oxidative stress and neuroinflammation (Iadecola and Gottesman, 2019; Ungvari et al., 2021). Hence, the structural changes induced by HT play an important role in the loss of functional integrity of the neurovascular unit. Natural frequency is assumed to represent the tonus and the speed of the system (Rosengarten et al., 2003), which would be altered by the increased rigidity of the vessels and endothelial dysfunction, and our results show that this parameter was the most sensitive in differentiating patients from controls.

Less effective NVC in the MCA territory has been associated with significant cognition, balance, and walking velocity changes in the elderly (Sorond et al., 2011; Purkayastha et al., 2014). Despite the presence and burden of WMD, normal NVC was associated with preserved walking speed, while slower walking is one of the earliest manifestations of CSVD. Further prospective work could use NVC for predicting symptomatic CSVD. Curiously, cocoa and deferoxamine have been demonstrated to reverse some of these changes suggesting the possibility for pharmacological modulation of the neurovascular function (Sorond et al., 2013, 2015).

Additional Contribute From Diabetes Mellitus

Comorbid diabetes associated with increased cerebrovascular dysfunction. NVC was worse in this subgroup of hypertensive patients, particularly in its oscillatory properties (natural frequency), in accordance to previous studies in early type 1 DM (Rosengarten et al., 2002). This might signify higher rigidity of the small arteries due to the accumulation of advanced glycated by-products. In fact, type 2 DM patients have particularly high incidence of lacunar stroke (van Harten et al., 2006; Brundel et al., 2014). Diabetes induced chronic vascular changes include not only macrovascular disorders, such as cardiovascular and cerebrovascular large vessel disease, but also microvascular disorders, with nephropathy, retinopathy and neuropathy (Chawla et al., 2016). Furthermore, studies have implicated DM as a risk factor for cognitive impairment, which may be related to CSVD. However, the mechanism by which cognitive decline occurs and whether it can be explained by dysfunction of the neurovascular unit remains to be elucidated (Mogi and Horiuchi, 2011). The sympathetic nervous system seems to attenuate the cerebrovascular response to hypercapnia, suggesting a direct effect on the cerebral vasculature (Jordan et al., 2000). NVC is also affected by autonomic dysfunction (Azevedo et al., 2011). Thus, differences in DM vs. nDM hypertensive patients could be related to DM associated dysautonomia in CSVD.

Overall, our study further supports cerebrovascular dynamics dysfunction as a major player in explaining the relationship between increased VRF burden and CSVD manifestations, independently of macroscopic white matter lesions.

Limitations

We acknowledge several methodological limitations. Due to the cross-section design, our study cannot provide evidence to support cerebrovascular dysfunction as an early predictor of CSVD. However, these patients had well-controlled HT, based on average ABPM values, with no clinical manifestations of cerebrovascular disease. All the patients were referred to the Hypertension Unit due to severe or difficult to manage HT, and we do not know the duration of the untreated disease, which could impact the degree of microvascular dysfunction.

Although the patient group's sample size is relatively large for hemodynamic physiology studies, this is an exploratory study and must be validated in larger, multicenter cohorts. Also, the control group is relatively small, not exactly age- and gender-matched, and the differences in comorbidities between the two groups can be potential confounders to the observed differences. Besides HT and DM, already discussed, other vascular comorbidities have been associated with the development of CSVD. Dyslipidemia plays an important role in the development of large vessel disease and stroke, but its role in CSVD is still controversial (Tsai et al., 2018). However, animal studies have demonstrated cerebral autoregulation impairment with hyperlipidemia and a relationship between hyperlipidemia and the development of CSVD (Ayata et al., 2013; Kraft et al., 2017). Obesity has been demonstrated to have an impact in the development of CSVD (Yamashiro et al., 2014), and it has been shown to affect cerebral vasoreactivity (Selim et al., 2008). Smoking appears to worsen the effects of hypertension in the cerebral microvasculature (Hara et al., 2019), and there is impaired neurovascular coupling in the PCA of young chronic smokers (Olah et al., 2008). Chronic heart failure can affect cerebral autoregulation, reduce cerebral blood flow (CBF) and has been shown to affect NVC in the PCA (Aires et al., 2020; Ovsenik et al., 2021). The prevalence of CSVD is higher in patients with coronary artery disease (Berry et al., 2019), and it has been identified as an independent risk factor for vascular dementia (Gorelick et al., 2011). It was recently demonstrated that atrial fibrillation reduces cerebral autoregulation and impairs neurovascular coupling responses to visual stimulation (Junejo et al., 2020). There is evidence that peripheral artery disease is associated with white matter disease and the development of vascular dementia (Bots et al., 1993; Gorelick et al., 2011). In patients with ischemic stroke, impaired renal function correlated with worse dCA and associated with an increased risk of hemorrhagic transformation (Castro et al., 2018). However, in addition to adjusting for age and gender, adjusting for the vascular comorbidities did not modify the results significantly.

Hypertension affects cerebral small vessels heterogeneously, and MRI seems the logical choice for its superior spatial resolution. However, MRI VRCO₂ protocols are more prone to failure (Moreton et al., 2018), expensive and not as

standardized as TCD (Malojčić et al., 2017). Moreover, TCD offers extraordinary time resolution (~5 ms) for studying the time behavior of CBFV activation in downstream cortical microvasculature. There are also inherent limitations for continuous blood flow monitoring by TCD, namely, providing a measure of blood flow velocity and not flow. However, the former is an adequate surrogate for the latter as long as the insonated artery diameter remains constant. Since data were acquired in a supine position and relied on spontaneous measurements, changes in artery diameter are not anticipated for NVC. Nevertheless, it has been demonstrated that with increasing partial pressure of arterial CO₂ there is noticeable increase in vessel diameter, which could lead to underestimation of cerebral blood flow in the VRCO₂ protocol (Coverdale et al., 2014).

We did not report the presence of cortical microinfarcts (CMI) in this study, despite having been found in cohorts of hypertensive patients (although not consistently) and despite their importance in cognitive decline (van Veluw et al., 2017). We did not detect CMI upon visual inspection of the MRI scans. However, the protocols for detecting CMI are mostly validated in 3T MRI scanners (Coverdale et al., 2014), and our 1.5T scan protocol is underpowered for their detection.

In conclusion, neurovascular coupling, and more specifically the modeled parameter natural frequency, seems to be particularly affected in hypertension and diabetes and could be useful as an early biomarker for microvascular dysfunction, future irreversible vascular damage, and cognitive decline. Additionally, our study supports TCD dynamic tests as useful tools for better understating microvascular damage associated with these diseases, but future research is warranted to confirm these results.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article may be made available by the authors upon request, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Comissão de Ética da U. L. S. Matosinhos, Hospital de Pedro Hispano, EPE, Unidade Local de Saúde de Matosinhos, Portugal. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

AM: conception and design, data collection, literature search, drafting the manuscript, and critical revision of the manuscript. PC: conception and design, data collection, critical revision of the manuscript, and supervision. GP, CF, and JP: data collection and critical revision of the manuscript. FS, AM, and JH: critical revision of the manuscript. EA: conception and design, critical revision of the manuscript, and supervision. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnagi.2021.728007/full#supplementary-material>

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Cerebral Vasoreactivity Changes Over Time in Patients With Different Clinical Manifestations of Cerebral Small Vessel Disease

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Objectives: Endothelial dysfunction (ED) has been linked to the pathogenesis of cerebral small vessel disease (SVD). We aimed to assess ED and cerebrovascular reactivity (CVR) in the patients with a diverse manifestation of SVD, with similar and extensive white matter lesions (WMLs, modified Fazekas scale grade ≥ 2), compared with a control group (CG) without the MRI markers of SVD, matched for age, gender, hypertension, diabetes, and to evaluate the change of CVR following 24 months.

Methods: We repeatedly measured the vasomotor reactivity reserve (VMRr) and breath-holding index (BHI) of the middle cerebral artery (MCA) by the transcranial Doppler ultrasound (TCD) techniques in 60 subjects above 60 years with a history of lacunar stroke (LS), vascular dementia (VaD), or parkinsonism (VaP) (20 in each group), and in 20 individuals from a CG.

Results: The mean age, frequency of the main vascular risk factors, and sex distribution were similar in the patients with the SVD groups and a CG. The VMRr and the BHI were more severely impaired at baseline (respectively, $56.7 \pm 18\%$ and 0.82 ± 0.39) and at follow-up (respectively, $52.3 \pm 16.7\%$ and 0.71 ± 0.38) in the patients with SVD regardless of the clinical manifestations (ANOVA, $p > 0.1$) than in the CG (respectively, baseline VMRr $77.2 \pm 15.6\%$, BHI 1.15 ± 0.47 , $p < 0.001$; follow-up VMRr $74.3 \pm 17.6\%$, BHI 1.11 ± 0.4 , $p < 0.001$). All the assessed CVR measures (VMRr and BHI) significantly decreased over time in the subjects with SVD (Wilcoxon's signed-rank test $p = 0.01$), but this was not observed in the CG ($p > 0.1$) and the decrease of CVR measures was not related to the SVD radiological progression ($p > 0.1$).

Conclusions: This study provided evidence that the change in CVR measures is detectable over a 24-month period in patients with different clinical manifestations of SVD. Compared with the patients in CG with similar atherothrombotic risk factors, all the CVR measures (BMRr and BHI) significantly declined over time in the subjects with SVD. The reduction in CVR was not related to the SVD radiological progression.

Keywords: neurovascular unit (NVU), cerebral small vessel disease, cerebrovascular reactivity (CVR), endothelial dysfunction, neurovascular coupling

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BACKGROUND

Small vessel disease (SVD) is one of the most important cerebral microangiopathy, responsible for the majority of lacunar stroke (LS), vascular dementia (VaD), and parkinsonism (VaP) cases (Pantoni, 2010). The basic mechanism of the cerebral vessels alterations in the SVD is linked to endothelial dysfunction (ED), but whether ED only reflects the load of atherothrombotic risk factors or if it is specific to SVD has not been clearly defined (Forsberg et al., 2018). The neurovascular unit (NVU) concept accentuates the symbiotic association between brain cells, cerebral blood vessels, and subsequently cerebral blood flow (CBF) (Muio et al., 2014). The endothelium and vascular smooth muscle within the NVU forms the basis of blood-brain-barrier (BBB), and contribute to the neurovascular coupling that is the response of the cerebral vessel to the changes in neural activity (Attwell et al., 2010). The SVD is a dynamic and progressive pathology involving variable components of NVU and BBB, however, the long-term clinical effects and outcomes usually differ between patients (Kisler et al., 2017). The recent studies have shown that the signaling pathways in the NVU control diverse processes, e.g., blood clotting and CBF, nevertheless it is not known precisely how CBF dysregulation translate to the disorders associated with neurovascular dysfunction, such as SVD. Cerebrovascular reactivity (CVR) is a measure of the capability of adaptive changes to vasodilatory stimuli (e.g., change in pCO₂ due to CO₂ inhalation, voluntary apnea, hyperventilation, or acetazolamide), and reflects the compensation of the collateral flow, therefore it can be used to indirectly assess and monitor the sequences of cerebral ED and the progression of vascular disease (Lavi et al., 2006). The reduced CVR indicates the impairment of NVU and regulatory mechanisms of CBF which result in neurovascular uncoupling. There are no biological markers to accurately assess brain vasoreactivity; however, CVR can be evaluated by various tools, such as single photon emission computed tomography (SPECT), PET, various MRI techniques, and transcranial Doppler ultrasound (TCD) (Terborg et al., 2000). Currently, one of the most widely used methods to measure the CVR is MRI, and it offers advantages over the use of radiolabeled products while maintaining regional specificity (Sleight et al., 2021). Though no direct anatomical information can be obtained, TCD permits for the evaluation of mean flow velocity (MFV) changes in the major cerebral arteries after a vasodilative stimulus and it provides complementary information of the brain hemodynamics (Ringelstein et al., 1988; Ebrahim et al., 2019; Burley et al., 2021a). The results of the TCD examination (MFV and pulsatility index) and the ventilation tests correlate with those obtained by other methods, and the value of TCD in the evaluation of CVR impairment, e.g., in the asymptomatic or symptomatic individuals with the brain white matter lesions (WMLs) has been established (Maeda et al., 1993; Marcos et al., 1997; Ghorbani et al., 2015; Fu et al., 2019). The assessment of CVR can be achieved with the bilateral recording of MFV in the middle cerebral artery (MCA) with acceptable reproducibility and inter-rater reliability using carbogen inhalation (McDonnell et al., 2013). Recently, we have found that the cerebral vasodilator

responses to breath hold and hyperventilation were abnormal in the patients with VaD, LS, and VaP caused by SVD, and they were severely impaired when compared with the controls matched for the main vascular risk factors and free from the cerebrovascular events (Staszewski et al., 2019). Most of the studies assessing CVR in SVD did not evaluate the CVR changes over time or had unmatched control groups (CGs) (Thrippleton et al., 2018). Therefore, our study aimed to investigate the CVR changes over 24 months in the subjects with diverse manifestations of SVD (LS, VaD, and VaP), with similar and extensive radiological burdens of the disease and compare with a carefully selected CG without MRI markers of SVD, free of cerebrovascular events, and matched for major vascular comorbidities.

MATERIALS AND METHODS

Participants

We analyzed the patients from the SHEF-CSVD Study in which the baseline and follow-up CVR and MRI imaging could be evaluated (Staszewski et al., 2013). The study protocol with detailed selection criteria and methodology has been described previously (Staszewski et al., 2019).

In brief: the SVD group consisted of ambulatory subjects above 60 years, enrolled between December 2011 and September 2015, with established LS (according to the OCSF criteria), VaP, or VaD (according to the Hurtig or NINDS-AIREN criteria) (Chui et al., 1992; Hurtig, 1993; Zijlmans et al., 2004). The patients with MRI contraindications, non-SVD-related WMLs, strategic single-infarct dementia, or post-stroke VaP or VaD, recurrent LS, carotid artery stenosis $\geq 50\%$, atrial fibrillation, chronic kidney disease (CKD) requiring dialysis, life expectancy < 6 months, and recent head trauma were not included. To maximize the statistical power, we decided to analyze the equal groups of subjects and we recruited the consecutive patients with VaP and matched them in a 1:1 ratio with VaD, LS patients, and CG (without known cerebrovascular disease or dementia) according to sex, age (± 5 years), and the presence of diabetes and hypertension. The signs of SVD are often seen in MRI in cognitively healthy elderly, and they are highly age-related. The studies showed that the WMLs are detected in 30–90% of cognitively healthy elderly (mean 72–74 years), therefore to maximize the homogeneity of the studied groups, we included only controls with no radiological signs of SVD in MRI (Fazekas 0) and the patients with SVD with extensive WMLs (Fazekas grade 2 or 3) (Longstreth et al., 2005; Gustavsson et al., 2015).

Of the 139 screened subjects (101 patients with SVD and 38 controls), 59 were excluded (18 controls, 15 LS, 9 VaD, and 17 VaP) due to inadequate acoustical bone window at the baseline ($n = 16$), the radiological markers of SVD in controls at baseline MRI examination ($n = 3$), withdrawal of consent during follow-up ($n = 7$), lack of follow-up MRI ($n = 22$), or TCD examination ($n = 11$). The patients with SVD and incomplete follow-up data ($n = 30$) had similar mean age (70.2 ± 7.6 vs. 72.6 ± 6.9 , $p = 0.14$) and the baseline SVD score (2.1 ± 0.6 vs. 2 ± 0.6 , $p = 0.6$) comparing with those included to the final analysis, however, more men dropped-out in comparison with women (48 vs. 9%, $p < 0.01$). The most common reported reason for lack of follow-up

was that the subjects felt asymptomatic or did not tolerate well to the baseline MRI or TCD examination. Finally, the study group comprised 60 patients with newly diagnosed symptomatic SVD (20 per group: VaP, VaD, and LS) and 20 controls. All the patients were functionally independent (modified Rankin Scale, mRS ≤ 3 and total Barthel Index ≥ 80) and without severe dementia (Mini-Mental State Examination, MMSE ≥ 12) (Schulc et al., 2015).

Study Procedures

All the patients signed informed consent before entering the study. The consent has been obtained prior to any study specific procedures. All the patients had TCD and MRI examinations performed at baseline and the 24-month follow-up visit (mean 23.1 ± 4 months; LS 22.5 ± 3 ; VaP 21.5 ± 4.6 ; VaD 23.3 ± 3.4 ; CG 23.6 ± 2 ; and ANOVA $p = 0.28$).

Ultrasound Examinations

The determination of the cerebrovascular reserve capacity is based on the ability of the intracranial arterioles to dilate. The CO₂ tests assume the correlation among the CBF, the CO₂ partial pressure, and the flow velocity in the basal cerebral arteries, and the reduction or elevation of pCO₂ leads to a decrease or increase in MFV. Since TCD measures flow velocity in the large arteries, this reflects flow in the combined gray and white matter. The TCD study included evaluation of both the MCAs with 2-MHz probes (Companion III, Nicolet) in fixed positions according to the standard protocol (Settakiss et al., 2002). The MFV values were averaged, and the interhemispheric differences for mean MFV did not exceed 15%. The recordings were considered acceptable when the velocities of blood flow could be detected bilaterally, and with a clear envelope of the MFV spectrum during the entire cardiac cycle. The CVR was measured as the breath-holding index (BHI), the ratio of the percentage MFV increase during hypercapnia, and vasomotor reactivity reserve (VMRr), the percentage change in MFV from hypo- to hypercapnia (Tsivgoulis and Alexandrov, 2008). Basing on the Markus and Harrison procedure, we measured the baseline MFV (baseMFV) following 10 min of rest with normal breathing (normocapnia), the minimal MFV value (minMFV) following 2 min of hyperventilation (hypocapnia), and the MFV value (maxMFV) subsequent to 30 s of breath holding (hypercapnia) and followed by a period of 4 min of normal breathing of room air (Markus and Harrison, 1992). To achieve the most reproducible results before proceeding to the definitive recording, the participants were trained to perform all the procedures correctly. All of them were able to hyperventilate and hold their breath for the required period. The exact length of breath-holding ranged from 29.8 to 30.5 s at the baseline and 30.1 to 30.3 s at follow-up, and it did not differ between the study groups (ANOVA, $p > 0.1$). The end-tidal CO₂ (etCO₂) concentration was monitored by capnograph (PC900A, Creative Medical, Shenzhen, China) during the examination. Blood pressure (BP) and heart rate (HR) were measured prior to and following the tests. Although McDonnell et al. found higher intra-rater reliability for the TCD measurements taken while the patients were sitting, we performed TCD examination in all the

subjects in the supine position in accordance with our standard protocol (McDonnell et al., 2013). Ultrasound examination was performed under the standardized conditions (same quiet room and time of the day; no sleep deprivation and no medication intake for at least 6 h were allowed) by a single experienced TCD sonographer unaware of the diagnosis of the subjects. All the patients with LS had the study procedures performed at least 3 weeks (mean 24 ± 2 days) after their index strokes.

MRI Evaluation (GE Healthcare 1.5 T Scanner, IL, USA)

The images were evaluated for the presence of acute LS, lacunes, deep WMLs (dWMLs), or periventricular (pWMLs), microbleeds (MBs), and enlarged perivascular spaces (PVS) according to the STRIVE guidelines and visual SVD scale (Wardlaw et al., 2013; Staals et al., 2014). The simple modified Fazekas rating scale was used to estimate the extent of pWMLs and dWMLs (Fazekas et al., 1987; Inzitari et al., 2009). The mild white matter lesions (Fazekas grade 1) were defined as the punctate lesions in the deep white matter with a maximum diameter of 9 mm for every single lesion and 20 mm for grouped lesions. The moderate white matter lesions (grade 2) were defined as early confluent lesions of 10–20 mm for single lesions and >20 mm for grouped lesions in any diameter, with no more than connecting bridges between the individual lesions. Severe white matter lesions (grade 3) were defined as single lesions or confluent areas of hyperintensity of 20 mm or greater in any diameter. The presence of each of the four MRI markers for SVD (WMH, lacunes, cerebral microbleeds, and perivascular spaces) was counted to retrieve a total SVD score (ranging from 0 to 4). One point on the visual SVD scale was awarded if confluent deep WMLs (Fazekas grade 2 and 3) or irregular periventricular hyperintensities extending into the deep white matter (Fazekas grade 3) were present or when one or more lacunes or MBs were present; 1 point was awarded if moderate (10–25) to extensive (>25) enlarged PVS were present (Kim et al., 2008). The presence of each marker produced a score of a minimum 0 and a maximum of 4, representing the total MRI load of SVD. At baseline MRI assessment, all the patients with SVD had at least Fazekas grade 2 WMLs, and controls did not have radiological markers of SVD in MRI. The visual rating of SVD radiological progression expressed by the WMLs progression or development of new lacunes was performed at a follow-up visit. As proposed by Prins et al., the WMLs progression and lacunes were rated on the FLAIR images and the presence or absence of progression was rated in the three periventricular regions, basal ganglia, infratentorial region, and four subcortical white matter regions (Prins et al., 2004). The images were reviewed by a neuroradiologist (E.S.) blinded to the clinical data.

Atherothrombotic Risk Evaluation

The vascular comorbidities were defined according to the current standards and evaluated based on the available medical data, physical examinations, widespread histories, and routine laboratory tests performed at baseline (Alberti et al., 2009; Goblirsch et al., 2013).

Statistical Analysis

All the demographic data were summarized, tabulated, and verified for normality with the Shapiro–Wilk test, and the homogeneity of the variances was assured by Levene's test. Categorical and continuous data are presented as frequencies or means \pm SD and analyzed using Fisher's exact tests, the chi-square test, paired *t*-tests, or non-parametric tests where appropriate.

The associations between MRI progression with CVR changes over time in the studied groups were compared using a linear mixed model. The model included study group (SVD, CG), Time (Baseline, Follow-up), MRI (progression, no progression), and

a Time*Group*MRI interaction for fixed effects. The subject variable was specified to assess possible individual variability. Per group, the Wilcoxon's signed-rank test was performed to evaluate the differences between CVR and other hemodynamic measures at the baseline and at follow-up visits. One-way ANOVA and chi-square tests were used to compare data between the study groups with *post-hoc* Tukey's honestly significant difference (HSD) tests for comparisons among the SVD subgroups. The effect sizes for the group differences were analyzed with partial eta squared (η_p^2) reflecting the proportion of the total variance attributable to the effect (and considered small if 0.01, moderate around 0.06, and high if >0.13). A probability value of $p < 0.05$ was

TABLE 1 | Main characteristics of the study population at baseline.

Variable	SVD (<i>n</i> = 60)	LS (<i>n</i> = 20)	VaP (<i>n</i> = 20)	VaD (<i>n</i> = 20)	CG (<i>n</i> = 20)	<i>p</i> [#]
Demographics						
Age (y)*	72.6 (± 6.9)	71.7 (± 7.7)	73.8 (± 6.2)	72.2 (± 6.8)	71.9 (± 3.2)	0.7
Female gender (%)*	30 (50)	10 (50)	10 (50)	10 (50)	10 (50)	1
Vascular risk factors						
Hypertension (%)*	54 (89)	18 (90)	18 (90)	18 (90)	18 (90)	1
Diabetes (%)*	30 (50)	10 (50)	10 (50)	10 (50)	10 (50)	1
CAD (%)	22 (37)	8 (40)	7 (35)	7 (35)	7 (35)	0.2
Current smoking (%)	22 (37)	7 (35)	6 (30)	9 (45)	8 (40)	0.5
Hyperlipidemia (%)	45 (75)	15 (75)	16 (80)	14 (70)	15 (75)	0.8
CKD (%)	8 (13)	2 (10)	2 (10)	4 (20)	3 (15)	0.3
PS (%)	28 (46)	10 (50)	10 (50)	8 (40)	8 (40)	0.8
Obesity (BMI > 30)	17 (28)	6 (30)	6 (30)	5 (25)	6 (30)	0.8
Examination findings						
Homocysteine ($\mu\text{mol/l}$)	14.4 \pm 6.1	13.2 \pm 3.7	18.1 \pm 8.3	12.03 \pm 3.5	13.1 \pm 2.7	0.01
hsCRP (mg/dL)	0.55 \pm 0.9	0.56 \pm 1.02	0.64 \pm 0.9	0.46 \pm 0.8	0.16 \pm 0.17	0.2
Uric acid (mg/dL)	5.4 \pm 1.5	5.8 \pm 1.6	5.5 \pm 1.3	4.8 \pm 1.5	4.6 \pm 1.3	0.05
eGFR (ml/min/1.73 m ²)	78.8 \pm 24.1	77.1 \pm 29	72.3 \pm 22.8	87 \pm 17.2	95 \pm 16	0.01
HbA1c (%)	6.2 \pm 1.03	6.3 \pm 1.2	5.9 \pm 0.5	6.3 \pm 1.2	5.9 \pm 0.1	0.26
FG (mg/dL)	120.1 \pm 43.2	128.8 \pm 48.6	112.8 \pm 32.3	120.6 \pm 47.2	106.3 \pm 18.7	0.21
LDL (mg/dL)	108.4 \pm 37.8	97.1 \pm 35.5	110.6 \pm 36.6	116.5 \pm 40.4	122.8 \pm 41.8	0.22
HDL (mg/dL)	52.2 \pm 17	46.7 \pm 10	55.2 \pm 16.9	54.1 \pm 22.5	60.9 \pm 19.3	0.12
TG (mg/dL)	124.5 \pm 57.9	125.6 \pm 60.1	132.1 \pm 64.2	115.5 \pm 50	115.1 \pm 39	0.7
TC (mg/dL)	185.2 \pm 48.1	166.8 \pm 39.5	194.4 \pm 54.7	193.4 \pm 54.7	198.7 \pm 31.7	0.1
BMI	26.4 \pm 5.6	27.7 \pm 7.6	25.1 \pm 4	26.3 \pm 4.5	27.8 \pm 3.8	0.3
MRI examination						
SVD score (mean \pm SD)	2 \pm 0.6	2.2 \pm 0.7	2 \pm 0.6	2.1 \pm 0.6	0	<0.01
Fazekas pWMLs (mean \pm SD)	1.45 \pm 0.9	1.5 \pm 1.1	1.4 \pm 1	1.5 \pm 0.7	0	<0.01
Fazekas dWMLs (mean \pm SD)	1.6 \pm 0.9	1.7 \pm 1.1	1.5 \pm 0.8	1.7 \pm 0.9	0	<0.01
Vascular treatment						
Statin treatment (%)	45 (75)	15 (75)	16 (80)	14 (70)	15 (75)	0.8
ACEI (%)	15 (25)	6 (30)	4 (20)	5 (25)	7 (35)	0.6
Beta blocker (%)	22 (37)	8 (40)	7 (35)	7 (35)	7 (35)	0.2
ARB (%)	19 (32)	6 (30)	7 (35)	6 (30)	7 (35)	0.4
Calcium channel blocker (%)	23 (38)	9 (40)	7 (35)	7 (35)	7 (35)	0.4
Diuretics (%)	12 (20)	4 (20)	3 (15)	5 (25)	4 (20)	0.2

Values represent the means (\pm SD) for continuously distributed data or the numbers (%) for categorical data.

*Matched factors; #ANOVA and χ^2 difference between the groups $p < 0.05$ compared with CG according to Tukey's honestly significant difference (HSD) test.

SVD, cerebral small vessel disease; LS, lacunar stroke; VaD, vascular dementia; VaP, vascular parkinsonism; CG, control group; CAD, coronary artery disease; BMI, body mass index; FG, fasting glucose; PS, polymeric syndrome; CKD, chronic kidney disease; IMT, intima-media thickness; ACEI, angiotensin-converting-enzyme inhibitors; ARB, angiotensin receptor blockers.

considered significant, all the analyses were made using the PQStat 1.8.0 software.

The study has been approved by the Internal Revision Board (Wojskowy Instytut Medyczny, 46/WIM/2010) and it was conducted in accordance with the Declaration of Helsinki.

RESULTS

The studied cohort consisted of 80 older adults (mean 72.4 ± 6.2 years) of both sexes (50% women). The baseline characteristic of the studied cohort is provided in **Table 1**. The prevalence of the main comorbidities was similar in groups, but there were some differences in the laboratory examination findings: the CG had lower concentrations of homocysteine, showed a trend toward lower concentration of uric acid, and had higher levels of estimated glomerular filtration rate (eGFRs) comparing with other subjects. The SVD subgroups had a similar prevalence of vascular risk factors and treatment use, laboratory, and MRI findings (Fazekas grade 3 was scored by 80% LS, 80% VaD, and 75% VaP subjects; $p = 0.7$). The mean etCO_2 concentrations and systolic blood pressure (SBP), diastolic blood pressure (DBP), and heart rate (HR) measures during hyperventilation and breath holding did not differ between the CG and the SVD group, and they were also similar between the SVD subgroups. In addition, these variables did not significantly differ between the baseline and follow-up assessments (Wilcoxon's sign rank test, $p > 0.1$). The reproducibility measurements of VMRr and BHI were performed in a sample of 12 individuals (four from CG,

eight with LS). The intraclass correlation coefficient for the two consecutive measurements was 0.90; $p < 0.01$ for VMRr and 0.88; $p < 0.01$ for BHI.

Vasomotor reactivity reserve and the BHI were more severely impaired at the baseline and at follow-up assessments in the patients in the SVD groups (regardless of the clinical spectrum of the disease) than in the CG (**Table 2**). The size effect of the difference was similar in all comparisons (eta-squared ranged between 0.25 and 0.29). For better visualization, the CVR change in the individual subjects is shown in (**Supplementary Figures 1, 2**).

Vasomotor reactivity reserve and the BHI were significantly lower in diabetic patients with SVD vs. diabetic controls at the two timepoints (baseline VMRr: 55.8 ± 16 vs. $74 \pm 18\%$, respectively, $p < 0.001$; BHI: 0.79 ± 0.39 vs. 1.1 ± 0.3 , $p = 0.02$; follow-up VMRr 52.7 ± 15 vs. 71.2 ± 20 , $p < 0.01$; BHI: 0.72 ± 0.4 vs. 1.04 ± 0.5 , $p < 0.01$ and the difference was also significant between the non-diabetic subjects from SVD group vs. CG (respectively; baseline VMRr: 54.2 ± 18.4 vs. 79.9 ± 13.7 , $p < 0.001$; BHI: 0.78 ± 0.39 vs. 1.16 ± 0.55 , $p = 0.01$; follow-up VMRr 52.1 ± 18 vs. $76.8 \pm 14.9\%$, $p < 0.01$; BHI: 0.70 ± 0.35 vs. 1.15 ± 0.2 ; $p < 0.001$).

Both the assessed CVR measures (VMRr and the BHI) significantly declined over time in the subjects with SVD (Wilcoxon's signed-rank test, $p = 0.01$), however, there was no significant decline of VMRr and BHI in CG over 24 months of observation (Wilcoxon's signed-rank test $p > 0.1$) (**Figure 1**). The mean VMRr and BHI values were also similar at the baseline and follow-up measurements

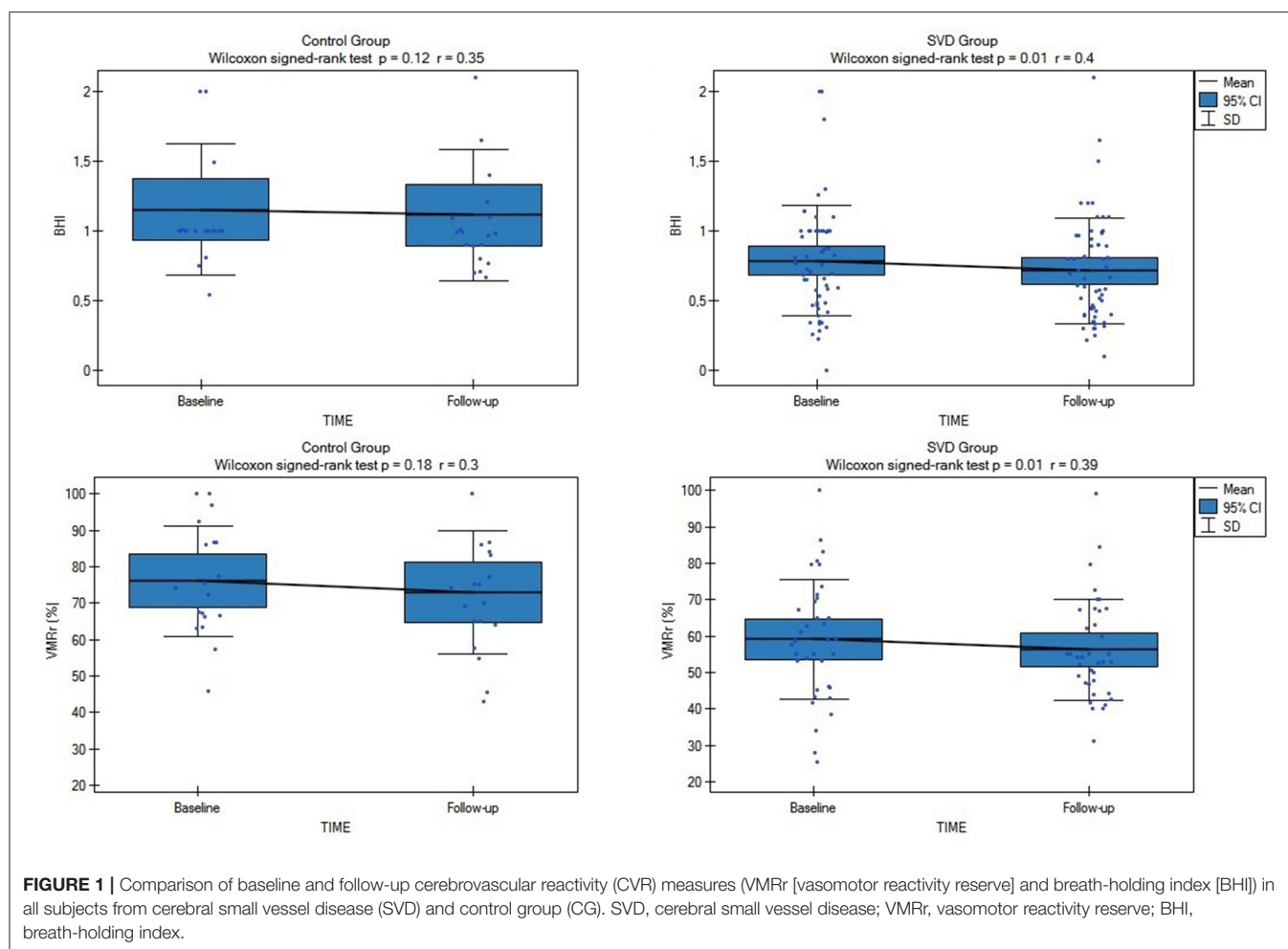
TABLE 2 | Basal and follow-up characteristics and vasodilatation responses of cerebral arteries in all the subjects.

Variable	SVD (n = 60)	LS (n = 20)	VaP (n = 20)	VaD (n = 20)	CG (n = 20)	p [#]
Baseline measures						
MFV (cm/s)	$41.5 \pm 12^*$	43.9 ± 15	42 ± 9.5	$38.4 \pm 10.4^*$	46.8 ± 8.7	0.03
VMRr (%)	$56.7 \pm 18.4^*$	$55.4 \pm 18.6^*$	$55.1 \pm 15.2^*$	$54.2 \pm 18.7^*$	77.2 ± 15.6	<0.001
BHI	$0.82 \pm 0.39^*$	$0.86 \pm 0.4^*$	$0.77 \pm 0.28^*$	$0.71 \pm 0.4^*$	1.15 ± 0.47	<0.001
Resting SBP, mmHg	127.2 ± 17.2	130.1 ± 16.2	126.1 ± 11.1	125.2 ± 16	124 ± 19	0.2
Resting DBP, mmHg	73.4 ± 10.1	72.4 ± 13.2	73.2 ± 12.4	74.4 ± 11	74.1 ± 8	0.7
Resting HR, beats/min	76.8 ± 6.6	76.8 ± 4.7	75.2 ± 8.2	78.3 ± 6.6	72.1 ± 6.1	0.8
Delta etCO_2 HV (%)	1.3 ± 0.15	1.4 ± 0.4	1.3 ± 0.2	1.1 ± 0.7	1.4 ± 0.5	0.6
Delta etCO_2 BH (%)	1.2 ± 0.1	1.3 ± 0.2	1.3 ± 0.6	1.1 ± 0.6	1.2 ± 0.4	0.8
Follow-up measures						
MFV (cm/sek)	$35.2 \pm 11.2^*$	37.1 ± 15.6	$34.3 \pm 9.2^*$	$34.2 \pm 8.3^*$	43.1 ± 8.2	0.06
VMRr (%)	$52.3 \pm 16.7^*$	$54.4 \pm 17.9^*$	$49.3 \pm 11.5^*$	$53.2 \pm 19.7^*$	$74.3 \pm 17.6^*$	<0.001
BHI	$0.71 \pm 0.38^*$	$0.72 \pm 0.5^*$	$0.71 \pm 0.24^*$	$0.69 \pm 0.4^*$	$1.11 \pm 0.4^*$	<0.001
Resting SBP, mmHg	124.8 ± 13.9	127.2 ± 17.1	125.3 ± 9.8	122.1 ± 15	127 ± 9	0.3
Resting DBP, mmHg	76.3 ± 11	74.1 ± 14.1	78.1 ± 7.7	76.7 ± 11	74.2 ± 10	0.6
Resting HR, beats/min	73.9 ± 2.2	72.6 ± 3	75.2 ± 1.2	74.1 ± 2.4	70 ± 10.5	0.2
Delta etCO_2 HV (%)	1.23 ± 0.6	1.3 ± 0.7	1.2 ± 0.5	1.2 ± 0.6	1.9 ± 0.7	0.3
Delta etCO_2 BH (%)	1.16 ± 0.7	1.1 ± 0.8	1.1 ± 0.4	1.3 ± 0.4	1.2 ± 0.7	0.8

[#]ANOVA difference between the groups, * $p < 0.05$ compared with the CG.

SVD, cerebral small vessel disease; LS, lacunar stroke; VaD, vascular dementia; VaP, vascular parkinsonism; CG, control group; MFV, mean flow velocity; BHI, breath holding index; VMRr, vasomotor reactivity reserve; PI, pulsatile index; RI, resistance index; SBP, systolic blood pressure; DBP, diastolic blood pressure; HR, heart rate.

$\Delta \text{etCO}_2 \text{HV}$, $\Delta \text{end-tidal CO}_2$ during hyperventilation; $\Delta \text{etCO}_2 \text{BH}$, $\Delta \text{end-tidal CO}_2$ during breath holding.



among the patients with LS, VaD, and VaP (ANOVA, $p > 0.1$).

Radiological progression during the study was observed in 28 of the 80 subjects (35%): significantly more often ($p = 0.03$) in the SVD subjects ($n = 25$ [42%]: eight subjects from LS [40%] group, eight with VaD [40%], and nine with VaP [45%], with no difference between the SVD groups [ANOVA $p > 0.1$]) than in the CG ($n = 3$; 15%). During the follow-up, only 4/80 patients (5%) experienced lacunar stroke (two patients from LS group and two from VaD group), no other vascular events were observed. Although in patients with SVD who experienced radiological progression, all the CVR measures were considerably impaired at the baseline and follow-up assessments compared with the no progression group (respectively, baseline VMRr $43.3 \pm 13.2\%$ vs. $63.3 \pm 14.9\%$, $p < 0.001$; eta-square 0.33; baseline BHI 0.63 ± 0.39 vs. 0.9 ± 0.36 , $p < 0.001$; eta-square 0.12; follow-up VMRr $41.2 \pm 13.4\%$ vs. $60.3 \pm 14\%$, $p < 0.001$; eta-square 0.32; BHI 0.52 ± 0.24 vs. 0.84 ± 0.39 , $p < 0.001$; eta-square 0.19), no significant decrease of CVR measures was observed during 24 months of observation (Wilcoxon's signed-rank test, $p > 0.1$) (Figure 2). These data did not change significantly when we analyzed all the patients (SVD and CG) with the radiological progression vs. no progression group (respectively, baseline VMRr $45.4 \pm 14.1\%$

vs. $68.6 \pm 16.9\%$, $p < 0.001$; baseline BHI 0.65 ± 0.37 vs. 1.0 ± 0.42 , $p < 0.001$; follow-up VMRr $43.1 \pm 14.3\%$ vs. $65.7 \pm 16.9\%$, $p < 0.001$; BHI 0.55 ± 0.24 vs. 1.0 ± 0.47 , $p < 0.001$) and there was also no significant decrease of CVR measures over time of observation (Wilcoxon's signed-rank test, $p > 0.1$). No interaction between CVR measures, time of assessment, and radiological progression in the SVD group were observed in the fixed linear analysis (Table 3).

DISCUSSION

This prospective study performed in symptomatic SVD has shown that the change in CVR measures is detectable following a 24-month period and that 42% of patients with SVD demonstrated radiological progression. We also confirmed that the cerebral vasoreactivity measures decreased in SVD regardless of the clinical manifestation of the disease. On the opposite, we did not demonstrate a significant CVR decrease in the subjects from CG who were neurologically asymptomatic, had no radiological markers of SVD in the baseline MRI, but who shared similar atherothrombotic risk factors as the SVD group. Although patients with SVD and radiological progression had

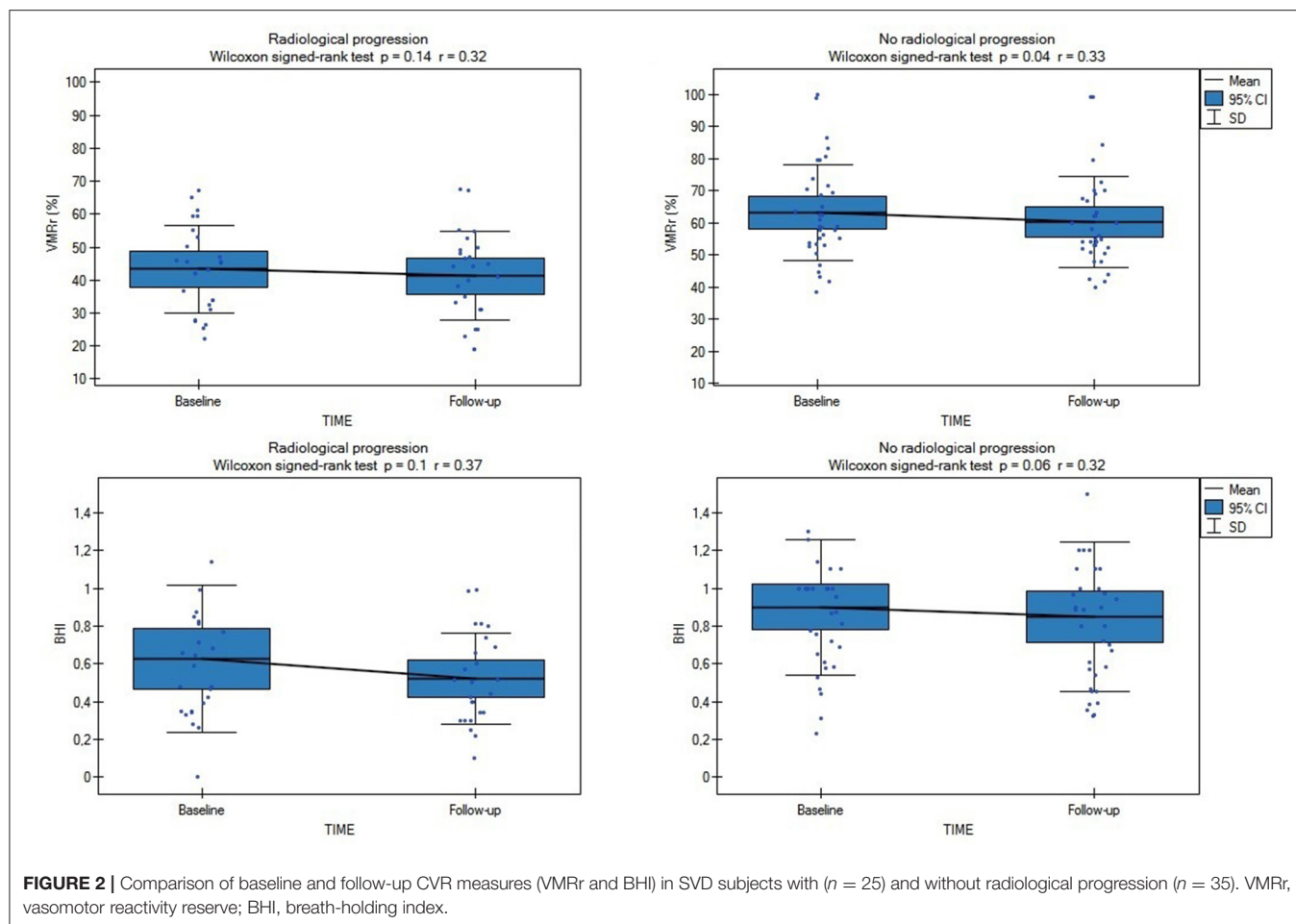


TABLE 3 | Association among the cerebral small vessel disease (SVD), assessment time point, and MRI progression on the cerebrovascular reactivity (CVR) measures.

Dependent variable	Fixed factor(s)	B	SE	t	P*	95% CI	
						Lower limit	Upper limit
BHI	Time	−0.01	0.05	−0.01	0.90	−0.11	0.11
	SVD	−0.47	0.11	−4.26	<0.01	−0.69	−0.25
	MRI progression	−0.60	0.22	−2.70	0.01	−1.04	−0.16
	Time*SVD	0.05	0.07	0.55	0.50	−0.20	0.38
	Time*MRI progression	0.08	0.18	1.20	0.24	−0.15	0.59
	SVD*MRI progression	0.29	0.24	1.23	0.22	−0.18	0.77
	Time*SVD*MRI progression	−0.02	0.16	−0.13	0.80	−0.34	0.30
VMRr	Time	−3.90	2.60	−1.40	0.20	−9.20	1.30
	SVD	−27.10	4.34	−6.20	<0.01	−35.70	−18.50
	MR progression	−28.80	9.20	−3.10	0.01	−47.10	−10.50
	Time*SVD	7.05	3.23	2.13	0.20	−0.20	5.97
	Time*MRI progression	9.04	6.80	1.32	0.20	−4.50	22.60
	SVD*MRI progression	9.80	9.90	0.90	0.30	−9.98	29.60
	Time*SVD*MRI progression	−10.07	7.40	−1.35	0.17	−24.80	6.70

*Linear mixed effects model.

BHI, breath holding index; VMRr, vasoreactivity reserve; SVD, small vessel disease.

TABLE 4 | Major transcranial Doppler ultrasound (TCD) studies assessing CVR in sporadic small vessel disease populations.

References	Study population	CVR assessment	Results
Molina et al. (1999)	46 patients with lacunar stroke and 46 CG	Acetazolamide test	Significant CVR reduction in SVD group
Terborg et al. (2000)	46 patients with SVD and 13 CG	Ventilation tests; NIRS	Significantly reduced VMR in TCD and NIRS assessments in severe SVD; a good correlation between the validity of TCD and NIRS
Kidwell et al. (2001)	55 patients with SVD	PI	Correlation between PI and severity of SVD ; PI cut points allowed discrimination of PVH with 89% sensitivity and 86% specificity and discrimination of DWMH with 70% sensitivity and 73% specificity.
Pánczél et al. (2002)	25 patients with lacunar stroke, 20 patients with leukoaraiosis and 35 CG	Tilting, ventilation, and acetazolamide tests	CVR in BA and MCA territory was impaired in hypercapnia in SVD; no significant differences between CVR measures in BA and MCA territory in acetazolamide test; significantly higher VRCO ₂ in MCA vs BA measurements
Ghorbani et al. (2015)	56 patients with SVD and 48 CG	PI	Good correlation between PI and SVD radiological manifestations: <ul style="list-style-type: none"> - In PVH with PI cut-off = 0.83, the sensitivity 90% and specificity 98% - In DWMH with PI = 0.79, the sensitivity 75% specificity 87.5% - In lacunar stroke with PI = 0.80, the sensitivity 73% and specificity 90%. - In PH with PI = 0.69, the sensitivity 92% and specificity 87.5%. - In PVH+ DWMH+ lacunar with PI = 0.83, the sensitivity 90% and specificity 96%.
Nam et al. (2020)	206 patients with lacunar stroke	PI	PI was positively associated with the WMHs volume and the presence of old lacunar infarcts.

TCD, transcranial Doppler ultrasound; PI, pulsatile index; MRI, magnetic resonance imaging; SVD, cerebral small vessel disease; CG, control group; PVH, periventricular hyperintensity; DWMH, deep white matter hyperintensity; CVR, cerebrovascular reactivity; BA, basilar artery; MCA, middle cerebral artery; VRCO₂, ventilatory response to CO₂; PH, pontine hyperintensity; NIRS, near-infrared spectroscopy; CT, computed tomography.

severely impaired CVR at the baseline and follow-up comparing with the subjects with no MRI progression, there was no significant interaction between that CVR decline and risk of radiological progression.

Besides of growing evidence of MRI techniques which increasingly play a more important role as a non-invasive tool capable to assess the cerebral reserve capacity in combination with a vascular challenge, the vasomotor reactivity testing with TCD is still a gold diagnostic standard and it has been proved to assess CBF and indirectly monitor the function of ED in patients with cerebral microangiopathy (Kozera et al., 2009; D'Andrea et al., 2016). The TCD studies demonstrated that CVR is negatively correlated with the duration of hypertension, patient age, and history of cerebrovascular events (Fujishima et al., 2001; Kozera et al., 2010). Some older studies using SPECT or Xenon CT techniques revealed that LS, particularly with leukoaraiosis was associated with the reduced CBF in the white matter and cortex in the subjects with severe WMLs, while the studies on VaD consistently showed reductions in both the white and gray matter (Markus et al., 2000). Many studies investigated CBF in the patients with SVD using TCD techniques, and in line with our data, the majority of them revealed impaired CBF comparing with the healthy controls in the patients with symptomatic and asymptomatic LS, more marked in multiple comparing with single infarcts and a reduction in reactivity that was correlated with the degree of leukoaraiosis (Table 4) (Maeda et al., 1993; Molina et al., 1999; Terborg et al., 2000). There is, however, a lack of studies on the dynamic CVR changes over time, especially in SVD, and in well-characterized SVD populations, e.g., in rarely studied patients with VaD and VaP. As SVD is a wide term and

it is used to describe the different disease processes ranging from asymptomatic WMLs in normal individuals through to the symptomatic SVD subjects, as we aimed in the presented report, our data added some further evidence that cerebral ED occurs and progresses in patients with LS, VaD, or VaP due to SVD, even if they had well-controlled hypertension (according to the baseline and follow-up BP measurements). Interestingly, a similar trend was not observed in CG with comparable atherothrombotic risk factors (age, sex, diabetes, hypertension, obesity, and smoking), and this may suggest that other vascular factors or duration of exposure to vascular risk factors in SVD are more important and responsible for a continuous decline in the CVR (Muio et al., 2014). For example, the studies on patients with cerebrovascular disease showed that the reduction in vasodilatory capacity in asymptomatic carotid disease can predispose for the development of cerebrovascular disease (Silvestrini et al., 2000). In 2020, Soman et al. (2020) using arterial spin labeling MRI to evaluate which cardiovascular risk factors alter the CVR found that higher SBP, chronic kidney disease, history of past stroke, and hypercholesterolemia are responsible for the impaired CVR. Although the controls and patients with SVD in our study were well-balanced with the main comorbidities, we did not control other important risk factors of SVD. For instance, chronic poor sleep patterns, depression, prediabetes, and unhealthy diet (dietary salt) can impact the endothelial function through immune influences (Hakim, 2019). Of notice, the Oxford Vascular (OXVASC) Study importantly showed that premorbid hypertension in midlife correlated more strongly with the global SVD burden than a single blood pressure measurement or a known history of hypertension (Lau et al., 2018).

Whether the progressive decrease of CVR is specific to the progressive nature of SVD is unknown (Stevenson et al., 2010). Importantly, we showed that there was no difference in the CVR at the baseline and follow-up assessments between the SVD clinical subgroups and this finding possibly could be related to similar WMLs burden or comparable prevalence of comorbidities. We also observed that the subjects with radiological progression had the most impaired CVR at baseline which, however, did not significantly decrease over time. These findings may indicate that MRI progression in that group may be related to the baseline severe CVR impairment or other factors, e.g., inflammatory or prothrombotic. It is also possible that the TCD measures are less sensitive to the more subtle changes, particularly in the white matter flow. Our results are, however, opposite to a large study in 628 asymptomatic individuals (mean 69 years) with WMLs which showed a progressive reduction in the CBF velocity measured using TCD as WMLs severity increased (Tzourio et al., 2001). Some similar data were demonstrated by Sam et al., who established areas of reduced CVR that preceded the development of WMLs by advanced MRI techniques, suggesting that CVR impairment contributes to the development and progression of SVD (Sam et al., 2016). On the opposite, in a PROSPER trial of statin therapy that investigated 390 individuals with cardiovascular risk factors, there was no association between CBF and the prevalence of WMLs at baseline, a decline of CBF, and risk of development of deep WMLs (Ten Dam et al., 2007). Our results are also in line with the study that showed significantly reduced CVR in LS than in the CG, which, however, could not be explained by the main atherothrombotic risk factors (Deplanque et al., 2013). It is important to appreciate that cerebral reactivity and autoregulation are not fully understood, although the neurogenic, metabolic, or myogenic factors were proposed. Of notice, CO₂-CVR is only one of the several mechanisms of cerebral autoregulation, thus preserved CO₂-CVR does not imply intact cerebral autoregulation (Klinzing et al., 2021). Therefore, CVR impairment could be due to different mechanisms, e.g., related to genetic factors or BBB dysfunction, resulting in microvascular dysfunction (Zlokovic, 2011). Some important aspects of the present study related to the patients with VaD and VaP should be underlined as these subjects probably share SVD pathology with neurodegenerative disease, and it is not established whether the microvascular changes reflect the diminished demand caused by the advanced neurodegeneration or whether SVD precedes and contributes to the neurodegeneration. For example, impaired cerebrovascular reserve capacity and vasoreactivity (identified using TCD along with the BHI) was found in the patients with mild cognitive impairment, Alzheimer's disease, and Parkinson's disease without severe underlying atherosclerosis (Shim et al., 2015; Urbanova et al., 2018). Although the contribution of CVR impairment to the pathogenesis of neurodegenerative diseases is not certain, it might be suspected that a reduced cerebrovascular reserve is an additional deteriorating factor in the neurodegeneration (Ojeda et al., 2017). One unanswered key question is that whether any reductions in the CBF are primary, or merely occurs secondary to the brain damage as due to vaso-neuronal coupling, the

reductions in brain metabolism are associated with reduced CBF (Markus et al., 2014).

Most studies in the literature have demonstrated the differences in CBF between the groups of patients with SVD and controls, with very few studies addressing the sensitivity and specificity of these parameters in TCD and MRI examinations (Panerai, 2009). The large variations in CVR across the patients and between different sessions of the same subject are often observed in the TCD and MRI studies, which hamper the ability of these measures in monitoring disease progression (Hou et al., 2020). It should be especially stressed that there are concerns with the application of the BHI as the relationship between breath-hold length and the pCO₂ stimulus remains unclear and this test has low reproducibility and high variability (Alwatban et al., 2018; Koep et al., 2020). The latest studies showed that the blood oxygenation level dependent MRI (BOLD-MRI) method is most reliable and reproducible because it has the advantage of mapping the whole brain with good spatial resolution, allowing investigation of CVR regional distribution (Leoni et al., 2012). Despite the relationships between the baseline CBF measures from TCD and MRI, the current studies showed no direct correlations between the CVR metrics calculated from TCD and MRI-BOLD measures during a 5%CO₂ challenge (Burley et al., 2021b). These variations in the CVR measures in different imaging modalities are important as they significantly reduce the statistical power to detect the pathology-related differences, preclude personalized determination of abnormalities, and challenge the validity of comparing CVR metrics across the studies.

Our study has several limitations, and the data should be cautiously interpreted owing to the single-center design and a small number of subjects included that may introduce bias. Even if we analyzed a relatively homogeneous cohort and highly selected SVD sample of patients, it should be considered a hypothesis-generating pilot study, because it may be less representative of the overall SVD population. However, the majority of the TCD studies on CVR in subjects with SVD had low patient numbers or controls (Table 4). Although the patients were prohibited from taking any drugs before the examinations, some medications could have affected the dilatatory responses of the arteries. In future studies, additional research may be needed to determine the effect of treatment as a function of CBF on the prognosis of patients. The other limitations are that we used 1.5 Tesla MRI which potentially limited the accuracy of the radiological assessment, and we did not account for etCO₂ for calculating the CVR. This could have influenced the results and limit the generalizability and reproducibility of the study, since PaCO₂ has a marked influence on CBF and autoregulation, evaluations could be compromised in situations where significant changes in PaCO₂ are undetected (Panerai et al., 1999). However, in our study, no significant difference between the subgroups and between the baseline and follow-up etCO₂ measures were noticed. Despite the multitude of alternatives to the gas challenges, existing literature lacks definitive conclusions regarding the best practices for the vasoactive modulation and associated analysis protocols for the TCD studies resulting in numerous metrics of cerebral

autoregulation (Valdúez et al., 2008; Sanders et al., 2018). Regardless of methodology, an assessment of cerebrovascular autoregulation is prone to moderate noise and artifact, with low reliability and reproducibility (Lee et al., 2020). In our study, CVR was assessed using TCD which additionally suffers from a limited field of view comparing with the MRI techniques and thus it may be less reflective of the local changes in the tissue blood supply. However, due to limited access and the high cost of MRI examination, the non-invasive, bedside, and acceptable repeatable assessments using TCD, it remains still the most utilized tool to study the CBF regulation in humans. As different techniques have low correlations and target different parts of the vascular tree, they should, however, be regarded as complementary and they are recommended to be used together (Purkayastha and Sorond, 2012; Burley et al., 2021b). Another important limiting factor is that the hemodynamic effect of breath holding is lower than that of CO₂ inhalation or acetazolamide injection, therefore the validity of our results should be confirmed in the future (Marcic et al., 2021). The TCD measurements are limited to the large basal arteries and can only provide an index of global rather than local CBF velocity, and it is highly operator dependent. However, all the TCD tests in our study were carried out by a single certified examiner who has 15 years of experience working with TCD, and thus we have minimized a possible interpersonal difference depending on the experience of the examiner.

The present study has, however, some important strengths. To the best of our knowledge, no data have been published on repeatedly assessed CVR measures over 24 months of observation in a well-phenotyped cohort of patients with SVD with extensive radiological markers of SVD. Ours is the first study to investigate all CVR by TCD in a wide range of participants, such as normal controls and patients with LS, VaD, and VaP. Our findings of impaired and progressive CVR alterations in the SVD subjects are important as this mechanism could contribute to exacerbating the clinical condition and may constitute a potential line of research for the treatment. A better understanding of the variations in CVR is of importance in both clinical and basic science applications (Hou et al., 2020). The majority of data on CVR are derived from the studies on individuals with asymptomatic WMLs, therefore, our study, which included homogenous groups of symptomatic SVD subjects, is important (Blair et al., 2016).

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- In conclusion, we provided further evidence that cerebral ED occurs and progresses over time in the patients with different clinical manifestations of SVD, however, we did not observe a significant CVR difference between the subjects with SVD and radiological progression comparing with no progression group and no significant CVR alterations over time in the CG who were neurologically asymptomatic and who shared similar comorbidities to the SVD group. The longitudinal studies with larger sample sizes are needed to definite the causal conclusions and confirm these findings.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Local Medical Ethics Committee, Military Institute of Medicine, Warsaw, Poland (46/WIM/2010). The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

JS: conceptualization, methodology, validation, data curation, and project administration. JS and ES: investigation. JS and AD: writing—original draft preparation. JS and AS: supervision. All authors have read and agreed to the published version of the manuscript.

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From Neurodevelopmental to Neurodegenerative Disorders: The Vascular Continuum

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Structural and functional integrity of the cerebral vasculature ensures proper brain development and function, as well as healthy aging. The inability of the brain to store energy makes it exceptionally dependent on an adequate supply of oxygen and nutrients from the blood stream for matching colossal demands of neural and glial cells. Key vascular features including a dense vasculature, a tightly controlled environment, and the regulation of cerebral blood flow (CBF) all take part in brain health throughout life. As such, healthy brain development and aging are both ensured by the anatomical and functional interaction between the vascular and nervous systems that are established during brain development and maintained throughout the lifespan. During critical periods of brain development, vascular networks remodel until they can actively respond to increases in neural activity through neurovascular coupling, which makes the brain particularly vulnerable to neurovascular alterations. The brain vasculature has been strongly associated with the onset and/or progression of conditions associated with aging, and more recently with neurodevelopmental disorders. Our understanding of cerebrovascular contributions to neurological disorders is rapidly evolving, and increasing evidence shows that deficits in angiogenesis, CBF and the blood-brain barrier (BBB) are causally linked to cognitive impairment. Moreover, it is of utmost curiosity that although neurodevelopmental and neurodegenerative disorders express different clinical features at different stages of life, they share similar vascular abnormalities. In this review, we present an overview of vascular dysfunctions associated with neurodevelopmental (autism spectrum disorders, schizophrenia, Down Syndrome) and neurodegenerative (multiple sclerosis, Huntington's, Parkinson's, and Alzheimer's diseases) disorders, with a focus on impairments in angiogenesis, CBF and the BBB. Finally, we discuss the impact of early vascular impairments on the expression of neurodegenerative diseases.

Keywords: cerebrovascular abnormalities, neurodevelopment and intellectual disabilities, aging, neurodegeneration, cerebral blood flow, angiogenesis, blood-brain barrier

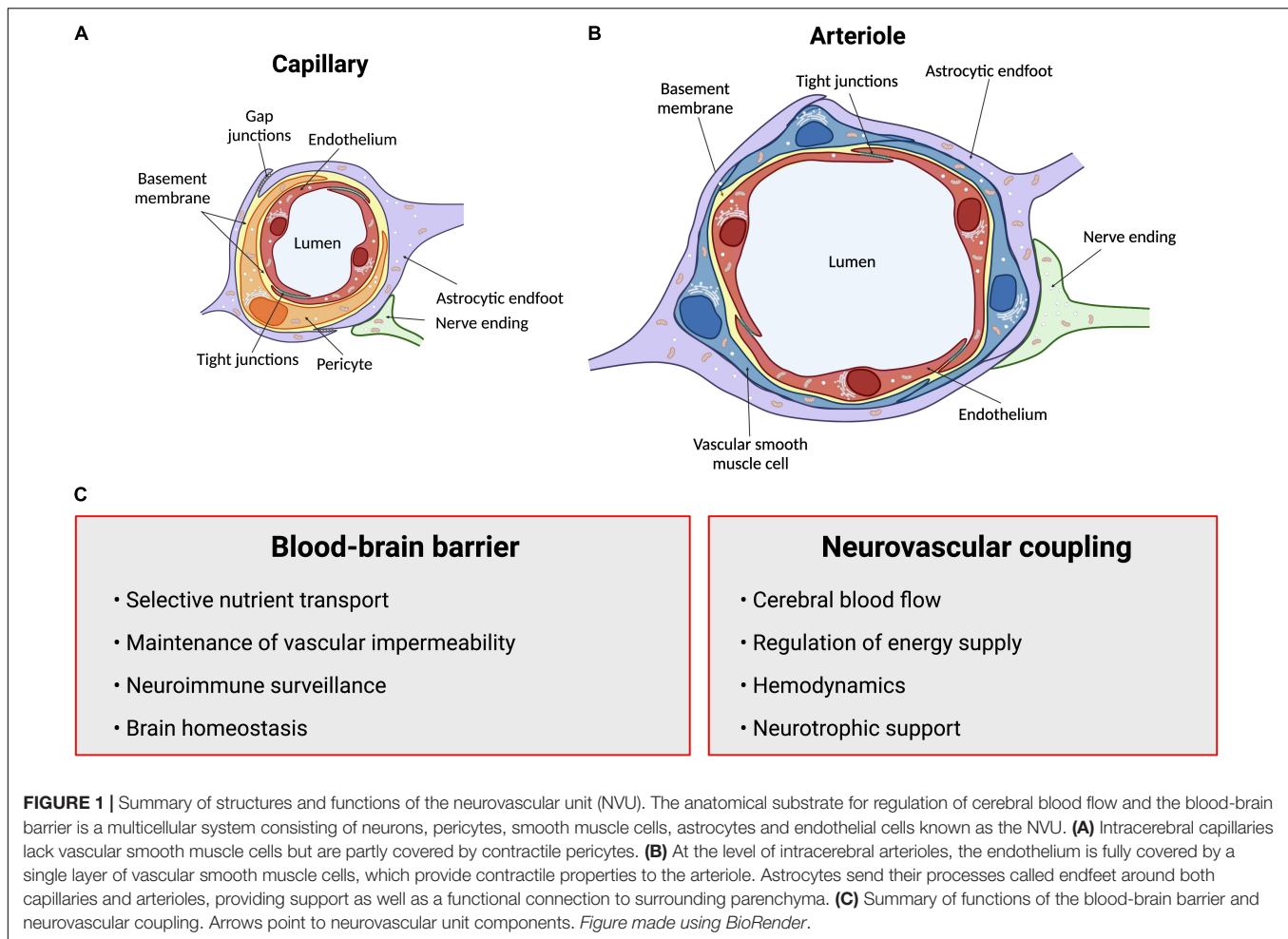
INTRODUCTION

The human brain contains approximately 100 billion vessels (~600 km), all of which are critical for the delivery of nutrients and oxygen to neural cells (Quaeghebeur et al., 2011). Although the brain accounts for only 2% of the body's mass, it consumes about a quarter of the body energy produced at rest (Attwell et al., 2010). This colossal energy consumption is elemental to maintain normal functioning of the brain. Such energy requirements make the brain heavily reliant on key vascular features: (i) a dense vasculature to sustain adequate perfusion, (ii) a functional blood-brain barrier (BBB) to maintain brain homeostasis, and (iii) the proper regulation of cerebral blood flow (CBF) to match metabolic demands (**Figure 1**). Thus, a healthy brain vasculature is essential to support neural cells and ensure normal brain maturation, function and aging (Attwell and Laughlin, 2001; Girouard and Iadecola, 2006; Andreone et al., 2015; Lacoste and Gu, 2015). This is accomplished in part via neurovascular coupling (NVC) mechanisms that regulate CBF to support energetic demands of brain cells (Hamel, 2006; Attwell et al., 2010; Kaplan et al., 2020). While most studies are describing neurovascular signaling at the level of the microvasculature, other vascular segments have received very little attention. There is evidence suggesting that different vascular segments play different roles during vascular responses which is involved in maintaining brain homeostasis. The concept of heterogeneous vascular modules has been extensively reviewed in Schaeffer and Iadecola (2021).

The close anatomical apposition between the nervous and vascular systems supports a functionally integrated network (Attwell et al., 2010; Lecrux and Hamel, 2011; Hillman, 2014; Huneau et al., 2015; Kaplan et al., 2020). This involves modulating vascular tone by secretion of vasoconstrictor and vasodilator molecules. Initially, it was proposed that local metabolic factors released by neurons modulate local CBF (Sherrington, 1890; Friedland and Iadecola, 1991). Since then, several studies have introduced other cellular mediators of NVC which altogether form the neurovascular unit (NVU). This anatomical substrate of NVC indeed involves a multicellular system consisting of neurons, pericytes, smooth muscle cells, astrocytes, microglia and endothelial cells (ECs) that together orchestrate CBF, and thus brain function (Attwell et al., 2010; Andreone et al., 2015; Grubb et al., 2021; **Figure 1**). The cerebral cortex is innervated by projection neurons that release neurotransmitters including, but not limited to, acetylcholine, noradrenaline, serotonin and glutamate, involved in the regulation of vessel diameter (Sandoo et al., 2010). Pericytes, while having debated roles in NVC, possess contractile properties and regulate blood flow around capillaries (Attwell et al., 2010, 2016; Fernandez-Klett and Priller, 2015; Sweeney et al., 2018; Watson et al., 2020; Hartmann et al., 2021). Capillary pericytes are α -smooth muscle actin (SMA)-negative and only partially cover the vessel, while ensheathing pericytes are α -SMA-positive, occupy proximal branches of penetrating arteriole offshoots, and fully cover the vessels. However, they are classified as different from smooth muscle cells as they display an ovoid cell body (Grant et al., 2019). Vascular smooth muscle cells (SMCs),

found on intracerebral arterioles and arteries, are absent from intracerebral capillaries. These cells are short, densely packed, ring-shaped, and essential for regulating vessel tone (Lacoste and Gu, 2015; Frosen and Joutel, 2018; Grant et al., 2019). Astrocytes occupy a critical position between blood vessels and neurons. They can modulate vessel tone via receptor-mediated increase in astrocytic Ca^{2+} , resulting in the release of astrocyte-derived prostaglandins (PGE_2), nitric oxide (NO), epoxyeicosatrienoic acids (EETs), glutamate, or adenosine, all of which can alter vascular diameter and tone (Attwell et al., 2010; Cauli and Hamel, 2010; Filosa and Iddings, 2013; Harada et al., 2015; Haidey et al., 2021), as reviewed in detailed elsewhere (Filosa and Iddings, 2013; Howarth, 2014; MacVicar and Newman, 2015; Mishra, 2017; McConnell et al., 2019; Stackhouse and Mishra, 2021). Whereas microglia are the main regulators of inflammatory processes in the brain, their role in NVC is not well defined. However, recently, they were suggested as essential in regulating CBF during neural activation (Császár et al., 2021). Brain ECs have unique morphological and functional features such as a lack of fenestration, the presence of tight junctions between cells, a low number of pinocytotic vesicles that limit transcytosis, hence forming the first limiting layer of the BBB (Reese and Karnovsky, 1967; Stamatovic et al., 2008; Rizzo and Leaver, 2010; Salmina et al., 2014; Andreone et al., 2015). This highly selective barrier promotes a tightly regulated brain homeostasis to ensure proper neuronal function, protecting the brain from toxins, pathogens, inflammation, and injury (Weiss et al., 2009; Larsen et al., 2014; Daneman and Prat, 2015; Van Dyken and Lacoste, 2018). Furthermore, brain ECs regulate vascular tone by releasing vasodilators including endothelial-derived NO, endothelium-derived EETs, PGE_2 and prostacyclin, as well as vasoconstrictors such as endothelin-1, thromboxane A_2 and prostaglandin $\text{F}_{2\alpha}$ (Mohan et al., 2012; Filosa and Iddings, 2013; Andreone et al., 2015; Kisler et al., 2016, 2017; Dabertrand et al., 2021). While the endothelium regulates vascular permeability and tone, it is also the main target of small vessel disease (SVD), which refers to a pathological process that damages arterioles, venules and brain capillaries. SVD has a major impact on CBF and cognition (Hakim, 2019). The NVU as a whole is also responsible for maintaining BBB integrity (Abbott et al., 2006; Zlokovic, 2008; Daneman, 2012; Kadry et al., 2020). Alterations in vascular patterning, CBF and BBB, either during development or later in life, contribute to the onset and/or progression of early- or late-onset neurological disorders (**Figure 2**).

Well-balanced vascular and neuronal interactions are required to support brain function from early life. The shared spatial and temporal patterns of vascular and neuronal networks suggest an integrative role for vessels in neural development, and *vice versa* (Gu et al., 2005; Carmeliet and Jain, 2011; Andreone et al., 2015; Lacoste and Gu, 2015). Neurovascular crosstalk, which initially takes place during embryogenesis, supports the rising oxygen and nutrient demand of immature neurons as they require extensive energy to maintain normal course of development (De Filippis and Delia, 2011). The increased energy consumption by neurons creates a hypoxic environment acting as a signal for boosting blood vessel production to upsurge delivery of oxygen and metabolic substrates to the brain (Stone et al., 1995;



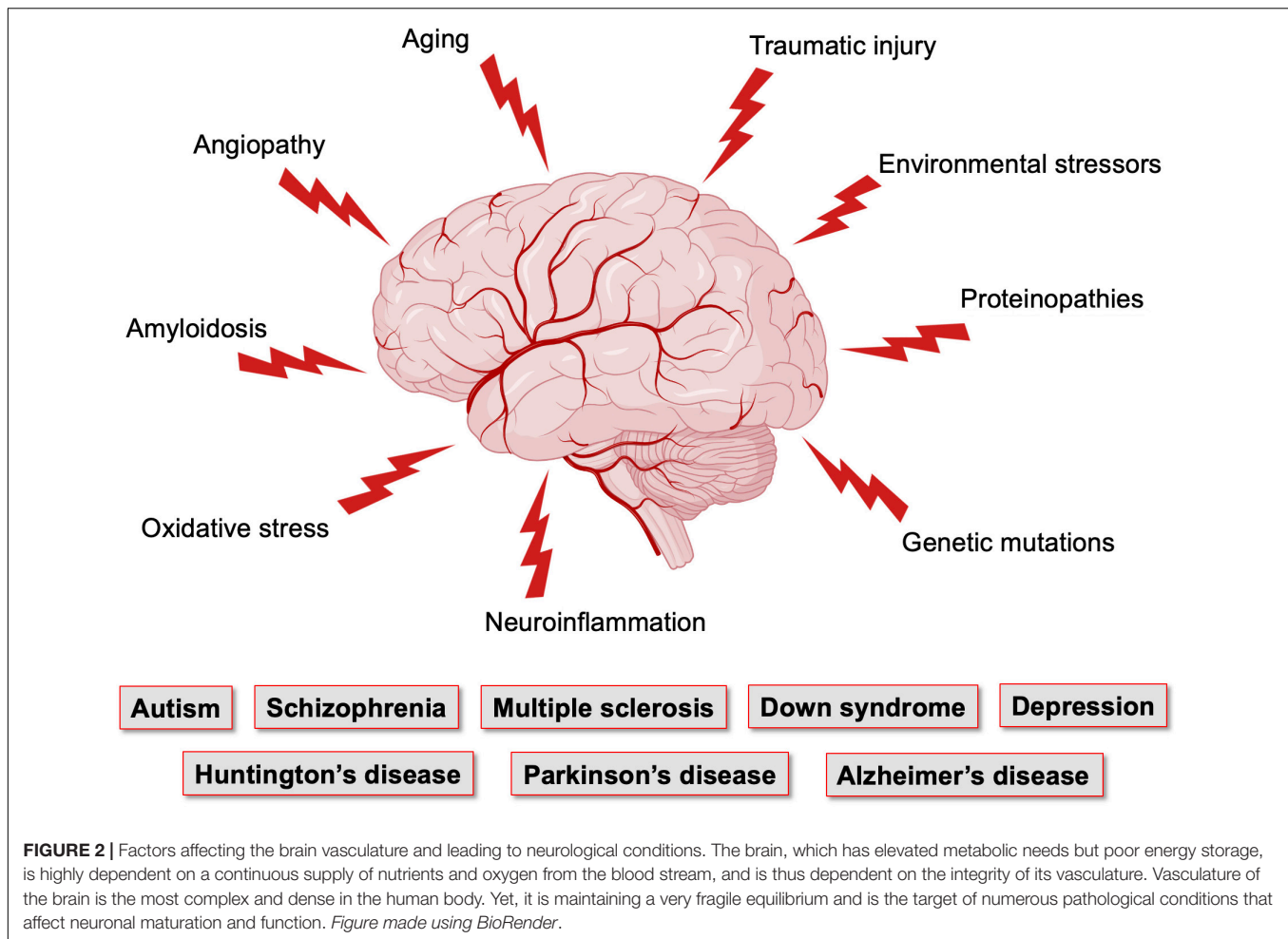
Lacoste and Gu, 2015; Peguera et al., 2021). Hypoxia initiates vessel ingression into deep brain structures, followed by usage of vascular patterning cues (Lacoste and Gu, 2015; Tata et al., 2015; Tata and Ruhrberg, 2018; Okabe et al., 2020). Comparably, ECs instruct neural progenitors into dividing, differentiating or migrating through release of paracrine signals that regulate neuronal development in vascular niches (Hogan et al., 2004; Shen et al., 2004; Daneman et al., 2009; Goldman and Chen, 2011; Delgado et al., 2014; Lacoste and Gu, 2015; Licht and Keshet, 2015; Walchli et al., 2015; Tata and Ruhrberg, 2018; Peguera et al., 2021). Moreover, neuronal activity plays important roles in modulating postnatal brain angiogenesis (Lacoste et al., 2014; Whiteus et al., 2014; Biswas et al., 2020). As the brain matures, vascular networks remodel until the system consists of an extensive network that actively regulates blood flow to adequately sustain energy demands. The functional relationships between neurons and blood vessels ensures that NVC mechanisms progressively develop (Lacoste and Gu, 2015; Coelho-Santos and Shih, 2020). NVC becomes fully functional ~3–4 weeks after birth in rodents, and 7–8 weeks in humans (Yamada et al., 2000; Muramoto et al., 2002; Kozberg et al., 2016).

These vascular features can become defective early in life, affecting brain maturation. Vascular susceptibilities can also

emerge later in life, taking part in neurodegenerative processes. Indeed, NVU deficits play a role in both early- and late-onset neurological disorders (**Figure 2**). Mounting evidence shows that vascular impairments contribute to the pathophysiology of neurological conditions throughout life, including neurodevelopmental, metabolic, and neurodegenerative disorders (Nicolakakis and Hamel, 2011; Van Dyken and Lacoste, 2018; McConnell et al., 2019; Ouellette et al., 2020; Sharma and Brown, 2021). This suggests the existence of a vascular continuum between developmental conditions and illnesses of aging, which will be the focus of this review (**Figure 3**). A better understanding of mechanisms and key players involved in cerebrovascular impairments may lead to transformative therapeutic strategies at different stages of life.

CEREBROVASCULAR DEFICITS ASSOCIATED WITH NEURODEVELOPMENTAL DISORDERS

Neurodevelopmental disorders are considered a group of conditions with onset/diagnosis during infancy, childhood, or adolescence (Morris-Rosendahl and Crocq, 2020). They are



defined by impairments in motor, social, cognitive, academic, and/or occupational functioning. Most studies focused on the neuronal contributions to these disorders; however, concomitant vascular impairments are starting to emerge (Ouellette et al., 2020). Here, we highlight vascular impairments identified in autism spectrum disorders (ASD) and schizophrenia.

Vascular Links to Autism Spectrum Disorders

ASD are pervasive neurodevelopmental disorders associated with social interaction deficits, speech and language impairments, as well as repetitive behaviors and restricted interests (Vijayakumar and Judy, 2016). These disorders have a prevalence of 1–2% in the general population and affect four times more boys than girls (Hogan et al., 2004; Daneman et al., 2009). Individuals with ASD show atypical behaviors associated with visual attention, imitation, social responses, and motor control by 12 months of age. By the age of 3, a child can be efficiently diagnosed with ASD (Park et al., 2016). While the underlying causes of ASD are enigmatic, both environmental and genetic origins have been found, leading to the identification of gene mutations within the ASD population (Hogan et al., 2004;

James et al., 2009; Emerson et al., 2017). Although most studies have been neurocentric, ASD are now being associated with vascular vulnerabilities.

Altered Cerebral Blood Flow in Autism Spectrum Disorders

Neuroimaging techniques can map changes in CBF or blood oxygenation during various activities. Morphological and functional investigations using functional magnetic resonance imaging (fMRI), positron emission tomography (PET), single-photon emission computed tomography (SPECT), or Arterial Spin Labeling (ASL) are used to measure CBF changes in ASD children. CBF disruptions have been demonstrated in ASD patients when compared to healthy controls in different regions of the brain (Björklund et al., 2018). It has also been suggested that perfusion alterations are more pronounced in older children diagnosed with ASD. Cerebral hypoperfusion has been detected in nearly 75% of ASD children (Zilbovicius et al., 2000). As CBF impacts the delivery of oxygen and nutrients to neurons, hypoperfusion in ASD children has been associated with key ASD-related behaviors such as language deficits, impaired executive function and abnormal responses to sensory stimuli, as well as difficulty in facial perception (Siegel et al., 1992;

The vascular continuum in neuronal dysfunction

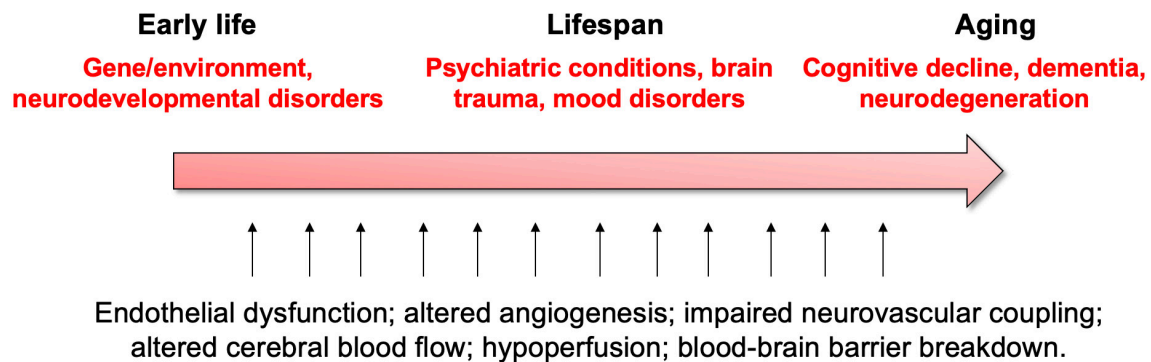


FIGURE 3 | Summary of vascular links to neurological disease throughout life.

Chiron et al., 1995; Ohnishi et al., 2000; Burroni et al., 2008; Reynell and Harris, 2013; Bjørklund et al., 2018; Yerys et al., 2018). These behaviors correlate with abnormal regional cerebral blood flow (rCBF) in the bilateral insula, superior temporal gyri and left prefrontal cortices, medial temporal lobe, supramarginal gyrus, right fusiform gyrus, and dorsal anterior cingulate cortex (Ohnishi et al., 2000; Zilbovicius et al., 2000; Burroni et al., 2008; Jann et al., 2015; Yerys et al., 2018). Studies are attempting to ameliorate these behavioral abnormalities using hyperbaric oxygen treatment (HBOT) to counteract cerebral hypoperfusion in children with ASD. There is some evidence that children who undertook 40 HBOT sessions of 60 min each showed improvements on selected psychosomatic parameters in the Autism Treatment Evaluation Checklist (ATEC) and Childhood Autism Rating Scale (CARS) (Kostiukow and Samborski, 2020). Currently, there is insufficient evidence to support the use of HBOT to treat children with ASD as there are many contradicting studies claiming no improvement in behaviors. Nevertheless, each study followed different protocols, consisted of patients with a large spectrum of behavioral impairments, and some lacked proper control groups, which could explain discrepancies. More research is required to determine if specific groups of children could benefit from HBOT treatment (Rossignol et al., 2012; Sakulchit et al., 2017).

ASL-based measurements of cerebral perfusion showed that children with ASD presented a pattern of widespread hyperperfusion in frontotemporal regions including medial orbitofrontal cortex, bilateral inferior frontal operculum, left inferior/middle temporal gyrus and the right precentral gyrus (Jann et al., 2015). The medial orbitofrontal cortex is known to have extensive connections with the limbic system involved in socio-emotional cognition. Furthermore, hyperperfusion was detected throughout the frontal white matter and subcortical gray matter in ASD children, which correlated positively with severity of social deficits (Peterson et al., 2019). As shown by these studies, CBF abnormalities appear linked to clinical manifestations. Although opposing observations of CBF in ASD patients were reported,

these further support the complexity of these disorders (Jann et al., 2015).

Neurovascular coupling alterations were also observed in ASD patients. Hemodynamic responses in children with ASD during a color-word task were significantly lower than the control group, especially in the dorsolateral prefrontal cortex (Uratani et al., 2019). Conversely, children displayed no difference in hemoglobin concentrations in the prefrontal cortex during a letter fluency task, while adults showed reduced responses (Kawakubo et al., 2009). Despite inter-study variability, there seems to be a consensus on the impact of altered CBF on the expression of behavioral impairments (Zilbovicius et al., 2000; Jann et al., 2015). But as ASD are heterogeneous, with various behavioral traits, genetic causes, medical co-morbidities and medications, these variables may have impacted neuroimaging results, which led to inconsistencies. Importantly, these studies take an important step toward the identification of key players in ASD pathophysiology, opening new opportunities for early diagnosis and treatment.

The relationship between CBF alterations and symptom profiles in ASD children provides insight into disease mechanisms that can be tested in animal models. As most pre-clinical studies have also focused on the neuronal aspects of ASD, very few have considered vascular contributions to these disorders in laboratory models. Recent studies using different ASD mouse models have reported alteration in CBF. A study by Abookasis et al. (2018) using inbred Black and Tan Brachyury (BTBR) *T+tf/J* reported decreased CBF in mutant mice using laser speckle imaging and laser Doppler flowmetry (LDF). Subsequently, work by Ouellette et al. (2020) using the 16p11.2 deletion mouse model of ASD (*16p11.2^{df/+}*; Horev et al., 2011) demonstrated an increase in resting CBF as well as neurovascular uncoupling in adult (P50) *16p11.2^{df/+}* mice compared to WT littermates using a combination of ultrasound imaging and LDF. No difference in CBF or NVC were observed between younger (P14) mutant and control mice (Ouellette et al., 2020). Results from this study in *16p11.2^{df/+}* mice revealed the cause of these functional cerebrovascular impairments: an endothelial

deficit. While normal vascular smooth muscle cell function was measured, defective endothelium-dependent vasodilation was found *ex vivo* following exposure to specific vasomodulators (Ouellette et al., 2020). This suggests that endothelial health plays an important role in the etiology of the 16p11.2 deletion ASD syndrome. Understanding the molecular and cellular factors that mediate CBF alterations in ASD could help design rescue approaches in animal models, as well as therapeutic approaches down the line.

Since MRI studies rely on Blood Oxygen Level Dependent (BOLD) signals as surrogates for neuronal activity (Hillman, 2014; Howarth et al., 2021; Moon et al., 2021), it is possible that changes in rCBF reflect changes in underlying neuronal activity. For instance, cerebral cortex hypoperfusion in ASD patients could reflect lower metabolic demands (Schifter et al., 1994). In the case *16p11.2^{df/+}* mice, however, it is interesting to note that a neurovascular uncoupling was measured, with enhanced neuronal activation yet reduced vascular responses to whisker stimulations, which led to the discovery of endothelium-dependent deficits (Ouellette et al., 2020).

Altered Blood-Brain Barrier and Angiogenesis in Autism Spectrum Disorders

Cerebral vessels are central for the maintenance of brain homeostasis, sustaining proper neuronal function, and providing an effective protection against toxins and pathogens (Profaci et al., 2020). The BBB consists of specialized ECs lining the vessel wall to separate the peripheral blood from cerebral tissue. Brain (central) ECs are distinct from peripheral ECs, as they produce specific proteins to control the flux (entry and exit) of metabolites across vessels, to maintain low rates of trans-endothelial vesicular transport, and to form tight junctions to limit the para-cellular flow of material between adjacent ECs (Andreone et al., 2015; Chow and Gu, 2015; Kealy et al., 2020). Alterations in the BBB are at the core of the onset and/or progression of numerous neurological disorders (Daneman and Prat, 2015; Van Dyken and Lacoste, 2018; Profaci et al., 2020). Only few studies have investigated the components of the BBB in the context of ASD. Children diagnosed with ASD have been associated with reduced levels of adhesion molecules such as soluble Platelet Endothelial Cell Adhesion Molecule-1 (PECAM-1, or CD31) and P-selectin. Since these molecules are essential to modulate BBB permeability through signaling and leukocyte infiltration, it suggests that crucial BBB components may be at play in ASD pathophysiology (Onore et al., 2012). Furthermore, a post-mortem study, with a small sample size, demonstrated altered BBB integrity in ASD with increased gene expression of matrix metalloproteinase (MMP)-9 (Fiorentino et al., 2016). Studies have shown that MMP-9 regulates cell proliferation, adhesion, degradation of laminin and collagen, angiogenesis, oxidative injury, and is implicated in BBB breakdown (Lepeta and Kaczmarek, 2015; Turner and Sharp, 2016). Additionally, important components of BBB integrity displayed altered expression in ASD patients, including claudin-5 (*CLDN5*) and claudin-12 (*CLDN12*), as well as tricellulin (*MARVD2*), a component of tight junctions involved in decreased permeability to macromolecules in brain ECs (Fiorentino et al.,

2016). In an older study, a small subset of ASD participants demonstrated higher levels of autoantibodies against brain ECs in the serum compared to typically developing individuals, suggesting an impact on the BBB (Connolly et al., 1999). Animal models have facilitated the study of BBB integrity in ASD. In a valproic acid rat model of autism, increased BBB permeability to Evans blue was found in the cerebellum, a phenotype attenuated by treatment with memantine, an NMDA receptor modulator. This BBB alteration was also attenuated using minocycline (antibiotic) and agomelatine (melatonin receptor) treatment (Kumar et al., 2015; Kumar and Sharma, 2016). Animal studies have investigated transendothelial transport mechanisms in ASD mouse models. Tarlungeanu et al. (2016) demonstrated that the large neutral amino acid transporter (LAT1, *Slc7a5*) localized at the BBB to maintain normal levels of brain branched chain amino acid (BCAA) was required for neurotypical development. Mice harboring an endothelial-specific deletion of *Slc7a5* (*Slc7a5^{ΔEC}*) displayed behaviors reminiscent of ASD, including motor dysfunctions consistent with a study in human patients harboring the constitutive mutation (Novarino et al., 2012; Tarlungeanu et al., 2016). Interestingly, administration of BCAA rescued ASD-like behaviors in *Slc7a5^{ΔEC}* mice (Tarlungeanu et al., 2016).

Recently, a post-mortem analysis of brain tissue from individuals diagnosed with ASD revealed significantly higher levels of markers associated with pericytes, as well as increased vascular tortuosity, indirectly suggesting impairments in angiogenesis, a process through which new blood vessels are formed (Azmitia et al., 2016). A more recent study in *16p11.2^{df/+}* mice revealed impaired cerebral angiogenesis in young (P14) *16p11.2^{df/+}* male mice compared to sex-/age-matched littermates, a phenotype which was absent in adult mice. Defective angiogenic activity was also measured using primary brain ECs from P14 *16p11.2^{df/+}* males or ECs derived from human-induced pluripotent stem cells (hiPSCs) of 16p11.2 deletion carriers (Ouellette et al., 2020). Moreover, RNA-sequencing analysis of *16p11.2^{df/+}* mouse brain EC transcriptome revealed changes in the expression of genes involved in angiogenesis (e.g., *Grem1*, *Apln*, *Angpt2*), while key genes involved in BBB regulation (e.g., *Pecam1*, *Mfsd2a*, *Cldn5*, *Slc2a1*) were not affected by the 16p11.2 deletion (Ouellette et al., 2020). Finally, this study generated a mouse model with endothelial-specific 16p11.2 haploinsufficiency which recapitulated ASD-related phenotypes, revealing a causal relationship between endothelial dysfunction and neuronal aspects of the 16p11.2 deletion syndrome (Ouellette et al., 2020).

Overall, these studies allude to the contribution (structural and functional) of a defective BBB and NVU in ASD, with an important role for endothelial impairments.

Vascular Links to Schizophrenia

Schizophrenia is a debilitating neurodevelopmental disorder affecting ~1% of the population. It is associated with behavioral and cognitive symptoms that arise progressively. Memory and attention deficits appear in childhood, while positive symptoms (psychotic episodes) and negative symptoms (social and motivational deficits) emerge later in adolescence or early adulthood (Stachowiak et al., 2013). Although the incidence of

schizophrenia is higher in men, women have a slightly later disease onset (Gogtay et al., 2011; Ochoa et al., 2012). While the behavioral aspects of schizophrenia have been described, the causes of this disorder are poorly known. As in ASD, both genetic and environmental origins are involved. Schizophrenia has been associated with genes essential for a wide range of functions including neuronal connectivity and patterning of brain structures, cell proliferation and differentiation, as well as cytoskeleton reorganization (Stachowiak et al., 2013; Clifton et al., 2019). As in most neurological disorders, the implication of neuronal alterations has been extensively studied, but research on vascular impairments in schizophrenia is starting to emerge.

Altered Cerebral Blood Flow in Schizophrenia

Cognitive impairments are often present before the first psychotic episode in patients with schizophrenia (Keefe and Harvey, 2012; Schuepbach et al., 2016) and deficits in executive functions are often parallel to changes in CBF. Several studies have linked altered CBF with schizophrenia-related symptoms (Sabri et al., 1997; Malaspina et al., 1999, 2004; Pinkham et al., 2011; Fujiki et al., 2013; Schuepbach et al., 2016; Stegmayer et al., 2017; Zhu et al., 2017). Interestingly, the manifestations of negative or positive symptoms correlate with different rCBF changes. In a study by Pinkham et al. (2011), CBF of 30 schizophrenia patients was measured using ASL perfusion MRI, which revealed a positive correlation between increased severity of positive symptoms and higher CBF in the cingulate and superior frontal gyri, but decreased CBF in precentral and middle frontal gyri. Patients who presented with severe negative symptoms also displayed reduced CBF in the superior temporal gyrus bilaterally, cingulate and left middle frontal gyri (Malaspina et al., 2004; Scheef et al., 2010; Pinkham et al., 2011; Liu et al., 2012). Most studies investigating CBF alterations in schizophrenia considered perfusion rates from medicated patients, and a small number of studies have measured CBF rates in neuroleptic-naïve patients. Using ASL in non-medicated patients, the schizophrenia group displayed resting-state hypoperfusion in the frontal lobes, anterior and medial cingulate gyri, as well as in the parietal lobes, while increased perfusion was measured in the cerebellum, brainstem and thalamus (Scheef et al., 2010). Sabri et al. (1997) measured rCBF using SPECT in non-medicated patients that have experienced positive symptoms, revealing that rCBF values varied depending on the severity of positive symptoms. Hyperperfusion was detected in the frontal, anterior cingulate as well as in both parietal and temporal cortices in patients who had scored high in severity for formal thought disorder (disturbance of the organization and expression of thought). In contrast, patients who scored high for delusions, hallucinations or distrust, with low scores for formal thought, displayed hypoperfusion in the same brain regions. No difference in rCBF was identified between control and schizophrenia groups after treatment (Andreasen et al., 1997; Sabri et al., 1997; Horn et al., 2009). Recent studies have detected hyperperfusion and hypoperfusion in brain regions from individuals with hallucinations. For instance, CBF increase was found in the right superior temporal gyrus and caudate nucleus, while CBF decrease was found

bilaterally in the occipital and left parietal cortices (Zhuo et al., 2017). In another study, patients were classified based on the severity of three behavioral dimensions (language, affectivity, and motor) according to the Bern Psychopathology scale. Patients with altered affectivity were associated with increased CBF in the amygdala, while changes in language dimension were linked to increased CBF in Heschl's gyrus (Stegmayer et al., 2017). While schizophrenia is classified as a neurodevelopmental disorder, its symptoms persist with age. Studies have identified significant bilateral temporal hypoperfusion related to aging and disease course. It has been suggested that this decrease in CBF with aging is paralleled with the degenerative changes observed in patients with schizophrenia (Schultz et al., 2002; Kawakami et al., 2014).

The polygenic risk of schizophrenia is an important dimension of this syndrome, and changes in CBF have been identified in patients diagnosed for either familial or sporadic schizophrenia. Sporadic schizophrenia patients were associated with hypofrontality (left frontal gyrus, orbitofrontal cortex, anterior cingulate, and paracingulate cortices), while familial schizophrenia patients had left temporoparietal hypoperfusion (posterior Sylvian fissure at the superior and inferior parietal lobules, angular, and supramarginal gyri). In both groups, positive symptoms are often associated with increased rCBF in the parahippocampal gyrus, cerebellum, and pons (Malaspina et al., 2004). Sporadic patients showed additional hyperperfusion in the fusiform gyrus, and familial patients the hippocampus, dentate, amygdala, thalamus, and putamen (Malaspina et al., 2004). In addition, the prefrontal cortex in schizophrenia has been associated with deficits of pericapillary oligodendrocytes, which could contribute to changes in CBF (Vostrikov et al., 2008; Uranova et al., 2010). Altogether, these studies support the idea that altered CBF is involved in schizophrenia pathophysiology.

In addition to studies investigating resting state CBF, there is evidence of altered NVC in schizophrenia whereby many reports demonstrate reduced hemodynamic response, reflecting reduced neuronal activity during processing of cognitive tasks, especially in the lateral prefrontal cortex and temporal regions (Ford et al., 1999, 2005; Mathalon et al., 2000; Carter et al., 2001; Hanlon et al., 2016; Pu et al., 2016). As with CBF reports, there are inconsistent hemodynamic responses associated with schizophrenia since increased hemodynamic responses in hippocampus, thalamus and prefrontal cortex have been identified (Tregellas et al., 2007). These conflicting results are translating to rodent models of schizophrenia whereby some models have revealed overall hypofrontality, hypoperfusion in the hippocampus or hyperperfusion in the somatosensory cortex (Finnerty et al., 2013; Song et al., 2013; Drazanova et al., 2019).

Altogether, these studies support the idea that altered CBF regulation is involved in schizophrenia pathophysiology. Moreover, it appears critical to consider the polygenic risk of disease, the category and severity of symptoms, as well as the age of patients when comparing CBF rates in schizophrenia. Although many studies have detected altered CBF using various methods, results thus far remain conflicting based on various stages of disease and pharmacological treatment (Drazanova et al., 2019).

Altered Blood-Brain Barrier and Angiogenesis in Schizophrenia

A dysfunctional BBB has been reported in schizophrenia, with increased permeability to damaging proteins (Müller and Ackenheil, 1995; Shcherbakova et al., 1999; Crockett et al., 2021). Studies are starting to decipher changes in cells associated with the BBB (for a detailed review, see Carrier et al., 2020). Briefly, evidence of schizophrenia-associated microvascular abnormalities in the neocortex include thickening and deformation of basal lamina, vacuolation of cytoplasm in ECs, basal lamina and astrocytic end-feet, swelling of astrocyte end-feet, activation of microglial cells in the prefrontal and visual cortex, as well as atypical vascular arborization (Uranova et al., 2010; Carrier et al., 2020).

Moreover, specific mutations are associated with schizophrenia, including alterations in the 22q11.2 deletion syndrome (22qDS) -strongest monogenic risk allele for this disorder, and polymorphisms in claudin-5, a densely expressed tight junction molecule (Gur et al., 2017; Greene et al., 2018; Carrier et al., 2020) altogether revealing barrier dysfunction in schizophrenia patients (Greene et al., 2018; Crockett et al., 2021). Post-mortem brain sections from 22qDS patients and animal models of 22qDS both demonstrate reduced claudin-5 expression in the BBB, which in turn compromised BBB function (Nishiura et al., 2017; Guo et al., 2020; Crockett et al., 2021; Usta et al., 2021). Additionally, altered levels of vascular endothelial (VE)-cadherin and occludin in ECs were identified in schizophrenia. These molecules regulate adherence of ECs and restrict movement of substances across the BBB (Cai et al., 2020). Furthermore, BBB hyperpermeability has been associated with another risk allele for schizophrenia. *NDST3*, expressed in the brain, encodes an enzyme involved in the metabolism of heparan sulfate, a component of basal lamina extracellular matrix that is required for BBB integrity (Khandaker et al., 2015).

Studies have documented primary vascular endothelial dysfunction in schizophrenia. Individuals carrying *MTHFR T* and/or *COMT Val* risk allele have been associated with cerebrovascular endotheliopathy, as well as lower frontal executive functions (Grove et al., 2015). While endothelial dysfunction is possibly associated with schizophrenia, many studies are using peripheral endotheliopathy as a surrogate marker for endothelial dysfunction. For example, studies are using non-invasive peripheral arterial tonometry (RH-PAT) to assess peripheral arteriole endothelial-dependent vasodilation and revealed impaired peripheral arterial vasodilation in schizophrenia (Ellingrod et al., 2011; Burghardt et al., 2014). Notably, brain ECs have unique properties to maintain BBB integrity and brain homeostasis. Although altered endothelial function was found in the periphery, it does not represent a definite marker of brain (central) endothelial dysfunction. A critical regulator of angiogenesis, *vascular endothelial growth factor* (VEGF), and its receptor (VEGFR2) have been found upregulated in the prefrontal cortex of individuals diagnosed with schizophrenia (Hino et al., 2016). Findings of elevated VEGF could also be linked to vascular hyperpermeability, as VEGF not only regulates angiogenesis but increases BBB leakage (Mayhan, 1999; Zhang et al., 2000). Conversely, a different group revealed

that a decreased production of VEGF predisposed individuals to develop this disorder and contributed to the severity of symptoms (Saoud et al., 2021). Another study investigated the impact of hiPSC-derived neural stem cells from schizophrenia patients on angiogenesis (Casas et al., 2018). This study found an imbalance in the expression and secretion of several angiogenic factors and non-canonical neuro-angiogenic guidance cues from neural stem cells from schizophrenic patients. Conditioned media from these cells induced impaired angiogenesis as evidenced by reduced number of sprouts and tubes formed in *in vivo* and *in vitro* models, as well as decreased neural stem cell migration compared to control conditioned media (Casas et al., 2018).

CEREBROVASCULAR DEFICITS ASSOCIATED WITH NEURODEGENERATIVE DISORDERS

CNS disorders are dichotomized as early onset neurodevelopmental disorders and late-onset neurodegenerative diseases (Taoufik et al., 2018). Neurodegenerative diseases consist of a group of heterogeneous disorders characterized by the progressive degeneration of structure and function in the CNS (Gitler et al., 2017). Although neurodegenerative and neurodevelopmental disorders are differentially classified, an accumulating body of work demonstrates significant similarities between these two groups of conditions. Here below, we cover cerebrovascular impairments reported in four neurodegenerative diseases that emerge throughout lifespan: multiple sclerosis (MS), Huntington's disease (HD), Parkinson's disease (PD), and Alzheimer's disease (AD).

Vascular Links to Multiple Sclerosis

MS is a chronic autoimmune disease of the CNS, occurring when the immune system attacks its own nerve fibers and myelin sheaths (D'Haeseleer et al., 2013). The pathological hallmark of MS consists of perivenular inflammatory lesions, leading to demyelinating plaques and diffuse axonal degeneration throughout the CNS (Dobson and Giovannoni, 2019). It is characterized by the infiltration of T cells reactive against myelin in the CNS (Schwartz and Kipnis, 2005). This demyelinating disease has key features including inflammation, BBB disruption and neurodegeneration. MS has a prevalence of 0.5–1.5 per 100,000 individuals, whereby women are three times more affected than men (Harbo et al., 2013). The age of MS onset is situated between 20 and 40 years of age (Ortiz et al., 2014). General symptoms related to MS include, but are not limited to, tremors, lack of coordination as well as weakness in limbs. There are various types of MS including relapsing-remitting MS (RR-MS), secondary progressive MS (SP-MS) and primary progressive MS (PP-MS). RR-MS consists of unpredictable relapses or inflammatory flare-ups during which new symptoms appear or existing ones worsen (Adhya et al., 2006). Most people with RR-MS, transition to a disease phase known as SP-MS. In this phase, there is progressive worsening and fewer relapses. Active lesions with profound lymphocytic inflammation are mostly found in RR-MS (Dobson and Giovannoni, 2019). PP-MS is

considered as a slow accumulation of disability without defined relapses. In this case, PP-MS is associated with an inactive lesion core surrounded by activated microglia and macrophages (Dobson and Giovannoni, 2019).

Altered Cerebral Blood Flow in Multiple Sclerosis

MS has been associated with functional cerebrovascular abnormalities including decreased cerebral perfusion and reduced CNS venous blood drainage, known as chronic cerebrospinal venous insufficiency (D'Haeseleer et al., 2011). SPECT, PET, and ASL imaging studies have reported decreased CBF in both gray and white matter of MS patients (Ge et al., 2005; D'Haeseleer et al., 2011). Widespread cerebral hypoperfusion has been revealed in SP-MS, RR-MS and PP-MS patients, while an ischemic threshold was not reached (Adhya et al., 2006; Ota et al., 2013; Monti et al., 2018). Gray matter hypoperfusion in MS suggests a reduction of metabolism due to the loss of cortical neurons (Peruzzo et al., 2013). Furthermore, studies have reported that CBF is globally impaired in normal appearing white matter (NAWM) of patients with early RR-MS (Law et al., 2004; Adhya et al., 2006). Of note, CBF was generally lower in PP-MS than in RR-MS in the periventricular and frontal white matter (Adhya et al., 2006). In the contrary, other studies have measured elevation of CBF and cerebral blood volume (CBV) in NAWM of patients with early RR-MS several weeks before signs of increased BBB permeability (Wuerfel et al., 2004). Although studies on different types of MS revealed changes in CBF, general active demyelinating lesion regions are associated with hyperperfusion while the more stable forms show hypoperfusion (Monti et al., 2018). Decreased CBF in cerebral NAWM, thalamus, and putamen was identified in patients whose symptoms emerged within the first 5 years of onset. This suggests that CBF alterations are present in the very early stages of the disease (Varga et al., 2009). Different mechanisms have been proposed to explain hypoperfusion in MS. A study suggested that decreased CBF is secondary to axonal degeneration, which leads to a decreased metabolic demand (Saindane et al., 2007). However, this hypothesis is yet to receive experimental support. A second mechanism that has been proposed is an impaired energy metabolism of astrocytes (De Keyser et al., 1999). In MS, astrocytes are deficient in β 2-adrenergic receptors which regulate high energy-consuming activities, such as glycogenolysis and phosphocreatine metabolism (De Keyser et al., 1999). Reduced energy production in astrocytes could be contributing to altered CBF. A third mechanism suggested was increased release of vasoconstrictor endothelin-1 (ET-1) from reactive astrocytes, found in a post-mortem study on white matter samples of RR-MS patients (D'Haeseleer et al., 2013; Hostenbach et al., 2019). Hence, elevated levels of ET-1 could be involved in dysregulating CBF in MS. Interestingly, administration of ET-1 antagonist Bosentan restored CBF to control levels in MS patients (D'Haeseleer et al., 2013).

Impaired cerebral vascular reactivity was evidenced in MS patients exposed to hypercapnia, which has been suggested to contribute to neuronal death identified in this disorder (Marshall et al., 2014). This global deficit is thought to be associated with elevated levels of NO reported in MS (Trapp

and Stys, 2009; Juurlink, 2013). These studies suggest that the overproduction of NO may desensitize endothelial and smooth muscle cell function, causing decreased vasodilatory capacity and limited blood supply for neurons that perform demanding tasks. Increased NO in MS may lead to neuronal activity-induced hypoxia leading to neurodegeneration (Marshall et al., 2014). Interestingly, high inflammatory MS lesion load has been associated with increased CBF. Therefore, perfusion changes may be sensitive to active inflammation (Bester et al., 2015). However, it remains unclear whether abnormal perfusion in MS is a precursor of lesions or occurs independently of lesion development (Marshall et al., 2014).

Notably, MS has been associated with cerebral SVD. It was demonstrated that younger MS cases are more severely impacted by cerebral SVD compared to older individuals (Geraldes et al., 2020). This suggests that the interaction between MS and cerebral SVD is affected by age, an assumption still under investigation (Geraldes et al., 2020).

Altered Blood-Brain Barrier and Angiogenesis in Multiple Sclerosis

BBB dysfunction is considered a major hallmark of MS and is deemed a trigger of disease onset (McQuaid et al., 2009; Cramer et al., 2014). Intense focal disruption of the BBB associated with inflammation (identified by gadolinium-enhanced MRI at acute and chronic MS lesion sites; Saade et al., 2018) and diffuse extensive BBB disruption with a long-term pathological activity, are both found in MS patients (Bennett et al., 2010). Hyperpermeability of the BBB was evidenced by leukocyte passage across the BBB (Cramer et al., 2014). Increased BBB leakage was associated with decreased expression of tight junction proteins in brain capillary ECs in patients with active lesions, inactive lesions, as well as NAWM associated with fibrinogen leakage (Kirk et al., 2003; McQuaid et al., 2009; Bennett et al., 2010). More specifically, dysregulation of tight junction adaptor protein ZO-1, occludin and claudin-5 have been reported in both primary progressive and secondary progressive disease states (Kirk et al., 2003; Leech et al., 2007). Experimental autoimmune encephalomyelitis (EAE) in rodents is a disease model with clinical and pathological characteristics relevant to the study of MS. This model revealed reorganization of ZO-1 and actin in the presence of inflammatory factors *in vitro*, associated with increased permeability of an endothelial monolayer (Bennett et al., 2010). The EAE model also revealed increased expression of VEGF in ECs, astrocytes, monocytes and activated TH1 lymphocytes, all of which contribute to BBB permeability during the early phase of disease, while decreased expression VEGF was evident in the late phase (Girolamo et al., 2014). The increase in VEGF expression was also found in the brain of MS patients (Girolamo et al., 2014). Furthermore, junctional adhesion molecule-A, a component of tight junctions, was found abnormally distributed in active and inactive MS lesions, although adherent junction proteins were normally expressed and localized in MS tissue (Padden et al., 2007). In addition, levels of PECAM-1 were found increased in active gadolinium-enhancing MS lesions (Ortiz et al., 2014). While BBB leakage is evident in MS, the complex network of cellular

and molecular players that lead to this dysfunction have yet to be fully understood. Targeting BBB defects in MS represent a therapeutic opportunity, for instance with MMP inhibitors, interferons, and corticosteroids (Minn et al., 2002; Ross et al., 2004; Pardridge, 2012; Ortiz et al., 2014). However, no current therapy addresses BBB deficits (Ortiz et al., 2014). For more details on BBB dysfunction in MS, the following reviews can be consulted (Girolamo et al., 2014; Kamphuis et al., 2015; Xiao et al., 2020).

ECs proliferation as well as an increase in vascular network density has been reported (Ludwin, 2006; Holley et al., 2010). Increased angiogenesis was suggested to contribute to disease progression as well as remission after relapses (Papadaki et al., 2014). In addition to increased VEGF levels, VEGFR2 is also expressed on ECs in active MS lesions (Seabrook et al., 2010). Other molecules, such as basic fibroblast growth factor, were increased in MS patients and involved in angiogenesis (Su et al., 2006). MS patients with activated lesions and NAWM show blood vessels with a glomeruloid morphology, hemorrhages and vessel wall hyalinization (Girolamo et al., 2014). Immunosuppressive therapies have been used in aggressive MS as they not only impact neuroinflammation but also have an anti-angiogenic effect. Further research is warranted to elucidate the vascular links to MS and identify new therapeutic targets, as disease modifying drugs have unfortunately little to no impact on MS progression (Girolamo et al., 2014).

Vascular Links to Huntington's Disease

HD is an hereditary, autosomal dominant and neurodegenerative disorder (Davenport, 1915; Wasmuth et al., 1988; Bano et al., 2011) leading to altered muscle coordination and declined mental abilities (Paulsen, 2011; Ha and Fung, 2012). An expansion of trinucleotide CAG repeats on chromosome 4 within the Huntingtin gene (*HTT*) results in the production of an altered Huntingtin (Htt) protein which accumulates in specific brain regions. Aggregation of mutant Htt (mHtt) leads to increased neurotoxicity (Zheng and Diamond, 2012), particularly in subcortical brain structures such as the neostriatum (caudate and putamen) where GABAergic medium-spiny neurons are particularly vulnerable (Sieradzan and Mann, 2001; Walker, 2007; Ross and Tabrizi, 2011; Drouin-Ouellet et al., 2015; McColgan and Tabrizi, 2018). At the cellular level, mHtt results in neuronal dysfunction and death through disrupted mechanisms involved in proteostasis, transcription and mitochondrial function as well as toxicity from the mutant protein (McColgan and Tabrizi, 2018). Worldwide, 2.71 per 100,000 individuals suffer from HD (Rawlins et al., 2016; Kounidas et al., 2021). Both men and women are affected equally, and heterogeneous symptoms emerge at around 40 years of age. However, functional and structural brain alterations emerge a decade before symptoms manifest (Snowden, 2017). Carriers of CAG repeat expansions in *HTT* can be identified decades before clinical manifestation, allowing researchers to identify possible biomarkers in the premanifest stage of HD (preHD). With this comes the increasing interest to study cerebrovascular abnormalities in HD (Snowden, 2017).

Altered Cerebral Blood Flow in Huntington's Disease

HD-related perfusion deficits have been mostly associated with cerebral hypoperfusion (Reid et al., 1988; Sotrel et al., 1991; Hasselbalch et al., 1992; Harris et al., 1999; Deckel and Duffy, 2000; Wild and Fox, 2009). There is evidence of reduced CBF in the basal ganglia in early HD, prior to gross structural changes and to motor symptoms. In these cases, severity of cortical hypoperfusion correlated with decreased functional capabilities (Sax et al., 1996; Harris et al., 1999). In preHD patients, classified as either near or far from motor symptom onset, displayed altered rCBF by MR-based perfusion imaging. Participants with preHDfar and preHDnear had lower rCBF in the medial prefrontal cortex and increased rCBF in the left precuneus. Of note, structure and function of the precuneus and hippocampus can be abnormal in very early HD (Feigin et al., 2006). PreHDnear participants had additional regions showing altered rCBF, including hypoperfusion in the medial and lateral prefrontal cortex and hyperperfusion in the right hippocampus (Wolf et al., 2011).

While resting CBF is affected, early manifest and premanifest HD patients also display altered neurovascular coupling during visual stimulation (Klinkmueller et al., 2021). After HD onset, a significant hypoperfusion in the HD group was identified in most of the cerebral cortex. During problem-solving activities, such as solving a maze or resting their eyes open while looking at a modified maze, patients with HD showed increased CBF in the caudate nucleus (Deckel and Duffy, 2000; Deckel et al., 2000). Following physical activity, HD patients were associated with CBF hyperperfusion compared to the control group (Stevenson et al., 2020).

Animal models of HD (e.g., gene knock-in of a human exon 1 CAG₁₄₀ expansion repeat) also revealed altered rCBF. In mice as in humans, different brain regions displayed either hypoperfusion (basal ganglia motor circuit, hippocampus and prefrontal area) or hyperperfusion (cerebellar-thalamic and somatosensory regions). This altered CBF was apparent at a presymptomatic stage (Wang et al., 2016).

While CBF is starting to emerge as a biomarker for HD, mounting evidence supports the utilization of CBV as an additional metric. Several studies have reported elevated CBV in preHD patients (Hua et al., 2014; Liu et al., 2020). In addition, there is evidence of increased CBV in cortical gray matter after HD onset (Drouin-Ouellet et al., 2015), suggesting that arteriolar CBV may be a sensitive biomarker for premanifest HD (Hua et al., 2014; Liu et al., 2020). From these studies it was suggest that imaging of CBF may be used to detect widespread functional abnormalities in HD, and possibly predict HD symptoms onset during premanifest stages.

Altered Blood-Brain Barrier and Angiogenesis in Huntington's Disease

Increases in vessel density, BBB permeability and VEGF-A release were observed in HD patients and animal models of HD (Stevenson et al., 2020). There is evidence that BBB leakage increases alongside disease progression (Drouin-Ouellet et al., 2015). Despite these observations, there seems to be discrepancies between mouse models of HD. For instance, the BACHD

transgenic mice, a well-known model of HD expressing the full-length mutant human *HTT*, failed to develop BBB breakdown at 12 months of age despite robust motor deficits (Lin et al., 2013; Mantovani et al., 2016). BBB dysfunction in HD patients has been associated with decreased tight junction molecules such as occludin and claudin-5 (Drouin-Ouellet et al., 2015). Moreover, other markers associated with BBB permeability, including hepatocyte growth factor, interleukin-8 and tissue inhibitor of MMP-1, were found elevated in HD patients (Drouin-Ouellet et al., 2015). A transgenic mouse model of HD (R6/2 mice) confirmed elevated tight junction molecules similar to HD patients. The R6/2 mouse model of HD is the most commonly studied and harbors a mutant *Htt* with CAG repeat expansion in exon 1 (Li et al., 2005). R6/2 mice also displayed increased transcytosis and paracellular transport across the brain endothelium compared to control mice (Drouin-Ouellet et al., 2015). In R6/2 mice, tight junction imbalance and perturbed BBB homeostasis were perceptible at very early stage of the disease, in absence of symptoms (Di Pardo et al., 2017). At the structural level, mHtt aggregates were found in the basal membrane of cerebral blood vessels in HD patients (Drouin-Ouellet et al., 2015). Interestingly, mHtt aggregates were localized in ECs, smooth muscle cells and perivascular macrophages, consistent with observations in R6/2 mice.

Further research is needed to determine BBB impairments in preHD patients. Lim et al. (2017) reported that iPSCs-derived brain microvascular endothelial cells (BMECs) from HD patients exhibit increased angiogenesis and altered barrier properties associated with elevated transcytosis and paracellular permeability. An increased and unregulated angiogenic activity may lead blood vessels to become more permeable with a potential role in neurovascular dysfunction in HD. RNA-seq analysis revealed a significant number of affected gene that regulate both clathrin- and caveolin- mediated endocytosis, which could lead to changes in endo- and transcytosis across the brain endothelium. These genes include *FABP4*, *DYNAMIN*, and *FILAMIN* that play a role in vesicle formation and scission. In addition, higher levels of transcytosis-related genes such as *CAV1* was detected in HD iPSCs-derived BMECs that also displayed impaired Wnt/ β -catenin signaling (Lim et al., 2017). The Wnt/ β -catenin pathway is essential for regulation of cell proliferation, cell determination and tissue homeostasis (Silva-Garcia et al., 2019). Furthermore, astrocytes from both HD patients and mouse models were associated with higher levels of VEGF-A, which may trigger proliferation of ECs and contributes to neurovascular changes in HD (Hsiao et al., 2015). Of note, sustained delivery of VEGF into the rat striatum via injectable hydrogels was neuroprotective in a lesioned model of HD; VEGF implants significantly protected against the quinolinic acid-induced loss of striatal neurons (Emerich et al., 2010). Moreover, neuroprotection induced by inhibition of hypoxia inducible factor (HIF) prolyl-4-hydroxylases in HD mice has been correlated with enhanced VEGF expression (Niatetskaya et al., 2010). In post-mortem tissue, cerebral blood vessel density was greater in HD patients while no differences in diameter of small- or medium sized blood vessels have been observed (Drouin-Ouellet et al., 2015). Post-mortem tissue of HD patients

revealed a higher proportion of small compared to medium-sized blood vessels in the putamen, an effect occurring in parallel with putamen degeneration. Notably, altered density of small blood vessels in HD patients was consistent with the R6/2 mouse model when brain vascular anomalies were restricted to smaller vessels (Drouin-Ouellet et al., 2015; St-Amour et al., 2015).

Vascular Links to Parkinson's Disease

PD is the second most common neurodegenerative disorder after AD (Antony et al., 2013). It is characterized by the progressive degeneration of the nigrostriatal system, resulting in rigidity, bradykinesia, postural instability, and resting tremor (Antony et al., 2013; Pagano et al., 2016). The most affected cells are dopaminergic neurons from the substantia nigra pars compacta (SNc). The pathological hallmark of PD is the formation of Lewy bodies containing aggregated α -synuclein (Hijaz and Volpicelli-Daley, 2020). While increasing age is a risk factor for PD, the average age of onset is after 60 years old (Hindle, 2010; Parkinson Canada, 2010). The etiology of PD is multifactorial where genetics (familial PD) and environmental (sporadic PD) factors take part in disease onset (Klein and Westenberg, 2012). Familial PD accounts for 10–15% of all PD cases whereas the remainder is classified as sporadic PD (Verstraeten et al., 2015). Genetically linked PD is inherited in an autosomal dominant or recessive fashion (Ball et al., 2019). Research has identified seven causal genes for familial PD including phosphatase and tensing homolog-induced Kinase-1 (PINK1), Parkinson protein 7 (PARK7), parkin RBR E3 ubiquitin protein ligase (PARK2), vacuolar protein sorting-associated protein 35 (VPS35), α -synuclein (SNCA), glucocerebrosidase (GBA) and leucine-rich repeat Kinase 2 (LRRK2) (Verstraeten et al., 2015; Kalinderi et al., 2016; Ball et al., 2019). Conversely, sporadic PD may develop from gene-environment interactions (Benmoyal-Segal and Soreq, 2006). Environmental factors associated with PD includes but are not limited to pesticides, heavy metals, and illicit drugs (Kwakye et al., 2017). Notably, individuals may respond differently to environmental factors which results in diverse symptomology of PD, thus adding to the complexity of the disease (Ball et al., 2019).

Altered Cerebral Blood Flow in Parkinson's Disease

Using non-invasive MRI in an heterogeneous PD patient population, studies revealed decreased CBF in the frontal, parietal and occipital areas, more specifically the posterior parieto-occipital cortex, cuneus, middle frontal gyri, putamen, anterior cingulate and post- and pre-central gyri (Kamagata et al., 2011; Melzer et al., 2011; Fernandez-Seara et al., 2012; Madhyastha et al., 2015). A study by Fernandez-Seara et al. (2012) reported a 20–40% decrease in CBF in PD patients compared to a control group. Studies are trying to determine if CBF changes are related to the presence of dementia in PD, or if it can be considered as a biomarker. Derejko et al. (2006) used SPECT in PD patients with dementia and demonstrated left temporo-parietal hypoperfusion compared to the group without dementia. This suggested that CBF differences between PD patients with or without dementia could represent a clinical biomarker for discriminating PD patients (Derejko et al., 2006). Another study revealed hypoperfusion in PD patients without dementia in posterior

cortical regions (posterior cingulate/precuneus) compared to healthy individuals (Syrini et al., 2017). Hypoperfusion was positively correlated with global cognitive performance and the level of motor impairment (Madhyastha et al., 2015; Syrini et al., 2017). Melzer et al. (2011) and Fernandez-Seara et al. (2012) reported CBF reduction with parietal cortex thinning in mild PD patients without dementia and proposed that CBF alterations occur in the early stages of PD.

Although studies have identified hypoperfusion in PD patients, the mechanisms underlying these changes are unknown (Biju et al., 2020). One study used a mouse model of PD (α -synuclein transgenic mice), which overexpress human WT α -synuclein. α -synuclein pathology develops before clinical symptoms and is present in both sporadic and familial forms. Using ASL-MRI analysis in this PD mouse model, authors reported a 36.6% reduction in cortical CBF in mutant mice accompanied by motor coordination impairments and olfactory bulb atrophy/dysfunction (Biju et al., 2020).

Altered Blood-Brain Barrier and Angiogenesis in Parkinson's Disease

The association of PD with altered vascular function has led studies to investigate possible players contributing to BBB (Al-Bachari et al., 2020). In animal studies, BBB disruption in the SNc has been reported (Barcia et al., 2005; Rite et al., 2007; Chao et al., 2009). While human studies investigating BBB in PD patients are sparse, there is evidence of BBB dysfunction with increased permeability in the post commissural putamen of PD patients (Kortekaas et al., 2005; Gray and Woulfe, 2015). Wardlaw et al. (2008) and Al-Bachari et al. (2020) revealed increased leakage of the BBB in PD using ASL and dynamic contrast enhanced -MRI (DCE-MRI). Authors compared PD patients with two other control groups: one with and one without known cerebrovascular disease. This comparison could determine if BBB changes are attributable to co-existing cerebrovascular disease in an aging population, or if a pattern of BBB alteration is specific to PD. Authors reported increased BBB leakage in the group with cerebrovascular disease compared to the group without cerebrovascular disease in regions previously associated with PD, including the substantia nigra, white matter, and posterior cortical regions (Al-Bachari et al., 2020).

Accumulation of α -synuclein in ECs may also contribute to BBB dysfunction and increased permeability (Elabi et al., 2021). Higher number of EC nuclei was found in the SNc of PD patients (Faucheux et al., 1999). Other EC dysfunctions were reported, such as down regulation of tight junction proteins (Kuan et al., 2016). In the 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) mouse model of PD, down-regulation of tight junction protein ZO-1 and BBB leakage were measured in the substantia nigra (Patel et al., 2011). There is also evidence of string vessel formation in brain capillaries from human PD. String vessels are described as collapsed basement membrane without endothelium and no circulatory function. An altered basement membrane was also observed in PD mice (Yang et al., 2015). VEGF, a prominent growth factor promoting angiogenesis and BBB permeability, was upregulated in the substantia nigra, but not the striatum, of

PD patients, while animal models of PD displayed parkinsonian traits following administration of exogenous VEGF into the substantia nigra (Barcia et al., 2005; Wada et al., 2006; Rite et al., 2007).

Guan et al. (2013) reported vascular degeneration in human PD, with formation of endothelial clusters, capillary network damage, and loss of capillary connections in the substantia nigra and brain stem nuclei. Authors found a larger vessel size in PD patients, while capillaries were shorter in average length, less in number and had fewer branches. These observations were also confirmed in an MPTP mouse model of PD (Guan et al., 2013; Sarkar et al., 2014). Furthermore, ultrastructural abnormalities were identified in cerebro-cortical microvessels of PD patients, including basement membrane thickening, vacuolization and pericyte degradation (Farkas et al., 2000). Structural alterations of the basement membrane can lead to pathophysiological consequences including compromised nutrient transport and cognitive disturbances (Farkas et al., 2000). Recently, a PD mouse model of α -synuclein overexpression was associated with altered vascular density at different stages of the disease (Elabi et al., 2021). The study reported that 8 month-old animals had increased vessel density compared to control mice, while 13 month-old PD mice displayed decreased vessel density, suggesting compensatory angiogenesis in the younger group (Elabi et al., 2021). Increased angiogenesis is considered an adaptative response to pathological conditions and is regulated by basement membrane proteins and their integrin receptors. These studies postulate that immature nascent vessels in PD could contribute to increased BBB permeability, as reviewed recently (Bogale et al., 2021).

Vascular Links to Alzheimer's Disease

AD accounts for 60–80% of all diagnoses of dementia (Alzheimer's Association, 2021). This progressive and debilitating neurodegenerative disease manifests with memory, attention, executive, visuospatial and perceptual impairments. AD is not only characterized by amyloid deposition, neuroinflammation, neurodegeneration and cognitive deficits, but also by cerebrovascular pathology. Indeed, an inadequate brain perfusion has been identified as an early event in the development and progression of AD (Nicolakakis and Hamel, 2011). The risk of developing AD is increased by age-associated vascular diseases such as hypercholesterolemia, hypertension, ischemic stroke, and diabetes (Kalaria, 1996; Roher et al., 2003; Casserly and Topol, 2004; Gorelick, 2004; Luchsinger et al., 2005). The AD brain is characterized by increased levels of soluble and insoluble amyloid-beta peptide ($A\beta$), derived from the amyloid protein precursor (APP), neurofibrillary tangles of hyperphosphorylated *tau* protein, neurodegeneration and neuroinflammation, and also linked with a cerebrovascular pathology (Selkoe, 2002; Iadecola, 2004; Querfurth and LaFerla, 2010). The latter is identified post-mortem by $A\beta$ deposition in brain vessels (cerebral amyloid angiopathy, CAA), $A\beta$ -induced oxidative stress, and alterations of the vessel wall that included fibrosis and degeneration of ECs (Buee et al., 1994; Vinters et al., 1994; Zarow et al., 1997; Farkas and Luiten, 2001; Humpel and Marksteiner, 2005). Various mouse models of AD have been developed, most mimicking

the overproduction of A β through transgene expression of mutated human APP (hAPP) combined or not with the amyloidogenic presenilin (PS1) or the pathologic *tau* (Mucke et al., 2000; Oddo et al., 2003; Gotz and Ittner, 2008). These models recapitulate AD's cerebrovascular pathology in addition to the cognitive deficits, senile plaques, A β -induced oxidative stress, neuroinflammation, cholinergic denervation, synaptic failure, and cerebral hypometabolism (Hsia et al., 1999; Palop et al., 2003; Aucoin et al., 2005; Tong et al., 2005; Nicolakakis et al., 2008; Iturria-Medina et al., 2016; Love and Miners, 2016; Liu et al., 2018; Czako et al., 2020). It is in fact estimated that up to 45% of all dementias worldwide are partly, or wholly, due to age-related SVD of the brain (Montagne et al., 2016; Clancy et al., 2021). This suggests that AD and vascular dementia share common grounds, which complicates their stratification. As such, it is of utmost importance to improve our understanding of vascular underpinnings of AD (Willis and Hakim, 2013). Clinical studies that attempted to reduce plaque load by blocking A β production, removing A β with antibodies, or preventing tau phosphorylation, have all failed to alleviate AD symptoms (Korte et al., 2020). However, mounting evidence demonstrates that the brain vasculature is the missing link (Sweeney et al., 2019). Early cerebrovascular dysfunction in AD leads to decreased A β clearance, vascular oxidative stress, inflammatory damage and impaired BBB function (Zlokovic, 2005). Here below we will succinctly describe vascular underpinnings of AD, from alterations in CBF to BBB dysfunction, topics that have been extensively reviewed elsewhere (Bell and Zlokovic, 2009; Zlokovic, 2011; Hamel, 2015; Hays et al., 2016; Nelson et al., 2016; Kisler et al., 2017; Korte et al., 2020; Solis et al., 2020; Soto-Rojas et al., 2021).

Cerebral Blood Flow Alterations in Alzheimer's Disease

Numerous investigations on individuals diagnosed with AD observed reduced CBF (Prohovnik et al., 1988; Montaldi et al., 1990; Bressi et al., 1992; O'Brien et al., 1992; Smith et al., 1992; Minoshima et al., 1997; Mattsson et al., 2014; Mielke et al., 2014; Smith and Verkman, 2018). CBF decline can be detected prior to cognitive decline, but also before plaque deposition. The accumulation of soluble A β prior to plaque deposition has early pathogenic consequences in AD (Suo et al., 1998). Studies have demonstrated increased levels of soluble amyloid species including A β_{1-40} and A β_{1-42} in AD cases compared to age-matched controls (Suo et al., 1998; Smith and Greenberg, 2009). Both soluble A β_{1-40} and A β_{1-42} have been associated to abnormal vascular reactivity in the absence of plaque deposition or vessel wall dysfunction (Smith and Greenberg, 2009; Dietrich et al., 2010). In particular, studies have revealed that application of exogenous A β_{1-40} to mouse neocortex *in vivo*, or to healthy bovine blood vessels *ex vivo*, leads to endothelium-dependent vasoconstriction (Thomas et al., 1996; Niwa et al., 2000). In addition, increased levels of soluble amyloid species (A β_{1-40} and A β_{1-42}) are associated with significantly reduced CBF, increased cerebral vascular resistance, decrease myogenic and vasodilator responses (Suo et al., 1998; Dietrich et al., 2010), where A β_{1-42} is equally potent to A β_{1-40} except at a higher

concentration (Dietrich et al., 2010). Soluble A β impacts vascular function through increased production of reactive oxygen species (ROS). The reaction of ROS superoxide and excess NO produces peroxynitrite. Peroxynitrite is commonly known as a toxic oxidant which contributes to endothelial dysfunction, a mechanism relevant to AD but also to other neuroinflammatory and metabolic conditions (Beckman et al., 1990; Paris et al., 1998; Tan et al., 2004; Dietrich et al., 2010; Kelleher and Soiza, 2013; Salisbury and Bronas, 2015; Incalza et al., 2018). Both A β_{1-40} and A β_{1-42} have been shown to acutely increase ROS production in cultured rat cerebral microvascular endothelial and smooth muscle cells in a dose dependent manner (Dietrich et al., 2010). Interestingly, this response was inhibited by the ROS scavenger MnTBAP (Dietrich et al., 2010). Notably, A β_{1-40} is the predominant isoform found in cerebral vessel walls and is commonly associated with vascular deposits in CAA, which will be discussed later, while A β_{1-42} is the major isoform deposited in senile plaques (Suo et al., 1998). Although this concept is still controversial, it is thought that A β_{1-42} acts as a "seed" which initiates the formation of vascular A β deposit in CAA (McGowan et al., 2005; Gireud-Goss et al., 2020).

Following A β deposition, reduction of CBF was found in the frontal, parietal and temporal cortices from individuals carrying Apolipoprotein E4 (APOE4) gene, most prevalent genetic risk factor for AD (Thambisetty et al., 2010; Michels et al., 2016). In addition, ApoE4 allele carriers displayed early impairments in cerebrovascular reactivity to a memory task (Suri et al., 2015). BOLD-fMRI, which uses blood flow changes as a surrogate to neuronal activity, detected decreased activation in areas engaged during naming and fluency tasks in AD patients compared to individuals with no risk factors (Smith et al., 1999). Decreased BOLD-fMRI responses to different cognitive tasks in early stage of AD are region-specific (Kisler et al., 2017). Most studies investigating perfusion in AD reported either CBF or CBV alterations. However, CBF alterations appear before CBV deficits during AD progression (Lacalle-Aurioles et al., 2014).

Decreased CBF is associated with poor cognitive function, and evidence suggested that lower CBF is linked with faster cognitive decline in patients with AD (Benedictus et al., 2017). Zheng et al. (2019) investigated rCBF, functional activity and connectivity in AD by combining resting-state BOLD fMRI and ASL techniques. ASL revealed decreased rCBF in AD patients in the left posterior cingulate cortex, bilateral dorsolateral prefrontal cortex, left inferior parietal lobule, right middle temporal gyrus, left middle occipital gyrus and left precuneus. In addition, they revealed decreased connectivity between regions in AD patients, which was associated with impaired cognitive performances (Alsop et al., 2000; Zheng et al., 2019). Brain regions affected by a reduction of CBF in AD patients (parietal, frontal, temporal and occipital cortices) are associated with cognitive impairment in all domains (language, global cognition, memory, attention, executive functioning and visuospatial functioning) (Leeuwis et al., 2017).

Blood flow reductions have also been identified in early preclinical AD, before A β plaque deposition (Nicolakakis and Hamel, 2011; Iturria-Medina et al., 2016; Szu and Obenaus, 2021).

Early reduction of CBF has been reported in mouse models of AD, such as mice overexpressing mutant forms of APP (Niwa et al., 2002; Ongali et al., 2010; Lacoste et al., 2013) and in mice expressing the *ApoE4* gene allele (Lin et al., 2017). In some brain areas, CBF reduction can reach over 50%. This CBF reduction has been associated with cognitive changes in mice, including a loss of ability to sustain attention (Marshall et al., 2001). Both *ApoE4* transgenic and APP/PS1 mice revealed CBF reduction prior to neuronal and synaptic dysfunctions (Guo et al., 2019; Montagne et al., 2021).

While decreased CBF in AD is widely accepted, studies are only starting to identify underlying mechanisms, for example the involvement of pericytes. Pericytes have been linked to hypoperfusion and increased capillary constriction in AD (Bell et al., 2010; Korte et al., 2020). Pericyte-deficient transgenic mice with no A β pathology develop early CBF reduction in the gray matter, even with normal neuronal activity, endothelial-dependent vasodilation, astrocyte number and blood vessels coverage (Bell et al., 2010; Kisler et al., 2017). As these pericyte-deficient mice age, neuronal dysfunction and degeneration start to emerge. Another underlying mechanism was reported by Cruz Hernandez et al. (2019), demonstrating that capillaries become blocked by neutrophils, while another study revealed increased formation of occlusive thrombi in AD mice (Cortes-Canteli et al., 2019). Inhibiting neutrophils adhesion using an antibody against neutrophil-specific protein Ly6G in the APP/PS1 mouse model led to rapid improvements in CBF (Cruz Hernandez et al., 2019). In a follow-up study, the same group assessed the impact of one treatment of anti-Ly6G on short-term memory function and reported increased CBF by 17% in 21–22 months old APP/PS1 mice. Furthermore, they suggested that increased CBF improved cognition into late stages of AD mice (Bracko et al., 2020). Reduced neurovascular coupling and cerebrovascular reactivity have also been reported in AD mice (Girouard and Iadecola, 2006; Tong et al., 2019). Recently, impaired capillary endothelial inward rectifying Kir2.1 channel, playing a role in mediating blood delivery, has been associated with AD (Mughal et al., 2021). In a model of familial AD (5xFAD) where Kir2.1 channel function is impaired, systemic administration of the co-factor phosphatidylinositol 4,5-bisphosphate (PIP₂), required for Kir2.1 activity, led to increased CBF and functional neurovascular coupling in 5xFAD mice (Mughal et al., 2021).

AD patients are often (80–90%) diagnosed with CAA, a vessel disorder (Gireud-Goss et al., 2020) and an important risk factor for intracerebral hemorrhage and cognitive impairment (Reijmer et al., 2016). CAA consist of vascular amyloid deposits similar to senile plaques in AD (Kumar-Singh et al., 2005). Neuropathological studies have revealed that CAA affects the outer leptomeningeal vessels on the surface of the brain as well as distal intraparenchymal arteries, arterioles, and capillaries (Gireud-Goss et al., 2020; Howe et al., 2020). APP23 mouse model and human AD brain revealed an association between CAA-related capillary occlusion with CBF disturbances, hypoperfusion, detected by magnetic resonance angiopathy (MRA), which could explain in part the changes in CBF measured in AD patients (Thal et al., 2009; Milner et al., 2014). As in AD,

patients with CAA have been linked to altered hemodynamics during visual stimulation as evidenced by reduced amplitude of BOLD response (Smith et al., 2008; Dumas et al., 2012; Switzer et al., 2020).

Altered Blood-Brain Barrier and Angiogenesis in Alzheimer's Disease

Early signs of BBB leakage in AD have been detected before dementia onset (Montagne et al., 2016). Neuroimaging techniques have evidenced BBB breakdown in AD in gray and white matter brain regions (Montagne et al., 2016; van de Haar et al., 2016). A β and tau pathologies contribute to increased BBB permeability in AD patients and mouse models (Park et al., 2011; Sagare et al., 2013; Alata et al., 2015). Several players involved in A β clearance, and closely related to the BBB, are reduced in AD patients, including phosphatidylinositol-binding clathrin assembly protein (PICALM), allows for A β exocytosis across the luminal part of the BBB, P-glycoprotein (expressed on both sides of the BBB) and glucose transporter (GLUT)1 (Mooradian et al., 1997; Chiu et al., 2015; Zhao et al., 2015). AD brain microvessel show diminished expression of LRP1, a major A β clearance receptor at the BBB (Deane et al., 2004; Donahue et al., 2006). LRP1 is an ApoE receptor and is expressed at the abluminal side of brain ECs and mediates the internalization of soluble A β (Deane et al., 2004). Endothelium-specific deletion of LRP1 leads to the acceleration of A β pathology in APP-overexpressing APP_{sw/0} mice (Storck et al., 2016). Moreover, studies have demonstrated low levels of GLUT1 in AD brain endothelium, which alters glucose transport (Kalaria and Harik, 1989; Simpson et al., 1994).

Several features lead to increased BBB permeability in AD, including reduced expression of tight junctions, perivascular accumulation of blood-derived products, degeneration of pericytes and ECs, as well as infiltration of circulating leukocytes (Sweeney et al., 2018; Huang et al., 2020). It was demonstrated that A β disrupts tight junctions and increases vascular permeability by suppressing expression of ZO-1, claudin-5 and occludin while increasing expression of MMP-2 and MMP-9 (Kook et al., 2012; Blair et al., 2015; Wan et al., 2015; Huang et al., 2020). Isolated rat cerebral cortical ECs treated with A β _{1–42} displayed decreased expression of occludin and redistribution of claudin-5 and ZO-2 in the cytoplasm while in untreated cells, both claudin-5 and ZO-2 were distributed along the plasma membrane at cell-cell contacts (Marco and Skaper, 2006). In addition, studies have reported leakage of blood-derived proteins (fibrinogen, thrombin, albumin, and IgG) around capillaries from post-mortem brain tissue in the prefrontal and entorhinal cortex as well as in hippocampus of AD patients (Ryu and McLarnon, 2009; Hultman et al., 2013; Sengillo et al., 2013). Furthermore, animal studies revealed that lacking pericyte-derived soluble factors, required for a healthy endothelium, can contribute to endothelial degeneration in AD (Bell et al., 2010). Finally, mouse models of AD have demonstrated that pericyte reduction is associated with BBB dysfunction as well as accelerated buildup of A β and tau pathology (Sagare et al., 2013). In human studies, there is also evidence of pericyte loss in the hippocampus and cortex of AD

patients due in part to prolonged exposure to A β peptides (Sagare et al., 2013; Sengillo et al., 2013; Huang et al., 2020). Of note, pericytes play a role in A β clearance by internalizing different A β peptides using the LRP1 pathway (Sagare et al., 2013).

Evidence of reduced capillary length and basement membrane changes in AD patients have been reported (Salloway et al., 2002; Sengillo et al., 2013; Halliday et al., 2016). It was shown that AD patients display abnormal angiogenesis due to low expression of MEOX2, a regulator of vascular differentiation, as well as premature pruning of capillary networks resulting in reductions of CBF (Wu et al., 2005; Grammas, 2011). Endothelial degeneration including reduction of EC thickness, length and density of blood vessels were reported in brain tissue from AD patients (Sweeney et al., 2018). An increase of pro-angiogenic factors in the AD brain, without the increase in vasculature, was also reported (Grammas, 2011). Notably, the increased A β species and plaques in AD have anti-angiogenic effects (Parodi-Rullan et al., 2020), and impaired angiogenesis was identified in transgenic AD mice (Grammas, 2011). Emerging evidence suggest that dysfunction of the VEGF-A/VEGFR2 pathway may play an aggravating role in neurodegeneration and AD. For instance, sustained brain delivery of VEGF via injectable hydrogels was protective against quinolinic acid-induced neurodegeneration (Emerich et al., 2010), and low VEGF levels have been associated to another debilitating neurological disorder, spinocerebellar ataxia type 1 (Cvetanovic et al., 2011). A β acts as an antagonist of VEGF signaling via sequestration of VEGF-A in senile plaques, and also via inhibition of VEGFR2 tyrosine phosphorylation (Patel et al., 2010). Moreover, implantation of VEGF secreting microcapsules on the cerebral cortex of APP/PS1 mice attenuated both brain A β burden and cognitive impairments (Spuch et al., 2010). Whether impaired neural perfusion and increased neurotoxicity in AD correlate to a loss of VEGF function, and whether VEGF overexpression is neuroprotective in transgenic AD mice remains to be explored.

CAA is associated with increased BBB permeability and arterial stiffness (Magaki et al., 2018; Gireud-Goss et al., 2020). A β deposition in CAA has been found to occur on the cerebrovascular basement membrane of arteries, arterioles and on the basal lamina of capillaries as shown by electron microscopy (Gireud-Goss et al., 2020). Moreover, ultrastructural studies of CAA demonstrated a thinned endothelium, shrinkage and degeneration of ECs, as well as vessel occlusion, all of which can lead to CBF disturbances and microinfarcts (Attems and Jellinger, 2004; Thal et al., 2009; Magaki et al., 2018). Tight junction proteins in CAA-laden vessels are found decreased (Tai et al., 2010). After exposure to exogenous A β , human ECs showed decreased expression of occludin, while post-mortem brain tissue of CAA patients revealed decreased expression of claudin-5, ZO-1, CD31 and basement protein collagen IV (Tai et al., 2010; Carrano et al., 2011; Magaki et al., 2018). In addition, CAA patients displayed increased expression of MMP-2 and MMP-9, which may lead to basement membrane degradation and increased BBB permeability (Carrano et al., 2011). In the Tg2576 mouse model of CAA, BBB integrity was compromised due to decreased expression of claudin-5 and claudin-1 (Carrano et al., 2011). Moreover, TgSwDI mice,

another model of CAA, revealed spontaneous hemorrhage and loss of BBB integrity (Davis et al., 2004). Soluble A β _{1–40}, predominant amyloid isoform in vessel walls, also leads to tight junction redistribution at the BBB and decreased transendothelial electric resistance (Hartz et al., 2012; Gireud-Goss et al., 2020). Understanding the impact of A β in CAA and AD is essential for slowing cerebrovascular disease progression.

ADDITIONAL REMARKS: VASCULAR DEFICITS IN DOWN SYNDROME, TRAUMATIC BRAIN INJURY AND DEPRESSION

In addition to neurodevelopmental disorders discussed earlier in this review, Down syndrome (DS), which results from trisomy of human chromosome 21, is a cause of early onset Alzheimer's disease-dementia (AD-DS) (Ballard et al., 2016; Tosh et al., 2021). Two-thirds of individuals with DS will develop dementia by the age of 65 (Tosh et al., 2021). The onset of AD in DS patients parallels the development of the classic brain pathological lesions seen in AD patients without DS (Salehi et al., 2016). DS and AD disorders have genetic similarities, as individuals with DS possess a triplication of the gene encoding APP, while patients with familial AD have an extra copy of the APP gene (Salehi et al., 2016). In rodent studies of DS-AD, triplication of chromosome 21 genes other than *APP* demonstrated increased A β aggregation deposition and cognitive deficits (Wiseman et al., 2018). A recent study, focused on a model of DS comprising of a mutation in a Down syndrome critical region (Hsa21) on chromosome 21 encompassing 21q21–21q22.3 (Li et al., 2016; Tosh et al., 2021). This study crossed an Hsa21 mouse model of DS with partial trisomies other than *APP* with a transgenic APP mouse model and revealed that an additional copy of genes of the Hsa21 region modulates APP/A β biology, including A β aggregation and mortality (Tosh et al., 2021). Despite striking similarities between AD and DS in terms of genetics and symptoms onset, neurovascular impairments in DS have been largely overlooked. As such, studies aimed at elucidating vascular abnormalities in DS represent an unmet clinical need.

Early vascular insults following a traumatic brain injury (TBI) can also increase the risk of late-onset neurological diseases (Brett et al., 2021). TBI is a significant public health problem associated with long-term disabilities. Early chronic TBI may lead to secondary injury with pathophysiological changes similar to those observed in neurodegenerative diseases (Impellizzeri et al., 2016). For instance, neuroinflammation plays a fundamental role in TBI, including reactive microglia and astrocytes, as well as release of pro-inflammatory cytokines and chemokines that may hinder the brain's ability to repair itself and lead to neurodegeneration following prolonged activation of these processes (Impellizzeri et al., 2016; Brett et al., 2021). Severe or repeated mild TBI can initiate long-term neurodegeneration with signs of AD (Mendez, 2017). For example, various contact-sport players developed TBI-associated dementia or parkinsonism years after retiring. TBI can induce acute BBB disruption through

TABLE 1 | Major altered features associated with CBF, BBB, and angiogenesis in neurodevelopmental and neurodegenerative disorders.

Disorder	Key features	Selected references
ASD		
Altered CBF	<ul style="list-style-type: none"> – Widespread cerebral hypoperfusion in 75% of ASD children associated with language deficits, impaired executive function and abnormal response to sensory stimuli. – Hyperperfusion identified in frontotemporal regions. – Reduced hemodynamic responses. – Cerebral hypoperfusion also identified in rodent models of ASD. – Increased resting CBF and decreased NVC in an adult mouse model of ASD associated with endothelial dysfunction. 	Ohnishi et al., 2000; Zilbovicius et al., 2000; Burrioni et al., 2008; Reynell and Harris, 2013; Jann et al., 2015; Ouellette et al., 2020; Uratani et al., 2019
Altered BBB and angiogenesis	<ul style="list-style-type: none"> – Reduced level of adhesion molecules (CD31 and P-selectin). – Increased MMP-9 which regulates cell proliferation, adhesion, angiogenesis, oxidative injury and BBB breakdown. – Altered expression of claudin-5 and claudin-12. – Increased BBB permeability and impaired angiogenesis in animal models. – Reduced angiogenesis found in a mouse model of ASD. 	Onore et al., 2012; Kumar et al., 2015; Azmitia et al., 2016; Fiorentino et al., 2016; Turner and Sharp, 2016; Ouellette et al., 2020
Schizophrenia		
Altered CBF	<ul style="list-style-type: none"> – Increased CBF in the cingulate gyrus and superior frontal gyrus associated with positive symptoms. – Negative symptoms associated with hypoperfusion in the superior temporal gyrus bilaterally and left middle frontal gyrus. – rCBF alterations depend on severity of positive symptoms. – Increased CBF in the right superior temporal gyrus and caudate nucleus. – Decreased CBF in the occipital and left parietal cortices. – Altered NVC including reduced amplitude of response and delayed hemodynamics. 	Sabri et al., 1997; Carter et al., 2001; Schultz et al., 2002; Malaspina et al., 2004; Ford et al., 2005; Pinkham et al., 2011; Liu et al., 2012; Kawakami et al., 2014; Pu et al., 2016; Zhuo et al., 2017
Altered BBB and angiogenesis	<ul style="list-style-type: none"> – Increased BBB permeability. – Thickening and deformation of basal lamina, vacuolation of EC cytoplasm, swelling of astrocyte end-feet, activation of microglial cells and atypical vascular arborization in prefrontal and visual cortices. – Decreased claudin-5 expression, altered level of VE-cadherin and occludin in ECs. – Impaired angiogenesis and VEGF upregulation in the prefrontal cortex linked to vascular hyperpermeability. 	Grove et al., 2015; Hino et al., 2016; Casas et al., 2018; Carrier et al., 2020; Cai et al., 2020; Guo et al., 2020; Crockett et al., 2021; Usta et al., 2021
MS		
Altered CBF	<ul style="list-style-type: none"> – Hypoperfusion in SP-MS, RR-MS and PP-MS patients. – Active demyelinating lesions associated with hyperperfusion and stable lesions linked to hypoperfusion. – CBF alterations present in early stages of disease. – Impaired cerebral vascular reactivity leads to neuronal death. – Overproduction of NO desensitize EC and smooth muscle cell function, leading to decreased vasodilatory capacity and limited blood supply to neurons. 	Ge et al., 2005; Varga et al., 2009; D'Haeseleer et al., 2011; Ota et al., 2013; Marshall et al., 2014; Bester et al., 2015; Monti et al., 2018; Hostenbach et al., 2019
Altered BBB and angiogenesis	<ul style="list-style-type: none"> – BBB hyperpermeability. – Decreased expression of TJ proteins (ZO-1, occludin and claudin-5) in ECs in patients with active and inactive lesions. – Rodent model of MS show increased expression of VEGF in ECs, astrocytes and monocytes. – Increased vascular network density and angiogenesis. 	Kirk et al., 2003; Bennett et al., 2010; Holley et al., 2010; Cramer et al., 2014; Girolamo et al., 2014; Papadaki et al., 2014
HD		
Altered CBF	<ul style="list-style-type: none"> – Altered CBF prior to structural changes and motor symptoms. – Cerebral hypoperfusion in the basal ganglia, medial and lateral prefrontal cortex. – Cerebral hyperperfusion in the cerebellar-thalamic and somatosensory regions. – Altered neurovascular coupling during visual stimulation. 	Hasselbalch et al., 1992; Sax et al., 1996; Deckel and Duffy, 2000; Wang et al., 2016; Klinkmueller et al., 2021

(Continued)

TABLE 1 | (Continued)

Disorder	Key features	Selected references
Altered BBB and angiogenesis	<ul style="list-style-type: none"> – Increased vessel density, BBB leakage and VEGF-A release. – Decreased TJ molecules including occludin and claudin-5. – Rodent model of HD revealed increased transcytosis and paracellular transport in brain ECs with TJ imbalance. – mHtt aggregates localized in ECs, smooth muscle cells and perivascular macrophages. – iPSCs-derived HD BMECs show increased angiogenesis, altered barrier properties and impaired Wnt/β-catenin signaling. 	Steventon et al., 2020; Drouin-Ouellet et al., 2015; Di Pardo et al., 2017; Lim et al., 2017
PD		
Altered CBF	<ul style="list-style-type: none"> – Decreased CBF in frontal, parietal and occipital areas. – PD patients with dementia show left temporo-parietal hypoperfusion. – PD patients without dementia display hypoperfusion in the posterior cortical regions. – Hypoperfusion is positively correlated with cognitive performance and motor impairment. 	Derejko et al., 2006; Kamagata et al., 2011; Fernandez-Seara et al., 2012; Madhyastha et al., 2015; Syrimi et al., 2017
Altered BBB and angiogenesis	<ul style="list-style-type: none"> – BBB disruption in the SNc with increased permeability in the post-commissural putamen. – Down regulation of TJ proteins (ZO-1) and higher number of EC nuclei in the SNc. – String vessel formation in brain capillary networks. – Upregulation of VEGF, and parkinsonian traits following VEGF administration in rodent models. – Formation of endothelial clusters, capillary network damage, loss of capillary connections in the SN, basement membrane thickening, vacuolization, and pericyte degradation. 	Farkas et al., 2000; Barcia et al., 2005; Kortekaas et al., 2005; Wada et al., 2006; Rite et al., 2007; Chao et al., 2009; Patel et al., 2011; Guan et al., 2013; Yang et al., 2015; Kuan et al., 2016
AD		
Altered CBF	<ul style="list-style-type: none"> – Reduced CBF prior to cognitive decline and plaque deposition. – Soluble Aβ_{1-40} and Aβ_{1-42} are associated with abnormal vascular reactivity and decreased myogenic responses in absence of plaque deposition. – Hypoperfusion detected following Aβ deposition in the frontal, parietal and temporal cortices and poor cognitive function. – BOLD-fMRI detected decreased activation in regions involved in naming and fluency tasks. – Hypoperfusion identified in rodent models overexpressing mutant forms of APP. – Rodent models show reduced NVC and cerebrovascular reactivity. – Parallel diagnosis of CAA linked with altered hemodynamics, capillary occlusion and hypoperfusion. 	Montaldi et al., 1990; Bressi et al., 1992; Smith et al., 1999; Marshall et al., 2001; Girouard and Iadecola, 2006; Smith and Greenberg, 2009; Dietrich et al., 2010; Ongali et al., 2010; Dumas et al., 2012; Lacoste et al., 2013; Mattsson et al., 2014; Milner et al., 2014; Benedictus et al., 2017; Smith and Verkman, 2018
Altered BBB and angiogenesis	<ul style="list-style-type: none"> – Aβ and tau pathologies contribute to BBB breakdown, reduced expression of TJ (ZO-1, claudin-5, occludin) and degeneration of pericytes and ECs. – Brain microvessel with diminished expression of LRP1. – Reduced level of GLUT1 in brain endothelium. – Reduced capillary length with basement membrane alterations. – Abnormal angiogenesis related to low expression of MEOX2. – Reduced EC thickness, and lower length/density of blood vessels. – Dysfunction of the VEGF-A/VEGFR2 pathway aggravates neurodegeneration. – Rodent models show pericyte loss. – Aβ deposition in CAA linked to decreased TJ proteins, increased expression of MMP-2 and MMP-9, thinned endothelium, degeneration of ECs and leaky BBB. 	Kalaria and Harik, 1989; Simpson et al., 1994; Emerich et al., 2010; Tai et al., 2010; Grammas, 2011; Carrano et al., 2011; Sagare et al., 2013; Halliday et al., 2016; Montagne et al., 2016; van de Haar et al., 2016; Magaki et al., 2018; Sweeney et al., 2018; Huang et al., 2020

Selected references are displayed. A β , β -amyloid peptide; AD, Alzheimer's disease; APP, amyloid precursor protein; ASD, autism spectrum disorders; BBB, blood brain barrier; BMECs, brain microvascular endothelial cells; BOLD-FMRI, blood oxygen level dependent imaging-functional magnetic resonance imaging; CAA, cerebral amyloid angiopathy; CBF, cerebral blood flow; ECs, endothelial cells; GLUT1, glucose transporter 1; HD, Huntington's disease; iPSC, induced pluripotent stem cells; LRP1, low-density lipoprotein receptor-related protein 1; MEOX2, Mesenchyme Homeobox 2; mHtt, mutant huntingtin; MMP, matrix metalloproteinases; MS, multiple sclerosis; NO, nitric oxide; NVC, neurovascular coupling; PD, Parkinson's disease; PP-MS, primary progressive-multiple sclerosis; rCBF, regional cerebral blood flow; RR-MS, relapsing remitting-multiple sclerosis; SN, substantia nigra; SNc, substantia nigra pars compacta; SP-MS, secondary progressive-multiple sclerosis; TJ, tight junctions; VE-cadherin, vascular endothelial cadherin; VEGF, vascular endothelial growth factor; VEGFR2, vascular endothelial growth factor receptor 2; Wnt/ β -catenin, Wntless-related integration site β -catenin; ZO-1, Zonula occludens-1.

vascular shear stress, hemorrhages, edema, alterations in CBF and chronic inflammation, which is known to contribute to A β deposition and tau pathology (Iadecola, 2013; De Silva and Faraci, 2016). Autopsies of TBI patients show diffuse A β plaques similar to those identified in AD, as reviewed by Perry et al. (2016). The formation of A β in perivascular spaces following TBI may lead to an injury cascade consisting of cerebrovascular damage, oxidative stress and ECs dysfunction (Ramos-Cejudo et al., 2018). Interestingly, alterations in EC survival, BBB integrity and neuroinflammation are considered early events after TBI, all of which are characteristic of cerebrovascular damage involved in the progression of AD and impairment of A β clearance. Thus, these early vascular impairments promote the onset of neurodegenerative diseases (Ramos-Cejudo et al., 2018). Considering early vascular injuries in TBI, biomarker studies are integrating a variety of neuroimaging and molecular techniques to better understand the incidence of cerebrovascular dysfunction and the onset of neurodegenerative diseases, and therapeutic investigations have looked at ways to improve cerebrovascular function (Graham and Sharp, 2019; Martinez and Stabenfeldt, 2019).

One of the leading causes of mental illness worldwide, depression, has a tremendous impact on psychosocial behaviors and vascular health (Knight and Baune, 2017; Menard et al., 2017). Chronic stress is the primary environmental risk factor for depression. The nucleus accumbens (NAc) is one of the main players in regulating stress response (Russo and Nestler, 2013). Menard et al. (2017) have demonstrated that chronic social stress induces BBB leakiness in the NAc of mice, which leads to circulating proinflammatory mediators and depression-like behaviors such as helplessness, social avoidance and anhedonia. As seen in neurodevelopmental and neurodegenerative disorders, the increase in BBB permeability in the rodent model of chronic social stress was facilitated by the loss of tight junction protein claudin-5 (Menard et al., 2017). Furthermore, stress-induced BBB permeability has been linked to inflammation of the endothelium and up-regulation of an epigenetic repressor, *hdac1*, which is involved in reducing claudin-5 expression and loosening of tight junctions (Dudek et al., 2020). Consequently, these studies are highlighting mechanisms by which chronic stress impacts vascular health, which could have long-term consequences on brain maturation and aging.

The vascular system, as any other system, undergoes aging. It has been hypothesized that vascular aging leads to a progressive functional deterioration (Grunewald et al., 2021). During aging, the brain vasculature undergoes several changes including decreased capillary density, attenuation of neovascularization potential, increased BBB permeability and decreased CBF as reviewed in Watanabe et al. (2020) and Banks et al. (2021). A suggested mechanism of typical vascular aging consist of the inability of VEGF to replenish vessel loss. The mechanisms by which VEGF is involved in vascular aging are unknown. However, mice treated with VEGF have been shown to live longer, with extended multiorgan functionality (Grunewald et al., 2021). Furthermore, aging is associated with several vascular changes including aortic stiffness which has been

linked to reduced blood flow in tissues leading to increased neuroinflammation and neurodegeneration later in life (Moore et al., 2021). Therefore, age-related changes in key vascular features may predispose to age-associated diseases (Banks et al., 2021). Improving early pathological conditions by protecting the brain vasculature is essential in preventing or modulating disease progression.

CONCLUSION

Vascular risk factors and co-morbidities take part in disease onset and/or exacerbate disease progression (Sweeney et al., 2018; Clancy et al., 2021). When it comes to alterations in CBF, BBB, and vascular patterning, neurodevelopmental and neurodegenerative disorders share interesting similarities (Table 1). While these disorders are siloed, mainly due to the age of onset, the commonalities in vascular alterations force to question the implication of early life vascular impairments on the expression of age-related neurodegenerative diseases. The vascular implications in middle-aged autistic adults have been largely overlooked, 10% of individuals diagnosed with ASD age between 40 and 60 years old will develop dementia, including AD within 15 years (Plana-Ripoll et al., 2019). In addition, there is a high frequency of parkinsonism among older ASD patients (Starkstein et al., 2015). The impact of altered brain perfusion and BBB integrity in ASD may contribute to the onset of neurodegenerative diseases due to the continuous vascular impairments associated with these diseases. Likewise, schizophrenia is associated with an elevated risk for developing Alzheimer's and Parkinson's diseases as they share core features including white matter abnormalities and cognitive deficits (Ribe et al., 2015; Kochunov et al., 2021; Kuusimäki et al., 2021).

Since fast-growing evidence demonstrates the role of early vascular impairments in the onset and/or progression of numerous neurological conditions, more work is needed to identify therapeutic targets to promote healthy cerebrovascular maturation and aging, as well as hinder the progression of age-related dementia and neurodegeneration. This is primordial considering recent findings that ECs show limited turnover compared to other cells in the human body (Sender and Milo, 2021). For instance, it was estimated that the turnover rate of ECs is 0.1% per day, as opposed to much higher rates for erythrocytes (65%), neutrophils (18%) or gastrointestinal epithelial cells (12%). In addition, the turnover rates of cellular mass in the human body were estimated at 0.4% for ECs, 4% for skin cells and adipocytes, and 42% for gastrointestinal epithelial cells (Sender and Milo, 2021). Hence, as ECs are long-lived, they may carry on early structural and functional impairments into adulthood and throughout aging, altering organ function in the long term. This concept emphasizes the importance of infant screening for cerebrovascular abnormalities, and of continuous management of vascular risk factors during lifespan. As such, the vascular continuum between neurodevelopmental and neurodegenerative disease should represent a growing focus in modern neuroscience (Figure 3).

AUTHOR CONTRIBUTIONS

JO wrote the draft following BL's instructions. BL chose the theme and edited the manuscript. Both authors contributed to the article and approved the submitted version.

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Carbonic Anhydrases as Potential Targets Against Neurovascular Unit Dysfunction in Alzheimer's Disease and Stroke

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The Neurovascular Unit (NVU) is an important multicellular structure of the central nervous system (CNS), which participates in the regulation of cerebral blood flow (CBF), delivery of oxygen and nutrients, immunological surveillance, clearance, barrier functions, and CNS homeostasis. Stroke and Alzheimer Disease (AD) are two pathologies with extensive NVU dysfunction. The cell types of the NVU change in both structure and function following an ischemic insult and during the development of AD pathology. Stroke and AD share common risk factors such as cardiovascular disease, and also share similarities at a molecular level. In both diseases, disruption of metabolic support, mitochondrial dysfunction, increase in oxidative stress, release of inflammatory signaling molecules, and blood brain barrier disruption result in NVU dysfunction, leading to cell death and neurodegeneration. Improved therapeutic strategies for both AD and stroke are needed. Carbonic anhydrases (CAs) are well-known targets for other diseases and are being recently investigated for their function in the development of cerebrovascular pathology. CAs catalyze the hydration of CO₂ to produce bicarbonate and a proton. This reaction is important for pH homeostasis, overturn of cerebrospinal fluid, regulation of CBF, and other physiological functions. Humans express 15 CA isoforms with different distribution patterns. Recent studies provide evidence that CA inhibition is protective to NVU cells *in vitro* and *in vivo*, in models of stroke and AD pathology. CA inhibitors are FDA-approved for treatment of glaucoma, high-altitude sickness, and other indications. Most FDA-approved CA inhibitors are pan-CA inhibitors; however, specific CA isoforms are likely to modulate the NVU function. This review will summarize the literature regarding the use of pan-CA and specific CA inhibitors along with genetic manipulation of specific CA isoforms in stroke and AD models, to bring light into the functions of CAs in the NVU. Although pan-CA inhibitors are protective and safe, we hypothesize that targeting specific CA isoforms will increase the efficacy of CA inhibition and reduce side effects. More studies to further determine specific CA isoforms functions and changes in disease states are essential to the development of novel therapies for cerebrovascular pathology, occurring in both stroke and AD.

Keywords: Alzheimer's disease, stroke, carbonic anhydrase (CA), neurovascular unit (NVU), cerebrovascular pathology, amyloid beta, inflammation, mitochondria

INTRODUCTION

Neurovascular dysfunction is an important, early and causative event in the pathogenesis of both Alzheimer's disease (AD) and Stroke (Iadecola, 2010, 2017; Hachinski et al., 2019; Sweeney et al., 2019a; Freitas-Andrade et al., 2020; Zlokovic et al., 2020). Indeed, the Neurovascular Unit (NVU) has recently gained a lot of momentum as a pharmacological target in cerebrovascular pathologies and neurodegeneration (Iadecola, 2017; Sweeney et al., 2019a; Zlokovic et al., 2020). The NVU is a functional multicellular structure composed of blood vessels and different cell types surrounding them within the central nervous system (CNS), and is instrumental in regulating CNS homeostasis (Andreone et al., 2015; Iadecola, 2017; Sweeney et al., 2019b; Freitas-Andrade et al., 2020). Important functions of the NVU include regulation of cerebral blood flow (CBF) and immunological surveillance, amongst others (Andreone et al., 2015; Cortes-Canteli and Iadecola, 2020; Freitas-Andrade et al., 2020). NVU dysfunction is observed in aging, AD and following neurological pathologies such as stroke (Iadecola, 2017; Cortes-Canteli and Iadecola, 2020; Freitas-Andrade et al., 2020; Sarvari et al., 2020) and traumatic brain injury (Xing et al., 2012; Lok et al., 2015), among others.

Stroke and dementia are the two most common neurological disorders. They confer risks for each other and share some, mostly modifiable, risk factors. Having a stroke doubles the chance of developing dementia (Savva and Stephan, 2010; Kuźma et al., 2018). Therefore, preventing stroke through management of hypertension and other risk factors could also decrease the incidence of dementia (Kuźma et al., 2018; Hachinski et al., 2019).

AD is the most common form of dementia and has been historically characterized by extracellular amyloid beta (A β) plaques and intracellular hyperphosphorylated tau tangles in specific brain regions (Day et al., 2015; Castillo-Carranza et al., 2017; Cortes-Canteli and Iadecola, 2020; Ojo et al., 2021). Interestingly, A β and tau intermediate aggregation species, such as oligomers, have been shown to have toxic effects on multiple cell types of the NVU (Fossati et al., 2010, 2012b; Parodi-Rullán et al., 2019; Canepa and Fossati, 2020). This toxicity, in association with the contribution of impaired clearance of undesired material from the brain, may lead to neurodegeneration and cognitive decline (Fossati et al., 2012b; Boland et al., 2018; Da Mesquita et al., 2018; Nortley et al., 2019; Provensi et al., 2019; Braun and Iliff, 2020; Carare et al., 2020; Cortes-Canteli and Iadecola, 2020; Nedergaard and Goldman, 2020; Parodi-Rullán et al., 2020; Quintana et al., 2021). Importantly, up to 90% of AD patients also present with cerebral amyloid angiopathy (CAA), defined as A β deposition around the brain vasculature and/or within the vessel walls (Iadecola, 2017; Sweeney et al., 2019a; Zlokovic et al., 2020). CAA is also common in the non-demented elderly population and constitutes an important contributor to NVU dysfunction in both normal aging and AD (Provensi et al., 2019; Cortes-Canteli and Iadecola, 2020; Ojo et al., 2021).

Stroke is classically defined as a neurological damage attributed to an acute focal injury of the CNS by a vascular cause, including cerebral infarction, intracerebral hemorrhage

(ICH), and subarachnoid hemorrhage (SAH), and is a major cause of disability and death worldwide (Sacco et al., 2013). The most common type of stroke is Ischemic Stroke (IS) which occurs when atherosclerotic plaques and fatty deposits cause vascular occlusions, interrupting blood flow in the brain. The blood vessel most commonly occluded is the middle cerebral artery (Kuriakose and Xiao, 2020). When IS occurs, it promptly causes multiple detrimental cerebral injuries due to both the lack of oxygen and glucose, as well as the associated pro-inflammatory signaling (Faraco et al., 2007; Freitas-Andrade et al., 2020; Kuriakose and Xiao, 2020; Sarvari et al., 2020). Following this event, there is a reperfusion injury phase, which occurs when oxygen and CBF are restored (Faraco et al., 2007; Freitas-Andrade et al., 2020; Sarvari et al., 2020). Another type of stroke is hemorrhagic stroke (HS), which occurs when a blood vessel, providing blood to the brain, ruptures (Corraini et al., 2017; Sarvari et al., 2020). HS is characterized by greater lesion volume, higher intracranial pressure and induce more severe brain injury than IS. Importantly, IS and HS affect different brain regions (Corraini et al., 2017).

Shared risk factors between AD and stroke are reduced CBF, cardiovascular diseases and age (Sarvari et al., 2020; Ojo et al., 2021). Cardiovascular risk factors like obesity, diabetes, hypertension and atherosclerosis have been observed to exacerbate cerebrovascular pathology as well as neurodegeneration, including AD (Girouard and Iadecola, 2006; de Bruijn and Ikram, 2014; Cortes-Canteli and Iadecola, 2020; Sarvari et al., 2020). The common underlying mechanisms involved in both AD and stroke include neuroinflammation, mitochondrial dysfunction, cell death, and blood brain barrier (BBB) dysregulation, indicating that the NVU is a target for both diseases (Fossati et al., 2012b; Alluri et al., 2014; Sekerdag et al., 2018; Eldahshan et al., 2019; Provensi et al., 2019; Parodi-Rullán et al., 2020). A vast amount of research has been invested into discovering new treatments for AD and stroke. In AD, this has recently led to the controversial FDA-approval of aducanumab (Ferrero et al., 2016; Sevigny et al., 2017; Knopman et al., 2021). However, more research needs to be done to develop successful disease-modifying therapies for both disorders. The scientific community is particularly encouraging the study of repurposed drugs, approved by the FDA for other disorders, which could be beneficial for AD and stroke, while allowing more rapid translation to clinical trials.

This review will introduce the idea of potentially repurposing carbonic anhydrase inhibitors (CAIs), many of which are already FDA-approved for other indications, for prevention of cerebrovascular and neurovascular pathology in AD and stroke and highlight the impact of carbonic anhydrase (CA) modulation in these two dominant neurological disorders.

CAs are a family of zinc metalloenzymes which catalyze the reversible hydration of carbon dioxide to produce bicarbonate and a proton ($\text{CO}_2 + \text{H}_2\text{O} \leftrightarrow \text{HCO}_3^- + \text{H}^+$) (Supuran, 2011; Mishra et al., 2020). This chemical reaction is essential for many physiological processes, such as pH and ion homeostasis, carbon dioxide transport, electrolyte secretion, gluconeogenesis, lipogenesis, and ureagenesis, water and sodium reabsorption in the kidney, bone reabsorption and calcification, cerebrospinal

fluid formation and turnover, amongst other processes (Provensi et al., 2019; Zamanova et al., 2019; Mishra et al., 2020). CAs have been studied as a well-known pharmacological target for many peripheral and CNS disorders (Bradwell et al., 1992; Ilies et al., 2004; De Simone and Supuran, 2007; Supuran, 2008, 2018; Akocak and Ilies, 2014; Provensi et al., 2019; Zamanova et al., 2019; Mishra et al., 2020). Interestingly, many recent studies revealed a common goal to further elucidate CAs involvement in both AD and stroke.

Humans have 15 CA isoforms, all with different expression patterns at a tissue and cellular level (Supuran, 2011; Provensi et al., 2019; Zamanova et al., 2019; Mishra et al., 2020). Many of these isoforms are expressed within the NVU and it has been hypothesized that each is involved in different functions (Draghici et al., 2014; Rasmussen and Boedtker, 2018; Provensi et al., 2019; Zamanova et al., 2019). It is known that some isoforms are extracellular, anchored to the plasma membrane (CA-IV, CA-IX, and CA-XII, CA-XIV), while others are cytosolic (CA-I, CA-II, CA-III, CA-VII, CA-XIII), two are found in the mitochondria (CA-VA and CA-VB) (Boriack-Sjodin et al., 1995; Nishimori et al., 2005), some are acatalytic isoforms (CA-VIII, CA-X, and CA-XI) (Aspatwar et al., 2014), and one isoform is secreted in saliva (CA-VI) (Nishimori et al., 2007; Alterio et al., 2012; Patrikainen et al., 2014; Provensi et al., 2019; Zamanova et al., 2019; Mishra et al., 2020). CA function has been linked to AD pathology as well as stroke, but the specific isoforms and the pathological mechanisms involved are not fully understood (Wang et al., 2009; Draghici et al., 2014; Fossati et al., 2016; Pollard et al., 2016; Solesio et al., 2018; Mishra et al., 2020). The determination of each CA isoforms' role in health and disease should be a priority for the development of novel and effective therapies in cerebrovascular pathology (Provensi et al., 2019; Mishra et al., 2020).

CAIs were developed first as diuretics and have become valuable in treating glaucoma, cerebral edema, epilepsy, as well as high altitude sickness (Mincione et al., 2007; Supuran, 2008; Ritchie et al., 2012; Akocak and Ilies, 2014; Zamanova et al., 2019; Mishra et al., 2020). The FDA-approved CAIs methazolamide (MTZ) and acetazolamide (ATZ) are the most studied pan-CAIs. Their activity in the NVU will be one of the focuses of this review. Other FDA-approved CAIs such as topiramate, which has some selectivity for the mitochondrial CA-VA and CA-VB isoforms, along with compounds with selectivity for CA-IX and CA-XII will also be discussed (Scozzafava et al., 2000; Supuran, 2012; Andring et al., 2020; McDonald et al., 2020). MTZ and ATZ, along with topiramate, have been observed to have protective properties on cerebrovascular pathology, as well as on mitochondria function, an important target for NVU integrity (Wang et al., 2009; Price et al., 2012; Fossati et al., 2016; Solesio et al., 2018; Salameh et al., 2019). A substantial amount of literature on the FDA-approved pan-CAIs indicates that they are safe and can pass the BBB. However, the development of new compounds targeting specific isoforms may improve the efficacy and reduce side effects of the already existing pan-CAIs (Provensi et al., 2019; Mishra et al., 2020).

This review will first illustrate the basic structure and functions of the cells composing the NVU, specifically describing how

they become dysfunctional in stroke and AD. We will then discuss the properties of multiple CA isoforms and highlight the available evidence showing how CA inhibition may be protective toward multiple dysregulated mechanisms in NVU-composing cells, pointing to CAs as potential targets for both stroke and AD therapy (Provensi et al., 2019; Mishra et al., 2020).

THE NEUROVASCULAR UNIT: FUNCTION AND DYSFUNCTION IN STROKE AND ALZHEIMER'S DISEASE

The Neurovascular Unit

The cell types that constitute the NVU (depicted in **Figure 1**) and collaborate to perform its functions are endothelial cells (ECs), pericytes, smooth muscle cells (SMCs), astrocytes, and microglia (Iadecola, 2017; Freitas-Andrade et al., 2020), which are functionally or physically connected to neurons (Andreone et al., 2015; Cortes-Canteli and Iadecola, 2020; Freitas-Andrade et al., 2020). The NVU is the morpho-functional unit including the BBB, which is important for the transport of nutrients and oxygen from the systemic circulation to the brain, for the clearance of toxic waste from the CNS, for the connection between blood flow and neuronal function, as well as for forming a physical barrier to prevent the entrance of pathogens and other harmful entities into the CNS (Sarvari et al., 2020).

Endothelial Cells

ECs are essential components of the blood vessel wall. Cerebrovascular ECs form tight and adherent junctions with each other to limit the entry of molecules and cells from the peripheral circulation into the CNS (Reese and Karnovsky, 1967). Transporters expressed on the plasma membrane of ECs specifically regulate what enters and exits the CNS. For example, glucose enters the brain exclusively via transporters, despite the brain being a highly metabolic organ responsible for up to 25% of total glucose consumption in the body (Tang et al., 2017; Parodi-Rullán et al., 2019). ECs also regulate blood flow by releasing vasodilators and vasoconstrictors, such as nitric oxide (NO) and endothelin-1, respectively (Morikawa et al., 1994; Biernaskie et al., 2001; Freitas-Andrade et al., 2020). Due to the fact that mitochondria are very abundant in cerebrovascular ECs (Oldendorf et al., 1977; Sarvari et al., 2020), these cells are particularly sensitive to oxygen deprivation (Pun et al., 2009; Freitas-Andrade et al., 2020). Hence, pathological conditions which cause oxygen-glucose deprivation (OGD) and prompt excessive reactive oxygen species (ROS) production trigger cerebral endothelial dysfunction, cell death and BBB breakdown (Schreibelt et al., 2007; Pun et al., 2009; Lochhead et al., 2010; Ghiso et al., 2014; Fossati et al., 2016; Parodi-Rullán et al., 2019; Freitas-Andrade et al., 2020), pointing to the mitochondria as critical targets for EC function and BBB integrity. Efficient communication and exchange of materials between ECs and other cell types of the NVU is essential for CNS homeostasis (Andreone et al., 2015; Freitas-Andrade et al., 2020; Sarvari et al., 2020).

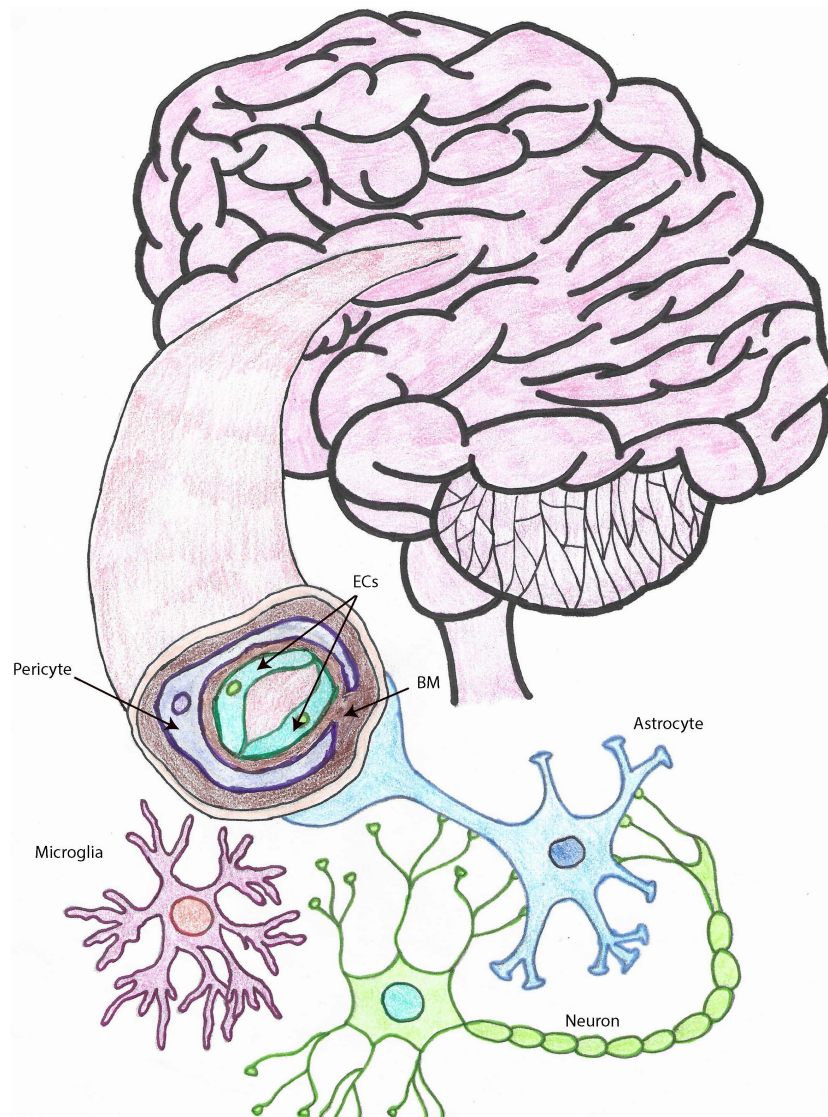


FIGURE 1 | The neurovascular unit. Drawing depicting the cell types that make up the NVU within brain capillaries. Brain capillaries are surrounded by pericytes, as shown in this figure, while arteries and arterioles are surrounded by SMCs. Other important cells associated with blood vessels and important for BBB and neurovascular functions, also represented in this drawing, are astrocytes and microglia. ECs, endothelial cells; SMCs, smooth muscle cells; BBB, blood brain barrier; NVU, neurovascular unit; MCs, microglia cells, BM, basement membrane.

Cerebrovascular dysfunction has been observed in both AD and IS (Fossati et al., 2010, 2012a,b; Guo et al., 2010; Lochhead et al., 2010; Jiao et al., 2011; Shin et al., 2016; Freitas-Andrade et al., 2020; Parodi-Rullán et al., 2020; Quintana et al., 2021). Aging, as well as cardiovascular risk factors, such as hypertension and diabetes, contribute to cerebrovascular pathology (Huang et al., 1995; Girouard and Iadecola, 2006; Price et al., 2012; Cortes-Canteli and Iadecola, 2020; Ojo et al., 2021). The reduction of tight junction proteins such as zona occludin-1 (ZO-1) and occludin is observed in models of stroke and AD (Marco and Skaper, 2006; Jiao et al., 2011; Engelhardt et al., 2014; Freitas-Andrade et al., 2020; Parodi-Rullán et al., 2020). An increase in adhesion molecules expression, such as intercellular adhesion

molecule-1 (ICAM-1) and vascular cell adhesion molecule-1 (VCAM-1), is observed in stroke, and recently, has been also associated with AD (Cruz Hernández et al., 2019; Sarvari et al., 2020). The reduction of tight junction proteins and the increase in adhesion molecules triggers the recruitment of peripheral immune cells into the brain and may be due to endothelial activation by danger associated molecular patterns (DAMPs), such as low glucose and oxygen, and aggregated proteins, like A β and hyperphosphorylated tau (Canepa and Fossati, 2020; Freitas-Andrade et al., 2020; Sarvari et al., 2020). An increase in BBB permeability allows peripheral substances to enter the CNS, leading to neuroinflammation and oxidative stress (Pun et al., 2009; Turner and Sharp, 2016; Yang et al., 2019). Increased

oxidative stress in ECs exacerbates mitochondrial dysfunction, leading to apoptosis (Lochhead et al., 2010; Fossati et al., 2012b, 2016; Solesio et al., 2018; Parodi-Rullán et al., 2019). During both ischemic injury and AD, ECs have been observed to decrease NO production, causing dysregulation of CBF (Morikawa et al., 1994; Biernaskie et al., 2001; Girouard and Iadecola, 2006; Austin et al., 2013; Parodi-Rullán et al., 2019; Freitas-Andrade et al., 2020). Interestingly, in endothelial nitric oxide synthase (e-NOS) knockout (KO) mice, deficiency of NO reduces the ability of ECs to neutralize ROS enhancing oxidative stress and neuroinflammation, likely exacerbating AD pathology (Austin et al., 2013). Accordingly, in models of AD and stroke, EC dysfunction has been shown to enhance neuroinflammation by increasing the production of ROS and enhancing BBB permeability, amongst other cellular mechanisms (Parodi-Rullán et al., 2019, 2020; Yang et al., 2019).

Pericytes and Smooth Muscle Cells

Pericytes and SMCs surround ECs within the vascular walls, wrapping around capillaries and arterioles/arteries, respectively. Both cell types have a vital role in the regulation of CBF and BBB integrity (Sagare et al., 2013; Hall et al., 2014; Freitas-Andrade et al., 2020). These cells regulate blood flow mainly through their contraction or dilation, which control the diameter of the blood vessel (Hall et al., 2014; Freitas-Andrade et al., 2020). Recently, the role of blood vessels in the clearance of CNS interstitial fluid and proteins has been thoroughly investigated (Aldea et al., 2019; Carare et al., 2020). Intramural periarterial drainage (IPAD) has been hypothesized as a route to drain the brain interstitial fluid and soluble proteins from the CNS into the cervical lymph nodes, by traveling along the capillaries and arterioles in the opposite direction of blood flow, within the basement membrane of the brain vasculature. The contraction of SMCs provides the motive force for IPAD (Carare et al., 2008; Aldea et al., 2019). This system is dysregulated in models of CAA and aging (Ojo et al., 2021). Another brain clearance pathway, the glymphatic system, has been observed to import cerebrospinal fluid along periarterial space and export interstitial fluid along perivenous spaces, in perivascular tunnels, formed by astroglial cells. This pathway has been proposed to be important in the clearance and regulation of toxic proteins in the brain such as tau and A β , and to be mediated by astrocytic aquaporin-4 (AQP4) (Braun and Iliff, 2020; Nedergaard and Goldman, 2020). Interestingly, the glymphatic system is most efficient during sleep (Boespflug and Iliff, 2018; Mestre et al., 2020).

During both stroke and AD, pericytes are observed to detach from the BBB (Sagare et al., 2013; Freitas-Andrade et al., 2020). In an AD mouse model, pericyte loss was observed to exacerbate AD pathology (Sagare et al., 2013). The loss of pericytes decreases vascular stability and decreases the ability for the NVU to regulate CBF (Sagare et al., 2013; Hall et al., 2014). On the other hand, a recent study concluded that A β induces pericyte-mediated capillary constriction, reducing CBF (Nortley et al., 2019). Pericytes have been observed to endure damage in models of diabetes as well as obesity, which are both risk factors of AD and IS (Price et al., 2012; Shah et al., 2013a). In cats exposed to hypoxia, pericytes exhibited detachment from the

microvasculature (Gonul et al., 2002). Along with dysregulation of CBF in models of AD and stroke, pericyte and SMC loss also contribute to impaired clearance in models of AD and CAA (Sagare et al., 2013; Aldea et al., 2019; Carare et al., 2020; Kim et al., 2020; Ojo et al., 2021).

Astrocytes

Astrocytes are essential cells for the NVU and provide a physical connection between blood vessels and neurons (Iadecola and Nedergaard, 2007; Braun and Iliff, 2020). Astrocytic end-feet wrap the blood vessels, helping to stabilize EC tight junctions and regulate CBF (Iadecola and Nedergaard, 2007). Astrocytes play a key role in providing metabolic and physical support to the CNS, along with the regulation of CBF, as well as brain clearance (Freitas-Andrade et al., 2020). AQP4 is a membrane protein that functions in water exchange within the CNS. In healthy individuals, AQP4 is a membrane channel localized at the astrocytic end feet, and it is important for clearance of toxic solutes from the brain (Smith et al., 2019). It is hypothesized that AQP4 is a part of the glymphatic system, mediating fluid exchange and the drainage of proteins, as well as the elimination of liquid from the CNS (Iliff et al., 2012; Boland et al., 2018; Braun and Iliff, 2020; Mestre et al., 2020). Despite the differential contributions of the glymphatic system and the IPAD pathway to the clearance of cerebral fluids and waste material are not completely grasped, it is accepted that astrocytes play a pivotal role in the exchange of fluids, based on studies showing that, in AQP4 KO mice, cerebrospinal fluid influx as well as CNS clearance were decreased (Iliff et al., 2012; Braun and Iliff, 2020). Another essential function of astrocytes is glutamate uptake and release, which is very important for the maintenance of CNS homeostasis (Dejakaisaya et al., 2021; van Putten et al., 2021; Verkerke et al., 2021). Astrocytes also have a critical role in the brain antioxidant system maintenance and in the production of glutathione, an important modulator of oxidative stress and aging (Bains and Shaw, 1997; Venkateshappa et al., 2012; Howarth et al., 2017; Verkerke et al., 2021).

In AD, astrocytes lose their polarization, detach from the BBB, become reactive and release inflammatory cytokines (Liddel et al., 2017; Sweeney et al., 2019b; Cortes-Canteli and Iadecola, 2020). The crosstalk between astrocytes and microglia needs further understanding, although it has been extensively shown that they modulate each other's activation state (Joshi et al., 2019; McConnell et al., 2019; McAlpine et al., 2021). Astrocytes have also been reported to lose their expression of AQP4 early in AD (Smith et al., 2019). Differently, in stroke, AQP4 expression correlates with cerebral edema, increasing neuronal damage (Manley et al., 2000). In IS, astrogliosis is very significant due to the presence of DAMPs during the initial/acute phase, as well as the secondary/late phase of injury. A major indicator of gliosis is an increase in the expression of glial fibrillary acidic protein (GFAP) by astrocytes, observed in stroke as well as AD models (Pekny and Lane, 2007; Oeckl et al., 2019; Freitas-Andrade et al., 2020). Upon ischemic injury, neurovascular coupling is lost, together with astrocyte mediated CBF (McConnell et al., 2019; Freitas-Andrade et al., 2020). This results in a loop of metabolic stress and inflammation, worsening mitochondrial

dysfunction in vascular cells, and ultimately leading to BBB breakdown (Girouard and Iadecola, 2006; Freitas-Andrade et al., 2020; Radenovic et al., 2020; Sarvari et al., 2020).

Microglia

Although not directly attached to ECs, microglia mediate BBB integrity (Eldahshan et al., 2019; Freitas-Andrade et al., 2020). They are considered the resident immune cells of the CNS, able to phagocytize neurotoxic substances, and, more recently, they have also been reported to participate in a series of cerebral homeostatic functions, including synaptic plasticity and brain development (Badimon et al., 2020; Willis et al., 2020). There are two well-studied activation phenotypes of microglia, referred as M1 and M2, which are classified as being pro-inflammatory and anti-inflammatory, respectively. The M1 phenotype is detrimental to the BBB, causing dysregulation of the NVU, while the M2 facilitates the endocytosis and the clearance of toxic substances and dying cells, limiting the amount of oxidative stress and promoting a more suitable environment for regeneration/healing, following CNS injury (Tang and Le, 2016; Freitas-Andrade et al., 2020; Jiang et al., 2020). Recently, a protein highly expressed in microglia, triggering receptor expressed on myeloid cells-2 (TREM2), has been observed to protect from neurodegeneration in models of AD. TREM2 is hypothesized to be neuroprotective, associated with the M2 phenotype, although there are variants that increase the likelihood of developing AD (Ulland and Colonna, 2018; Gervois and Lambrichts, 2019; Zhou et al., 2020). The protective mechanisms of TREM2 have also been observed in IS (Gervois and Lambrichts, 2019).

Following IS or HS, microglia transition into a reactive state, causing the release of pro-inflammatory cytokines, exacerbating neuroinflammation and neurovascular dysfunction (Eldahshan et al., 2019). Microglia activation is also observed in models of AD and is now considered one of the major hallmarks of the disease (Villegas-Llerena et al., 2016). Inflammasome activation has been observed to occur in aging, neurodegeneration, as well as in stroke (Mohamed et al., 2015; Liddelow et al., 2017; Hu et al., 2019; Yang et al., 2019). In the brain, inflammasome activation occurs in many different cell types. Particularly, microglia, being the most sensitive cells to pathogen associated molecular patterns (PAMPs) and DAMPs, trigger the activation of caspase-1 and the release of cytokines IL-1 β along with IL-18 (Hu et al., 2019). The cytokine receptors on vascular cells are then activated, triggering detrimental processes which cause NVU dysfunction and disruption of neurovascular coupling (Eldahshan et al., 2019). Among other cells, microglia have also been observed to secrete metalloproteases (MMPs), such as matrix metalloproteinase-2 (MMP-2) and matrix metalloproteinase-9 (MMP-9), which have been reported to be activated in AD and IS, degrading the basement membrane, therefore increasing BBB permeability and NVU dysfunction (Lorenzl et al., 2003; Hernandez-Guillamon et al., 2010; Yang and Rosenberg, 2015; Turner and Sharp, 2016; Montaner et al., 2019; Freitas-Andrade et al., 2020; Sarvari et al., 2020; Carcel-Marquez et al., 2021).

Basement Membrane

The basement membrane is an essential part of the BBB and is composed of numerous proteins such as laminins, collagen,

nidogen, and heparin sulfate proteoglycans (Carare et al., 2020). The different proteins that make up the basement membrane are secreted by the cell types composing the NVU such as ECs, pericytes and astrocytes. These different proteins support cell-cell interactions, and thus BBB integrity (Carare et al., 2020; Freitas-Andrade et al., 2020).

Destruction of the basement membrane is severely apparent after middle cerebral artery occlusion (MCAO) (Sarvari et al., 2020). This degeneration occurs through different mechanisms of neuroinflammation such as ROS production as well as MMP secretion (Mohamed et al., 2015; Yao, 2019; Freitas-Andrade et al., 2020). Furthermore, the secretion and activation of MMPs interferes with the composition of the BBB, causing it to become leaky, and further exacerbating already existing oxidative stress and neuroinflammation (Fujimura et al., 1999; Hernandez-Guillamon et al., 2010; Montaner et al., 2019; Kang and Yao, 2020). Basement membrane composition and structure has also been observed to change in models of CAA (Morris et al., 2014).

Neurons

Blood vessel-coupled-neurons communicate with ECs and other cells of the NVU to modulate vascular structure, dilation or constriction, and to provide nutrients based on neuronal need (Andreone et al., 2015; Iadecola, 2017; Kaplan et al., 2020). Neurovascular coupling is essential to maintain the proper influx of nutrients and the proper efflux of toxic waste from the brain (Girouard and Iadecola, 2006; Andreone et al., 2015; Kaplan et al., 2020). Neuronal activity has been observed to participate in both angiogenesis and neurovascular coupling (Girouard and Iadecola, 2006; Huneau et al., 2015).

Upon CNS injury and cell death of NVU cells, dysregulation of neurovascular coupling occurs (Iadecola, 2017; Sweeney et al., 2019a; Freitas-Andrade et al., 2020; Kaplan et al., 2020). The absence of communication between the CNS and the systemic blood circulation ultimately leads to metabolic failure, oxidative stress, mitochondrial dysfunction and synaptic loss (Freitas-Andrade et al., 2020; Kaplan et al., 2020). It is therefore conceivable that, when cerebrovascular integrity and function are preserved, neurodegeneration is less severe and CNS homeostasis is better maintained.

CARBONIC ANHYDRASES

CAs are a family of zinc metalloenzymes catalyzing the reversible reaction $\text{CO}_2 + \text{H}_2\text{O} \rightleftharpoons \text{HCO}_3^- + \text{H}^+$. All four species in this chemical reaction are essential for CNS homeostasis. Therefore, it is expected that CAs influence CNS and further NVU function. The CO_2 hydration reaction can occur spontaneously (uncatalyzed), but the reaction rate is too slow for the dynamic of living cells. Consequently, CA is ubiquitously spread in all living organisms, including humans, where it accelerates this reaction millions of times, making the interconversion of CO_2 and HCO_3^- almost instantaneous. In humans, there are 15 CA isoforms, either acatalytic (CA-VIII, CA-X, and CA-XI) with exact function presently unknown, or catalytically active. The latter are found in the subcellular locations where $\text{CO}_2/\text{HCO}_3^-$ interconversion is required, from the site of production in

the mitochondria (mitochondrial CA-VA and CA-VB), moving into the cytosol (CA-I, CA-II, CA-III, CA-VII, CA-XIII), then to the plasma membrane (CA-IV, CA-IX, CA-XII, CA-XIV) and finally extracellularly (CA-VI secreted in saliva (Supuran, 2008; Zamanova et al., 2019; Mishra et al., 2020)). As genetic manipulation and selective inhibitors (Supuran et al., 1998; Ilies et al., 2003; Smaïne et al., 2007; Supuran, 2008; Güzel et al., 2009; Winum et al., 2009; Alterio et al., 2012; Akocak et al., 2016; Angeli et al., 2020) have become more available, there has been an increase in studies to determine the cell-specific function of different CA isoforms (Ghandour et al., 1992; Sly and Hu, 1995; Kida et al., 2006; Pan et al., 2006, 2012; Imtaiyaz Hassan et al., 2013; Shah et al., 2013b; Akocak and Ilies, 2014; Angeli et al., 2017; Waheed and Sly, 2017; Haapasalo et al., 2020; Mishra et al., 2020). One of the most ubiquitously expressed and catalytically active isoform is CA-II, having a turnover rate for CO₂ hydration approaching diffusion limit ($K_{cat} = 1.4 \times 10^6 \text{ s}^{-1}$). It is a cytosolic enzyme and has the widest distribution in the human body, being expressed in cells from virtually every tissue or organ. In the brain, it is found in large amounts in oligodendrocytes and epithelium of the choroid plexus. Subjects suffering from CA-II deficiency syndrome, a human autosomal recessive disorder, display osteopetrosis, bone fragility, renal tubular acidosis, and, importantly, cerebral calcification and cognitive defects, developmental delay and usually a short stature (Sly and Hu, 1995; Supuran, 2008; Zamanova et al., 2019; Mishra et al., 2020). This isoform has been observed to translocate to the mitochondria, upon aging and neurodegeneration in a Purkinje cell degeneration mouse model (Pollard et al., 2016). This study also showed that *C. elegans* exposed to CA-II have a shorter lifespan, suggesting that high CA-II levels are involved in cell life/cycle-limiting mechanisms. Future studies need to further elucidate why and how this occurs.

CA-I isoform was first identified in red blood cells, where is five to six times more abundant than CA-II, although it has only about 15% of the CA-II activity (Supuran, 2008; Zamanova et al., 2019; Mishra et al., 2020). CA-I is the most abundant non-hemoglobin protein in erythrocytes and, together with CA-II, contributes to equilibration of dissolved CO₂/HCO₃⁻ pools in blood and maintains the pH blood homeostasis, also facilitating the CO₂ transport from brain and metabolizing tissues to lungs (Sly and Hu, 1995; Supuran, 2008; Zamanova et al., 2019; Mishra et al., 2020). CA-I is hypothesized to contribute to cerebral edema due to the observation that CA-I is increased in the brain following HS (Guo et al., 2012). One of the most important cytosolic isozymes in the brain is CA-VII, a fast isozyme ($K_{cat} = 9.5 \times 10^5 \text{ s}^{-1}$), found especially in the neurons of hippocampus, together with CA-II. Interestingly, CA-VII is not found in glial cells, which contain just CA-II in the cytosol. Besides CNS, CA-VII is found in skeletal muscles, stomach, duodenum, liver, colon (Sly and Hu, 1995; Supuran, 2008; Zamanova et al., 2019; Mishra et al., 2020). In the brain, Kaila's group has shown that CA-VII acts as a molecular switch in hippocampal CA1 pyramidal neurons in the development of synchronous gamma-frequency firing in response to high-frequency stimulation. This

finding makes CA-VII an important modulator of long-term potentiation, synaptic plasticity, memory, and learning processes (Ruusuvuori et al., 2004). The same group, using a novel CA-VII (*Car7*) KO mouse model, as well as a CA-II (*Car2*) KO, and a CA-II/VII double KO mouse models, has shown that in mature hippocampal pyramidal neurons CA-VII and CA-II isozymes enhance bicarbonate-driven GABAergic excitation during intense GABA_A-receptor activation. The expression of these two cytosolic isozymes was detected at a very early age in the animals (10- and 20-days post-birth), pointing toward CA-VII and CA-II being key molecules in age-dependent neuronal pH regulation (Supuran, 2008; Ruusuvuori et al., 2013; Ruusuvuori and Kaila, 2014; Zamanova et al., 2019; Mishra et al., 2020).

Membrane-bound isozymes also play an important role in the brain, especially for the regulation of extracellular pH (Chesler, 2003; Shah et al., 2005). Thus, the phosphatidylinositol glycan (GPI)-anchored isozymes CA-IV (Stams et al., 1996) is a fast isozyme ($K_{cat} = 1.1 \times 10^6 \text{ s}^{-1}$) found on the plasma face of the cortical capillaries. It is more resistant to inhibition by halide ions than CA-II, being adapted to perform the CO₂/HCO₃⁻ interconversion in the extracellular space that contains a higher concentration of Cl⁻ ions than the cytosol. The isozyme is also expressed in the choriocapillaries of the eye, in skeletal and cardiac muscles, lungs, kidneys, gastrointestinal and reproductive tracts (Sly and Hu, 1995; Supuran, 2008; Zamanova et al., 2019; Mishra et al., 2020). Another isoform localized within the plasma membrane is CA-IX, a transmembrane isozyme, possessing an N-terminal proteoglycan domain, the catalytic domain, a single-pass transmembrane region, and an intracellular tail. It is a dimeric protein with a low expression pattern in most organs, except for the digestive system (largest amount) and CNS, where it can be found mainly in the ventricular-lining cells and in the choroid plexus (Saarnio et al., 1998; Hilvo et al., 2008; Supuran, 2008; Alterio et al., 2009; Pastorek and Pastorekova, 2015; Zamanova et al., 2019; Mishra et al., 2020). The expression of CA-IX is up-regulated in hypoxia by the transcription factor hypoxia inducible factor-1 (HIF1 α) and is associated with the Warburg effect in cancer pathology (Ivanov et al., 2001; Supuran, 2008; Pastorek and Pastorekova, 2015; Shabana and Ilies, 2019). In models of glioblastoma, CA-IX has been observed to increase cell migration, motility, and adhesion in monocytes (Huang et al., 2020). Currently, a compound partially selective to inhibit for CA-IX, referred to as SLC-0111, is in clinical trials, and it is likely it would operate as a therapy for multiple cancers, including glioblastoma (Boyd et al., 2017; Ilies and Winum, 2019; Shabana and Ilies, 2019; McDonald et al., 2020). This isoform has also been suggested to be involved in heart fibrosis in a rat cardiac ligation model (Vargas et al., 2016). Interestingly, in humans, CA-IX is expressed also within atherosclerotic plaques, where it is suggested to be a marker of necrotic tissue (Demandt et al., 2021). Recent studies provided evidence that CA-IX likely exacerbates cerebral ischemia outcomes (Mishra et al., 2020). CA-XII is another medium-fast ($K_{cat} = 4.2 \times 10^5 \text{ s}^{-1}$), membrane-bound isozyme, similar in general structure with CA-IX, but without the proteoglycan domain. It is also dimeric, with the two active sites oriented toward the extracellular milieu

(Ivanov et al., 1998; Türeci et al., 1998; Whittington et al., 2001). It is found in many hypoxic tumors, alongside CA-IX, including gliomas, hemangioblastomas, and meningiomas. However, opposite to CA-IX, CA-XII is highly expressed in many normal tissues including breast epithelium and non-pigmented ciliary epithelial cells of the eye, in brain (small peripheral capillaries), esophagus, pancreas, colon, rectum, kidney, prostate, ovary, testis, endometrium, sweat glands. CA-XII was shown to play a key role in epithelial cell electrolyte homeostasis via activation of the ductal $\text{Cl}^-/\text{HCO}_3^-$ exchanger AE2, and association with $\text{Na}^+/\text{HCO}_3^-$ cotransporter kNBC1 (Proescholdt et al., 2005; Purkerson and Schwartz, 2007; Supuran, 2008; Hong et al., 2015; Waheed and Sly, 2017; Zamanova et al., 2019; Mishra et al., 2020). The membrane-bound isozyme CA-XIV possesses an extracellular catalytic domain, a single transmembrane helix, a short intracellular polypeptide segment, and has a moderate catalytic activity. It is highly expressed in the kidney, retina and the heart, as well as in brain, skeletal muscles, liver, and lungs. It was reported that CA-XIV is interacting with bicarbonate transporters and is involved in acid–base balance in muscles and erythrocytes in response to chronic hypoxia, and hyperactivity of the heart (Mori et al., 1999; Supuran, 2008; Mboge et al., 2018; Zamanova et al., 2019; Mishra et al., 2020).

The mitochondrial isoforms CA-VA and CA-VB are isozymes with medium-high activity ($K_{catVA} = 2.9 \times 10^5 \text{ s}^{-1}$, $K_{catVB} = 9.5 \times 10^5 \text{ s}^{-1}$), important for gluconeogenesis, lipogenesis, ureagenesis, and other anabolic pathways. They supply HCO_3^- to pyruvate carboxylase during gluconeogenesis and lipogenesis pathways, and to carbamoyl phosphate synthetase in ureagenesis pathway. CA-VA is found mainly in the liver, while CA-VB is found in skeletal and heart muscles, kidneys, pancreas, gastrointestinal tract, brain and spinal cord (Shah et al., 2000; Supuran, 2008; Scozzafava et al., 2013; Zamanova et al., 2019; Mishra et al., 2020). Due to their involvement in these anabolic pathways, they have been studied in models of obesity and type 2 diabetes (De Simone and Supuran, 2007; Supuran, 2012; Scozzafava et al., 2013; Salameh et al., 2019; Mishra et al., 2020). The function of CA-V and the difference between CA-VA and CA-VB isoforms has been examined by analyzing the phenotypical differences between CA-VA and CA-VB KO, as well as the double KO mouse models, indicating their role in ammonia detoxification (Shah et al., 2013b). The function of CA-V in the brain is extremely understudied, however, CA-VA has been reported to be expressed in both neurons and glial cells (Ghandour et al., 2000). Interestingly, the effect of CA-V has been also investigated in cerebral pericytes (Price et al., 2017). It has been observed that silencing of both isoforms protect against high-glucose-induced cell death and ROS production, CA-VA to a more significant degree (Price et al., 2017), confirmed also by increased oxidative stress and apoptosis in models of CA-VA over expression (Price et al., 2017).

Although extensive studies reported the multiple functions of the different CA isoforms in a variety of tissues, further investigation regarding CAs cell-specific expression/activity, especially within the CNS, and in CNS disorders, is needed. In particular, CAs impact on cerebrovascular dysregulation occurring during IS and AD must still be elucidated. It is

therefore crucial to identify the cell- and isoform-specific roles, and to design isoform-selective inhibitors, which may be likely to ameliorate the cell/tissue specificity and to decrease the observed side effects of the FDA-approved pan-CAIs (Provensi et al., 2019; Zamanova et al., 2019; Mishra et al., 2020).

Below, we summarize the available literature highlighting the positive effects of CA inhibition on neurovascular dysfunction in stroke-, AD-, CAA-, and diabetes-induced cerebrovascular pathology.

Carbonic Anhydrase Inhibitors and Cerebrovascular Pathology

CAIs have been studied for decades (Maren, 1967; Bertini and Luchinat, 1983; Silverman and Lindskog, 1988; Supuran et al., 2003; Krishnamurthy et al., 2008; Supuran, 2008; Winum et al., 2009; Alterio et al., 2012; Akocak and Ilies, 2014; McKenna and Supuran, 2014; Supuran and Winum, 2015; Ilies and Winum, 2019; Angeli et al., 2020; Supuran and Capasso, 2020). They were first developed as diuretics due to their function on the reabsorption of sodium and water within the kidney (DuBose, 1984; Purkerson and Schwartz, 2007; Supuran, 2018), and are currently used to treat glaucoma as they decrease intraocular pressure, and for the prevention of high altitude sickness (Bradwell et al., 1992; Mincione et al., 2007; Ritchie et al., 2012), as they diminish pulmonary vasoconstriction, increase CBF likely controlling cerebral oxygenation, and reduce cerebral edema (Mishra et al., 2020). However, the molecular mechanism responsible for the effects of CAIs, including some of the most used pan-CAIs such as ATZ and MTZ, are multiple, and still under investigation.

MTZ, for example, was recognized as one of a few compounds that had the ability to inhibit cytochrome c release from the mitochondria under oxidative stress (Wang et al., 2008). Over 1000 compounds of the NINDS drug library were first screened in isolated mitochondria from mouse liver and further confirmed in striatal cells in models of Huntington's disease. MTZ, inhibiting cytochrome c release from challenged mitochondria, also resulted in the reduction of caspase-9 and caspase-3 activation (Wang et al., 2008; Fossati et al., 2016; Sekerdag et al., 2018). Following this study, FDA-approved CAIs such as MTZ, ATZ, topiramate and more recently developed non-FDA approved selective inhibitors, are starting to be applied in models of cerebrovascular pathology, ischemia, CAA and AD, and their positive effects have been attributed, at least in part, to their ability to prevent mitochondrial dysfunction in cerebrovascular cells (Wang et al., 2009; Shah et al., 2013a; Fossati et al., 2016; Solesio et al., 2018).

CA inhibition has been shown to contribute to cerebrovascular tone during transient phases of pH change, but not in steady-state conditions, in rat arteries (Rasmussen and Boedtker, 2018). This study used two different CAIs, ATZ and 4-aminomethylbenzenesulfonamide (AMB). It was revealed that only the pan-CAI ATZ, potent against most of intracellular CA isozymes, had an effect on intracellular acidification mechanisms. Based on the observed results, it was concluded that intracellular CAs are responsible for modifying the rate of intracellular pH and vascular tone within the arteries (Rasmussen and Boedtker, 2018). Other studies have observed the vasodilator

abilities of ATZ, such as increased NO production leading to increased blood flow in the cortex of rats (Tuettenberg et al., 2001). It is likely that the production of NO is not the only mechanism by which ATZ mediates vasodilation (Kiss et al., 1999). Systemic administration of ATZ has also been widely used as a short test to increase CBF in human studies (Okudaira et al., 1995; Grossmann and Koeberle, 2000; Russell et al., 2008). Interestingly, its stimulatory effect on CBF is reduced in patients with AD or vascular dementia compared to healthy controls (Stoppe et al., 1995; Pavics et al., 1999).

Carbonic Anhydrase Inhibition in Models of Ischemic Stroke

The neuroprotective effects of MTZ, in both *in vitro* and *in vivo* models of ischemic injury, displayed in Table 1, were first explored by Wang et al. (2009). Exposing primary cortical neurons to OGD and H₂O₂ resulted in necrosis, which

was rescued by MTZ treatment. Similarly, OGD increased cytochrome c release and apoptosis inducing factor (AIF) release from the mitochondria, as well as the activity of caspase-3 (Wang et al., 2009), and MTZ attenuated these effects (Wang et al., 2009), suggesting that MTZ does not only prevent necrosis, but also mitochondria-mediated apoptosis, induced by OGD. In the same study, primary cortical neurons exposed to OGD presented inflammasome activation, measured by the activation of caspase-1 and release of IL-1 β . MTZ inhibited both IL-1 β release as well as caspase-1 activation *in vitro* (Wang et al., 2009). This finding supports the hypothesis that CAs may mediate both apoptotic and inflammatory mechanisms. In a mouse model of MCAO, MTZ-treated mice (20 mg/kg) had a smaller infarct size, improved neurological score, and decreased cytochrome c release and caspase-3 activation, compared to non-treated mice (Wang et al., 2009). A more recent study performed in rats determined the effectiveness of ATZ alone, as well as in conjugation with head-down tilt, a way to physically promote CBF, in a transient MCAO rat model. The results indicated that ATZ was protective following transient MCAO, significantly decreasing infarct size (Han et al., 2020). Furthermore, ATZ reduced AQP4 expression, compared to rats with no treatment, suggesting a reduction in brain edema (Han et al., 2020). In a rat model of permanent MCAO, ATZ was used in comparison with selective inhibitors for CA-VII, CA-IX, and CA-XII to determine whether these

TABLE 1 | CA inhibition in models of IS.

Model	Mechanism	CA Inhibitor/Isoform	References
Mouse primary cortical neurons	Inhibition of OGD induced necrosis Inhibition of OGD induced mitochondria mediated apoptosis Inhibition of OGD induced inflammasome activation	MTZ	Wang et al., 2009
C57BL/6J Mouse pMCAO	Reduction of infarct size Improvement of neurological score Reduction of caspase 3 activation Decrease of cytochrome C release	MTZ	Wang et al., 2009
Wistar rat tMCAO	Decrease of infarct size Reduction of AQP4 expression Reduction of brain water content and sodium accumulation	ATZ	Han et al., 2020
Sprague dawley rat pMCAO	Improvement of neurological score Reduction of infarct size	CA-VII inhibition, CA-IX/XII inhibition ATZ, CA-IX/XII inhibition	Di Cesare Mannelli et al., 2016
Rat hippocampal slices	Inhibition of OGD induced anoxic depolarization	ATZ, CA-IX inhibition, CA-XII inhibition	Dettori et al., 2021
Wistar rats pMCAO	Reduction of infarct size Improvement of neurological score Attenuation of microglia activation	ATZ, CA-IX/-XII inhibition	Dettori et al., 2021

MTZ, Methazolamide; ATZ, Acetazolamide; HS, hemorrhagic stroke; p/Tmcao, permanent/transient middle cerebral artery occlusion; CA, carbonic anhydrase; AQP4, Aquaporin-4; OGD, oxygen glucose deprivation.

TABLE 2 | CA inhibition in models of HS.

Model	Mechanism	CA Inhibitor/Isoform	References
Sprague-dawley rats intracaudate blood injection	Improvement of neurological outcome Reduction of neuronal death Exacerbation of brain water content Increase neurodegeneration	ATZ CA-I injection	Guo et al., 2012 Guo et al., 2012
Mouse primary cortical neurons	Inhibition of blood/hemoglobin induced cell death Inhibition of blood/hemoglobin induced ROS production	MTZ	Li et al., 2016
C57BL/6J Mice SAH	Reduction in caspase-3 activation and cell death in hippocampus/cortex Improvement in neurological outcome	MTZ	Li et al., 2016
New Zealand White Rabbits SAH	Reduction of neurodegeneration/apoptosis in hippocampus	Topiramate	Seçkin et al., 2009

ATZ, acetazolamide; MTZ, methazolamide; CA-I, carbonic anhydrase-1; SAH, subarachnoid hemorrhage; ROS, reactive oxygen species.

isoforms are involved in the pathology. Interestingly, the pan-CAI ATZ did not improve neurological score 24 h following occlusion, however, the CAI with selectivity for CA-VII, as well as the compound more selective for membrane isoforms CA-IX and CA-XII, improved the neurological deficits observed in the occluded untreated rats. In the group treated with the CA-IX/CA XII potent and medium selective inhibitor, 6-(benzyloxy)benzo[d]thiazole-2-sulfonamide (BBT), infarct size was significantly reduced, while no reduction was observed

in the CA-VII inhibitor group. However, rats treated with 50 mg/kg of ATZ also displayed diminished infarct volume. This study suggests that CA-IX and possibly CA-XII may exacerbate neuronal loss along with neurological and vascular deficits, following ischemic insult (Di Cesare Mannelli et al., 2016). Very recently, another study evaluated different CA-selective inhibitors, in both *in vitro* and *in vivo* models of IS (Dettori et al., 2021). This study analyzed a new generation of CAIs, which are lipophilic and selective for the hypoxia-associated (CA-IX and CA-XII) and cytosolic (CA-II and CA-I) isoforms. Importantly, ATZ was used as a reference compound at a much lower dose in this study (4.4 mg/kg). *In vitro*, the lipophilic CAIs and ATZ protected against OGD-induced anoxic depolarization, providing mechanistic information. *In vivo*, the CAIs reduced infarct volume, neurological deficits, neuronal damage and microglial activation (Dettori et al., 2021). This study also measured TNF- α and IL-10 plasma levels, showing differences between the sham surgery and the occluded groups, without any affect in the treated groups, likely due to the short time window of treatment (24-h). As proposed in the discussion, it is possible that, with a longer treatment, CA inhibition may mediate inflammatory pathways (Bulli et al., 2021; Dettori et al., 2021). Indeed, CAs have been also associated with inflammatory pathologies, such as rheumatoid arthritis and cancer metastasis (Supuran, 2008; Alver et al., 2011; Liu et al., 2012). Overall, both MTZ and ATZ are protective in models of IS, however, the development of CA-IX and potentially CA-XII inhibitors may be beneficial for the treatment of both neurological and vascular disorders, following IS. The protective mechanisms observed following CA inhibition in models of IS, summarized in Table 2, seem to be superior with compounds selective for membrane-bound isoforms CA-IX and CA-XII (Bulli et al., 2021).

Carbonic Anhydrase Inhibition in Models of Hemorrhagic Stroke

The protective mechanisms of CA inhibition in models of HS are summarized in Table 2. Inhibition of CA reduced brain injury after ICH in Sprague-Dawley rats (Guo et al., 2012). This study focused on CA-I, which is highly expressed in red blood cells. Upon intracaudate injection of blood, CA-I levels were increased in the ipsilateral basal ganglia, for as long as 3 days post-injection. Moreover, following intracaudate injection of CA-I, increased brain water content, microglial activation and neuronal cell death were detected, suggesting that CA-I expression may contribute to cerebral edema and neuroinflammation. ATZ-treated rats exhibited reduced perihematoma edema, as well as sodium accumulation, together with decreased neuronal deficits (Guo et al., 2012). However, the cellular mechanisms responsible for the effects of CA-I on BBB integrity require further investigation. Lately, the scientific community is beginning to hypothesize that CA-I, in both the retina and brain, contributes to vascular permeability, suggesting that specific inhibitors of CA-I could be beneficial to minimize NVU dysfunction, and thus neuroinflammation around the barrier (Gao et al., 2007). Furtherly, in a SAH mouse model, mice treated with the pan-CAI, MTZ, had decreased caspase-3 activation and

TABLE 3 | CA inhibition in models of AD and CAA.

Model	Mechanism	CA Inhibitor/Isoform	References
hCMEC/D3	Inhibition of A β induced DNA fragmentation Inhibition A β induced of cytochrome c release Inhibition of A β induced H ₂ O ₂ production Inhibition of A β induced caspase-9 activity	MTZ, ATZ	Fossati et al., 2010; Solesio et al., 2018
Human primary brain SMC	Inhibition of A β induced DNA fragmentation Inhibition A β induced of cytochrome c release	MTZ	Fossati et al., 2010
Normal Human Astrocytes	Inhibition of A β induced DNA fragmentation	MTZ	Fossati et al., 2016
Human Glioma M059K	Inhibition of A β induced DNA fragmentation Inhibition A β induced of cytochrome c release Inhibition of A β induced H ₂ O ₂ production Inhibition of A β induced caspase-9 activity	MTZ	Fossati et al., 2016
Human Neuroblastoma (SHSY5Y)	Inhibition of A β induced DNA fragmentation Inhibition A β induced of cytochrome c release Inhibition of A β induced H ₂ O ₂ production Inhibition of A β induced caspase-9 activity Activation of Nrf2	MTZ, ATZ MTZ	Fossati et al., 2016; Solesio et al., 2018 Sotolongo et al., 2020
Rat primary cortical neurons	Activation of Nrf2	MTZ	Sotolongo et al., 2020
C57BL/6 Mice A β hippocampal injection	Inhibition of caspase-3 activation Increase of NeuN expression in hippocampus Reduction of caspase-3 activation in reactive microglia	MTZ	Fossati et al., 2016

hCMEC/D3, human cerebral microvasculature endothelial cells; A β , amyloid-beta; MTZ, methazolamide; ATZ, acetazolamide; SMC, smooth muscle cell; SHSY5Y, human neuroblastoma cells; M059K, glioblastoma cell line; Nrf2, nuclear factor erythroid 2-related factor.

apoptosis in the hippocampus and cortex, compared to the non-treated group, ameliorating neurological deficits (Li et al., 2016). In primary cortical neurons, MTZ reduced hemoglobin and blood induced cell death along with production of ROS (Li et al., 2016). Topiramate, another CAI active on multiple CA enzymes, including CA-V, reduced SAH injury in rabbits (Seçkin et al., 2009).

Carbonic Anhydrase Inhibition in Models of Alzheimer's Disease and Cerebral Amyloid Angiopathy

Recently, the effects of CA inhibition on mitochondrial and neurovascular cell function have been tested in models of amyloidosis (Table 3). In AD and CAA, A β toxic aggregates accumulate around and within the cerebral vessel walls, as well as in parenchymal plaques in specific brain regions, such as the hippocampus and cortex (Ghiso et al., 2014; Cortes-Canteli and Iadecola, 2020). Our lab has shown that A β oligomers and protofibrils induce cytochrome c release from the mitochondria in multiple cell types of the NVU, including ECs and SMCs, neurons and glial cells, leading to caspase activation and cell death (Fossati et al., 2010, 2012a,b, 2016; Parodi-Rullán et al., 2020). In all of these neurovascular cell types, MTZ reduced mitochondrial mediated cell death triggered by A β (Fossati et al., 2010, 2016). Moreover, MTZ attenuated caspase-3 and caspase-9 activation in ECs, neuronal and glial cells *in vitro*, as

well as caspase-3 activation *in vivo* (Fossati et al., 2016; Solesio et al., 2018). In an *in vivo* study, A β was injected into the hippocampus of wild-type mice, in the presence or absence of a previous intraperitoneal MTZ injection. Interestingly, MTZ treatment attenuated the activation of caspase-3 in microglia and increased the amount of NeuN positive neurons within the hippocampus, indicating that MTZ treatment had a protective effect against neurodegeneration following A β -injection (Fossati et al., 2016). To elucidate the specific mechanisms responsible for these protective effects, our group showed that A β -induced mitochondrial dysfunction, mitochondrial membrane depolarization, cytochrome C release, and H₂O₂ production were attenuated in neuronal and ECs, not only by MTZ, but also by ATZ, which was effective at lower concentrations (Solesio et al., 2018). Moreover, both ATZ and MTZ inhibited caspase-9 activation caused by A β in cerebrovascular EC, and the resulting apoptosis (Solesio et al., 2018). MTZ has been also observed to increase the activation of nuclear factor-related factor 2 (Nrf2), in models of high-altitude sickness and more interestingly, in human neuroblastoma cells and primary cortical neurons challenged with A β *in vitro* (Lu et al., 2020; Sotolongo et al., 2020). Nrf2-activation by MTZ increased the activity of antioxidant enzymes, such as superoxide dismutase-1 and heme-oxygenase-1, pointing to the potential downstream effects of MTZ (Sotolongo et al., 2020). These studies support the hypothesis that CA inhibition is protective to multiple cell types of the NVU, in models of amyloidosis, and highlight the necessity to test

TABLE 4 | CA isoforms in CNS and cerebrovascular pathology.

CA isoform	Cellular localization	Cell type/Brain area	Cellular function	Neurological disorder	References
CA-I	Cytosol	Red blood cells	-pH homeostasis in the blood -Edema/sodium accumulation in the brain	IH	Guo et al., 2012
CA-II	Cytosol	-Epithelium of choroid plexus -Glial Cells -Neurons	-Intracellular ion homeostasis -Cell-life/life cycle	Loss of expression associated with cognitive abnormalities	Sly and Hu, 1995; Mishra et al., 2020
CA-III	Cytosol	MCA	N/A	N/A	Rasmussen and Boedtkjer, 2018
CA-IV	Plasma membrane	-Cortical capillaries -MCA	Extracellular pH	N/A	Chesler, 2003; Shah et al., 2005; Rasmussen and Boedtkjer, 2018
CA-VA	Mitochondria	-Cerebrovascular pericytes -Neurons -Glial cells	-High-glucose induced Apoptosis -High-glucose induced ROS production -Cell viability -Biogenesis reactions	Type-2 Diabetes induced cerebrovascular pathology	Ghandour et al., 2000; Patrick et al., 2015; Price et al., 2017
CA-VB	Mitochondria	-Cerebrovascular pericytes -MCA -CNS cells	Cell Viability	Type-2 Diabetes induced cerebrovascular pathology	Price et al., 2017; Rasmussen and Boedtkjer, 2018; Mishra et al., 2020
CA-VII	Cytosol	Hippocampal neurons	-Gamma-frequency firing -Long-term potentiation	Epilepsy	Ruusuvuori et al., 2004, 2013
CA-IX	Plasma membrane	-Glioblastoma -Choroid plexus -Ventricular linings	-Extracellular pH -Monocyte adhesion/cell migration -Tumor progression	Glioblastoma-IS	Di Cesare Mannelli et al., 2016; Huang et al., 2020; Mishra et al., 2020
CA-XII	Plasma membrane	-Glioblastoma -Peripheral capillaries -MCA	Extracellular pH	Glioblastoma	Proescholdt et al., 2005; Rasmussen and Boedtkjer, 2018

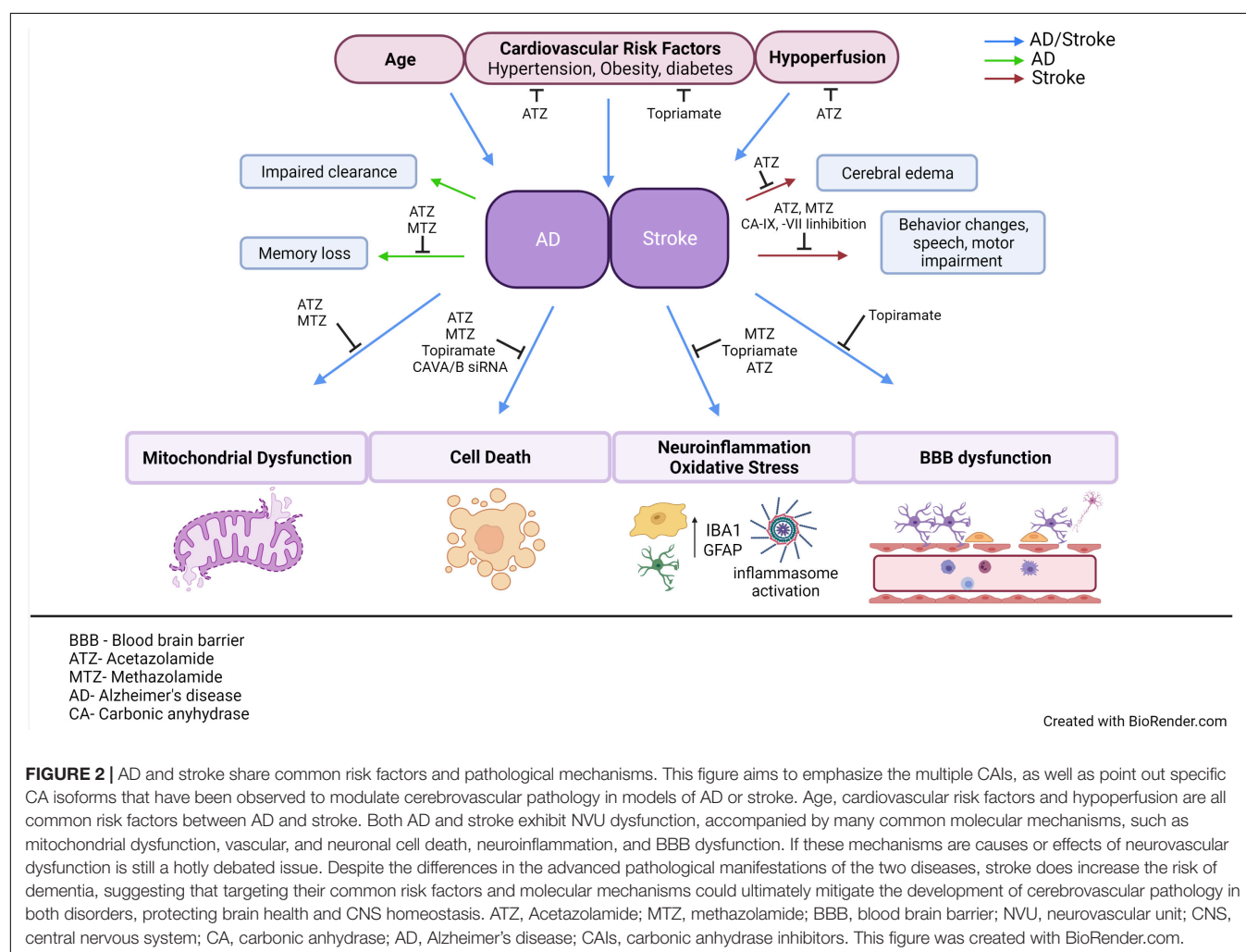
CA, carbonic anhydrase; MCA, middle cerebral artery; IH, intracerebral hemorrhage; IS, ischemic stroke.

these and/or isoform-specific compounds in clinical trials for AD and CAA. New studies currently performed in our lab are also confirming the positive effects of CAIs on cognitive performance in mouse models of amyloidosis (Angiulli et al., 2018). Additional studies are in process to investigate chronic treatments with pan-CAIs in multiple models of CAA and AD, in parallel with the assessment of specific isoform inhibitors, to further elucidate the role of different CA isoforms in AD and CAA pathology.

Interestingly, although CAIs are proving to be beneficial in preventing CAA and AD pathology, CA activators have also showed beneficial effects for memory retention in acute models (Ilies et al., 2002; Canto de Souza et al., 2017; Sanku et al., 2018; Provensi et al., 2019; Blandina et al., 2020; Schmidt et al., 2020). These findings point to a relevant role of CA enzymes in the modulation of multiple pathways in contextual fear memory extinction, neurodegenerative and neurovascular pathology. They also highlight the need for a careful and specific modulation of CA enzyme's activity, and for a further understanding of the effects of each isoform in different brain areas, cell types, and pathological conditions.

Carbonic Anhydrase Inhibition in Models of Diabetes-Induced Cerebrovascular Pathology

Some studies have also explored the specific inhibition of the mitochondrial isoforms CA-VA and CA-VB, which recently gained interest for their role in the prevention of cerebrovascular pathology, in different models of diabetes (Price et al., 2012, 2017; Shah et al., 2013a), which contributes to AD and stroke pathogenesis (Kuzma et al., 2018; Cortes-Canteli and Iadecola, 2020; Freitas-Andrade et al., 2020; Zlokovic et al., 2020). In a streptozotocin induced diabetic CD-1 mouse model it was reported that topiramate treatment for 3 weeks increased the levels of glutathione and reduced oxidative stress in the brain. It was also observed that diabetes-induced loss of cerebral pericytes was attenuated by topiramate (Price et al., 2012). The same group confirmed these mechanisms in the brains of CA-V double KO mice, where increased levels of glutathione and decreased oxidative stress was observed compared to wild-type (Price et al., 2012). Hyperglycemia-treated cerebrovascular pericytes increased ROS production, as well as the rate of



mitochondrial respiration. The inhibition of CA (particularly CA-V), with topiramate, reduced ROS production (Shah et al., 2013a). Although topiramate has a high inhibitory activity on the CA-V isoforms, it also has affinity for the other isoforms, as well as multiple off-target effects in the brain, as shown by studies that analyzed topiramate's functions on GABA and glutamate receptors (Johnson, 2005; Mao et al., 2015). Ideally, more specific inhibitors should be developed to further investigate the functions of CA-VA/B. The same group used a plasmid to overexpress CA-VA in cerebral pericytes exposed to high-glucose to confirm the hypothesized molecular mechanisms involving CA-VA, such as ROS production and cell death, observing that topiramate was protective against these mechanisms (Patrick et al., 2015). As a proof of concept, the same group genetically knocked down CA-VA and CA-VB in cerebral pericytes (Price et al., 2017). CA-VA and CA-VB knockdown improved cell viability, compared to the control, when exposed to hyperglycemia, although the specific knockdown of CA-VA was even more effective than CA-VB knockdown in decreasing ROS production and apoptosis, in a hyperglycemic environment (Price et al., 2017). These studies reveal potential differences between CA-VA and CA-VB function in brain pericytes, which need further investigation. In a streptozotocin induced diabetic mouse model, cerebrovascular pathology was also characterized using electron microscopy. The breakdown of the BBB and its dysfunction was attenuated by topiramate treatment *in vivo* (Salameh et al., 2016). More recently, a study using a high-fat-induced diabetes mouse model, focused on the disruption of the hippocampal BBB. BBB tight junction proteins, such as ZO-1 and claudin-12, were reduced with a high-fat diet, while topiramate treatment increased their expression, along with the attenuation of oxidative stress (Salameh et al., 2019).

CONCLUSION

The NVU is an important functional structure of the CNS, and its failure participates in the development of AD, vascular dementias, and exacerbates stroke outcomes. Multiple pathological and cellular mechanisms leading to NVU pathology need clarification, and new therapeutic strategies should be further investigated and developed. As scientific techniques improve, the ability to understand the functions of this unit increases. Recent studies support that CA inhibition is protective to the NVU, and the role of these enzymes should be further investigated. **Table 4** emphasizes the CA isoforms that have been discussed throughout the review, their cellular localizations, their association with specific neurological disorders, and their

expression in neurovascular cells. Clarifying how the cells of the NVU interact with each other, as well as the roles of CAs within each cell type, is critical for targeting cerebrovascular pathology in IS, AD, as well as other neurodegenerative diseases. The pan-CAIs MTZ and ATZ, as well as topiramate, have shown protective effects in models of stroke, cerebrovascular pathology, type II diabetes, and AD, summarized in **Figure 2**. Although these compounds are FDA-approved, facilitating translation to clinical trials, these CAIs are not specific drugs. Isoform-specific CAIs are of increasing interest, as 15 isoforms with different functions and localizations have been identified in humans. Further preclinical and clinical studies to assess the efficacy of CA inhibition in AD as well as IS are essential. To confirm the role of specific CA isoforms as pharmacological targets, genetic studies, such as isoform KO and knock in in different NVU cell types, would be beneficial. Overall, the studies discussed above provide evidence for CAs as important potential mediators and targets in neurovascular pathology for AD, stroke and related cerebrovascular disorders.

AUTHOR CONTRIBUTIONS

NL and SF designed and conceptualized the review. NL wrote the review draft and did the literature search. SF critically revised, edited the manuscript, provided relevant insights, additional literature search, and acquired funding. EC and MI revised the manuscript and provided additional literature search. All authors contributed to the article and approved the submitted version.

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Insulin-Like Growth Factor-1 Differentially Modulates Glutamate-Induced Toxicity and Stress in Cells of the Neurogliovascular Unit

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The age-related reduction in circulating levels of insulin-like growth factor-1 (IGF-1) is associated with increased risk of stroke and neurodegenerative diseases in advanced age. Numerous reports highlight behavioral and physiological deficits in blood-brain barrier function and neurovascular communication when IGF-1 levels are low. Administration of exogenous IGF-1 reduces the extent of tissue damage and sensorimotor deficits in animal models of ischemic stroke, highlighting the critical role of IGF-1 as a regulator of neurovascular health. The beneficial effects of IGF-1 in the nervous system are often attributed to direct actions on neurons; however, glial cells and the cerebrovasculature are also modulated by IGF-1, and systemic reductions in circulating IGF-1 likely influence the viability and function of the entire neuro-glio-vascular unit. We recently observed that reduced IGF-1 led to impaired glutamate handling in astrocytes. Considering glutamate excitotoxicity is one of the main drivers of neurodegeneration following ischemic stroke, the age-related loss of IGF-1 may also compromise neural function indirectly by altering the function of supporting glia and vasculature. In this study, we assess and compare the effects of IGF-1 signaling on glutamate-induced toxicity and reactive oxygen species (ROS)-produced oxidative stress in primary neuron, astrocyte, and brain microvascular endothelial cell cultures. Our findings verify that neurons are highly susceptible to excitotoxicity, in comparison to astrocytes or endothelial cells, and that a prolonged reduction in IGFR activation increases the extent of toxicity. Moreover, prolonged IGFR inhibition increased the susceptibility of astrocytes to glutamate-induced toxicity and lessened their ability to protect neurons from excitotoxicity. Thus, IGF-1 promotes neuronal survival by acting directly on neurons and indirectly on astrocytes. Despite increased resistance to excitotoxic death, both astrocytes and cerebrovascular endothelial cells exhibit acute increases in glutamate-induced ROS production and mitochondrial dysfunction when IGFR is inhibited at the time of glutamate stimulation. Together these data highlight

that each cell type within the neuro-glio-vascular unit differentially responds to stress when IGF-1 signaling was impaired. Therefore, the reductions in circulating IGF-1 observed in advanced age are likely detrimental to the health and function of the entire neuro-glio-vascular unit.

Keywords: astrocytes, endothelial cells, neurotoxicity, somatomedin C, reactive oxygen species

INTRODUCTION

Aging is a primary risk factor for a number of diseases and pathologies, including stroke, which is recognized as one of the leading causes of death and disability globally (Johnson et al., 2016; Benjamin et al., 2019). Ischemic stroke is particularly devastating as the long-term phenotypic deficits in cognitive behavior and sensorimotor function build in the days following the insult, even if blood flow to the ischemic tissue is restored [as reviewed by George and Steinberg (2015)]. Ischemia precipitates cellular damage in the brain within minutes, as the loss of blood flow induces metabolic and oxidative stress in the energy-demanding neurons, and nearby glial and cerebrovascular cells. Immediate restoration of blood flow is critical, as the length of ischemia correlates with the extent of tissue damage (Terasaki et al., 2014). To date, the most effective ischemic stroke therapy is fibrinolytic breakdown of the clot with administration of tissue plasminogen activator. Unfortunately, this therapy is time-restricted and requires confirmation of ischemic vs hemorrhagic manifestation, which restricts clinical utility and has limited its use to around 5% of stroke patients each year (National Institute of Neurological Disorders and Stroke rt-PA Stroke Study Group, 1995). Longitudinal observation-based clinical assessments have identified multiple possible biomarkers of stroke risk and severity, including insulin-like growth factor-1 (IGF-1), which may serve as a point of therapeutic intervention in the days surrounding insult (Katzan et al., 2000; California Acute Stroke Pilot Registry Investigators, 2005; Qureshi et al., 2005; Kleindorfer et al., 2008).

Insulin-like growth factor-1 is a pleiotropic neuroendocrine modulator that has been assessed in both clinical and preclinical studies of stroke [as reviewed by Hayes et al. (2021)]. Both humans and rodents exhibit age-related circulating declines of IGF-1 (O'Connor et al., 1998; Sonntag et al., 2013). IGF-1 levels decline in advanced age, and the extent of this decline is associated with increased risk of stroke incidence, increased mortality, and worsened functional outcomes post-stroke in humans (Johnsen et al., 2004; De Smedt et al., 2011; Tang et al., 2014; Armbrust et al., 2017; Saber et al., 2017). Moreover, altered IGF-1 levels are also associated with increased risk of multiple age-related neuropathologies marked by neurovascular distress, excitotoxicity, and oxidative stress (Watanabe et al., 2005; Sonntag et al., 2013; Ghazi Sherbaf et al., 2018; Gubbi et al., 2018). While many studies associate reduced IGF-1 with increased neurodegeneration, some clinical studies of dementias and stroke see increased IGF-1 levels at the start of pathology, which may be an attempted compensatory mechanism designed to protect the brain (Gubbi et al., 2018; Castilla-Cortazar et al., 2020). Preclinical studies suggest that IGF-1 plays a causal role

in the stroke risk and recovery processes by directly regulating cell damage during ischemia [as reviewed by Hayes et al. (2021)]. IGF-1 supplementation in rodent models of middle cerebral artery occlusion and photothrombotic stroke reduces the size of infarcted tissue and the accompanying sensorimotor deficits, even when administered in the days after insult (Liu et al., 2001; Bake et al., 2014, 2016; Parker et al., 2017; Serhan et al., 2019; Hayes et al., 2021). Because IGF-1 regulates numerous cells throughout the body, the cellular mechanism by which IGF-1 protects viability and function in the brain remains unclear. It is commonly inferred that the beneficial effects of IGF-1 are mediated by direct regulation of neurons. However, neurons, astrocytes, endothelial cells, and perivascular cells each express the receptor for IGF-1, IGFR. Together, these cells work to compose the neurovascular unit (also known as neuro-glio-vascular unit) which tightly coordinates physiological responses within the nervous system and is a central target for dysfunction in neurodegenerative diseases and pathologies (Stanimirovic and Friedman, 2012). Additional work is needed to tease apart the impact of each cell type when exogenous IGF-1 is administered to protect against the damage induced by stroke and neurodegenerative diseases.

Animal models of circulating IGF-1 deficiency exhibit impaired neurovascular coupling, compromised blood-brain barrier integrity, and increased prevalence of micro hemorrhages (Toth et al., 2015; Tarantini et al., 2016b, 2017, 2021a; Hayes et al., 2021). Increased reactive oxygen species (ROS) production, increased inflammation, and reduced glutamate and glucose handling machinery accompany the loss of circulating IGF-1 in these models (Toth et al., 2015, 2017). Together, these data suggest that the age-related reduction in IGF-1 likely leads to severe consequences on the structure and function of the neuro-glio-vascular unit. Indeed, *in vitro* studies have highlighted roles for IGF-1 in promoting specific functions of each cellular component of the neuro-glio-vascular unit. Pharmacological and genetic manipulations of IGF-1 alter the viability and oxidative stress levels of cultured neurons (Wang et al., 2014; Li et al., 2017; Chen et al., 2019). Additionally, IGF-1 signaling in brain vascular endothelial cell cultures promotes angiogenesis and tight-junction formation/integrity (Lopez-Lopez et al., 2004; Bake et al., 2016; Higashi et al., 2020). Genetic reductions of IGFR specifically in endothelial cells impairs stimulation-induced cerebral blood flow, indicating that the loss of IGF-1 signaling in one component of the neurovascular unit results in phenotypic changes commonly observed in the aged brain (Tarantini et al., 2021b). We recently reported that deficiency in IGF-1 signaling in astrocytes impairs glutamate handling *in vitro* and *in vivo*, by reducing glutamate transporter expression and availability at the cell surface (Prabhu et al., 2019). Although

there are thought to several key drivers to aging pathologies and disease, glutamate over excitation is a central contributor to tissue damage following ischemic stroke (Choi and Rothman, 1990; Moskowitz et al., 2010), it is likely that the age-related loss of IGF-1 may compromise the neuro-glio-vascular unit by weakening the glutamate-buffering capabilities of astrocytes which ultimately exacerbates calcium imbalances, mitochondrial dysfunction, and oxidative stress. While the aforementioned studies highlight the ability of IGF-1 to regulate neurons, astrocytes, and brain-derived endothelial cells, each of these studies was performed under varied conditions. A comparative approach is needed to better-understand how the age-related loss of IGF-1 influences cells within the neuro-glio-vascular unit. Thus, we exposed individual cell cultures and co-cultures to excitotoxic levels of glutamate when IGF-1 signaling was impaired. This approach allows for determination of the cellular mechanism(s) by which IGF-1 exerts its protective effects during ischemic stroke-induced stressors. In addition, it highlights cell-specific responses to a common driver of neuronal dysfunction- glutamate excitotoxicity- which has applications to numerous age-related neurodegenerative disease states.

MATERIALS AND METHODS

Animals

All procedures were approved by the Institutional Animal Use and Care Committees of the University of Mississippi, and performed in accordance with their approved guidelines. Timed-pregnant Sprague Dawley rats (16–18 days postictal plug) were purchased (Envigo) and were temporarily housed in 19 × 11.5 × 11 inch polycarbonate cages until the time of euthanasia. Rats were given access to standard rat chow (Teklad 7001) and water *ad libitum*. At the time of euthanasia, pregnant dams were anesthetized with isoflurane and cervical dislocation was completed. Embryonic pups (E17–19) were excised and rapidly decapitated. Due to the non-proliferative nature of neurons, the number of cells needed seeded per well, and the replication of results across multiple primary cell culture preparations, multiple pregnant dams were needed for these studies (approximately 7). The average litter size of a Sprague Dawley female is 10–11 pups, and cerebral tissues of all pups were pooled at the time of euthanasia. Isolated embryonic rat pups were not separated based on sex; thus, our studies are indicative of cell responses from both sexes combined. To reduce animal number, astrocytes and neurons were both cultured from the pooled tissues at the same time, in varying selection medias, as described below.

Neuron Cultures

Primary rat neuron cultures were established following previously described protocols (Ashpole and Hudmon, 2011). Cell culture dishes and coverslips were coated with Poly-D-Lysine (PDL; Sigma P6407) for at least 2 h at 37°C in a humidified incubator containing 5% CO₂. Individual cultures were plated in 96-well plates, while 15 mm glass coverslips were used to plate neurons for the triple co-cultures. Following PDL

coating, cell culture surfaces were washed with 1X PBS (Gibco 10010-023) and dried. Primary neuronal cultures were derived from the cortex and hippocampus of E17–19 Sprague Dawley rat pups. External cerebrovasculature was removed, the tissue was minced and subsequently enzymatically and mechanically digested with papain (Worthington Biochemical Corporation, LS003126) and glass-blown pipettes, respectively. Dissociated cells were pelleted with centrifugation (500–1,000 g, 5 min) and suspended in neuron growth media [Neurobasal medium (Gibco 21103-049) containing B27 (Life Technologies, 17504044), penicillin/streptomycin (10 units/mL; Life Technologies, 15140122), and 1x L-glutamine (250303-081)] with a target density of 2,500,000/mL. As neurons are non-dividing, the density of cells plated is indicative of the density at the time of experimentation; less than 10% of the cells in these cultures are GFAP + or IBA + positive, indicating they are predominantly neurons (Ashpole and Hudmon, 2011). Partial media changes occurred every 3–4 days post seeding to replenish nutrients and remove debris.

Astrocyte Cultures

Tissue from male and female rat pups was isolated, digested, and pelleted following the methodology described above for neuron cultures, and neurons and astrocytes were derived from the same isolated tissues. To promote astrocyte selection after digested tissue was centrifuged, cell pellets were suspended in astrocyte growth media containing Neurobasal medium (Gibco 21103-049) with 10% fetal bovine serum (Corning 35-010-CV), penicillin/streptomycin (10 units/mL; Life Technologies, 15140122), and 1x L-glutamine (250303-081). Cells were plated on PDL-coated 10 cm dishes and media was completely exchanged the day following plating. To further select for astrocytes, the cells were sub-cultured every 3–4 days (90–100% confluent). For this, the growth media was removed, cells were washed with 1X PBS, and 0.05–0.25% of Trypsin-EDTA 1X (Gibco 25200-072) was added. To avoid excess debris, trypsin was applied twice during the first passage- once for 2 min to dislodge neurons and microglial cells from the upper layer (vacuumed off), and a second time to release the astrocytes underneath. Once the astrocytes dislodged, the cells were centrifuged (1,000 × g; 5 min) and suspended in growth medium for further plating. All astrocyte cultures were passaged multiple times prior to experimentation (as described above), and the growth rate was continuously monitored to determine if cells were exhibiting replicative senescence. Cells were not passaged more than 6 times to avoid confounds of senescent phenotypes. The resulting cultures are >90% GFAP+, indicating they are astrocytes (Prabhu et al., 2019).

Endothelial Cell Culture

Rat brain microvascular endothelial cells (RBMVECs) and medium were purchased from Cell Applications Inc. (San Diego) and cultured following the manufacturer's guidelines. A single vial of RBMVEC were used to conduct all included studies. RBMVEC were continually passaged and 20% of cells were frozen at each passage throughout the study for replicate, follow-up, and future studies to reduce variance from a different lot of cells

originally isolated. In brief, cells were grown on a T-75 flasks pre-coated with Cell Attachment Factor Solution (123-100, Cell Applications), in 15 mL of RBMVEC growth medium (R819-500). Growth medium was changed 24 h after initial seeding, and cells were sub-cultured every 4–6 days or when at 80–100% confluency. Media was changed every other day. RBMVECs were passaged using the same protocol as astrocytes, using the proprietary trypsin solution (Cell Applications Inc).

Co-cultures

Triple cultures were developed by plating primary neurons on glass coverslips, RBMVECs on transwell inserts, and astrocytes on the surface of a multi-well plate. Astrocytes were cultured and passaged at least 1 week prior to combination with the other cells, to allow for selection of astrocytes away from neurons and microglial cells. Neurons were also cultured 7–8 days prior to combination and experimentation, to allow for development/expression of ionotropic glutamate receptors (GluNR) responsible for excitotoxic signaling. Coverslips were elevated off the surface of the multi-well plate and astrocytes were seated in the outer 1/3 of the wells. The media on all 3 cell types was exchanged for neuron growth media 24 h prior to combination of the co-cultures to allow for uniformity between media types.

Cells Treatments

Glutamate stimulation was performed using L-glutamic acid and glycine (FisherScientific) in a ratio of 10:1 glutamate:glycine diluted from a PBS stock solution. Increasing concentrations of glutamate were prepared in the appropriate cell growth media, and administered as a 2x solution to the cells plated in their respective media (final concentration 1x upon application). Cells were incubated for 1 h at 37°C. Following treatment, glutamate media was removed, cells were washed in fresh growth media twice, and incubated in new growth media until the designated time point. All cell viability studies were conducted 24 h after cells were stimulated with glutamate. All other study time points are described in the respective sections. In one experimental series, media lacking the B-27 supplement was administered 24 h prior to glutamate stimulation.

Stock solutions of picropodophyllin toxin (PPP; Sigma T-9576) and MK-801 (Sigma M107) were prepared in dimethyl sulfoxide (DMSO; Fisher BP321-100) and administered to cells at concentrations and time-points described. Treatment groups were block randomized to ensure equal sampling. Individual wells in the multi-well plates were treated as independent samples; and assays with neurons and astrocytes were further repeated across multiple tissue preparations to verify reproducibility of findings in cells isolated from separate rat pup litters.

Cell Viability

Following treatments, cell viability/cytotoxicity was quantified using Live:Dead viability/cytotoxicity kit (Thermo Fisher Scientific L3224). Calcein-AM and ethidium homodimer were diluted (1:1000–2000; 1:500–1000) were diluted in PBS. Growth media was removed and cells were washed with 1X PBS prior

to adding the live/dead labels. After 20 min of incubation, cells were imaged on the Nikon Ti2-E HCA inverted fluorescent microscope using the JOBS automated image acquisition (Nikon). Magnification was set to 200x (20x, extra-long working distance objective) and samples were excited with the LED Triggered acquisition exposures using excitation/emission filters for 470-FITC and 540-TRITC. The JOBS program selected 6 fields at random per well for imaging, and the total number of live and dead cells were quantified in 2–3 images using the Nikon Elements Cell Analysis plug-in features. The total number of cells and percent toxicity were calculated by observers blinded to treatment groups.

Reactive Oxygen Species Quantification

Following treatment, ROS production was quantified in astrocytes and endothelial cells. Media was removed, and cells were washed with 1X PBS. Cells were then loaded with 3 μ M of 2',7'-dichlorodihydrofluorescein diacetate (DCFDA; Thermofisher, D399) for 20 min and incubated at 37°C in a humidified incubator containing 5% CO₂. DCFDA was then imaged using the FITC filter on the Nikon Ti2-E microscope as described above. Six images at random through the JOBS interface and 4–6 non-adjacent cells per field were selected at random to be quantified by a blinded observer. DCFDA intensity of each cell was normalized to the background of that image.

Seahorse

Cells were seeded on the XFe96 growth plates 2–3 days prior to extracellular efflux assessment. Treatments were administered at the desired time points and mitochondrial response was assessed using the Cell Mito Stress Test, per manufacturer recommendations. The sensor cartridge was hydrated in water overnight at 37°C and then equilibrated with the recommended equilibration buffer. Oligomycin (1.5 μ M), FCCP (1 μ M), and rotenone (0.5 μ M) were loaded into the sensor plates and injected into the wells by the XFe/XF Seahorse system. Basal and maximal respiration, as well as proton leak were auto-calculated by the Seahorse program using an average of 3 readings per stage.

Data Analysis

Statistical analysis and graphical analyses were performed using Sigma Plot version 14 software and R studio. One-way and two-way ANOVAs was used when appropriate (defined in the figure legend). *Post hoc* comparisons were selected based upon the experimental question and fulfillment of normality and equal distribution. Details for each are provided in figure legend. All data were expressed with mean \pm standard error. Sample sizes were estimated using previous variance observations with these experimental methodologies. With $\beta = 0.8$, $n = 4–6$ was required to achieve power for most studies. *F* values for two-way ANOVAs are described in the results and main effects are noted in figures with a pound sign (#). For all studies, $p < 0.05$ was used as the statistical significance value and denoted using an asterisk (*).

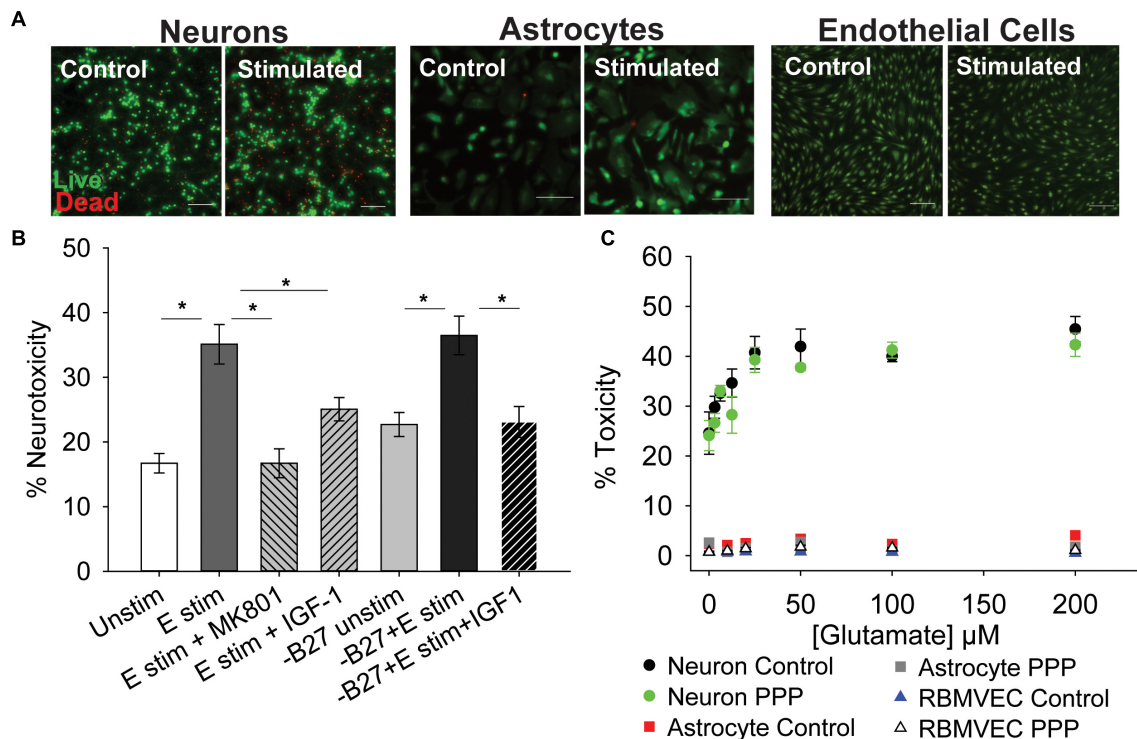


FIGURE 1 | IGF-1 protects against excitotoxicity in glutamate-sensitive neurons. **(A)** Representative images of neurons (left two panels), astrocytes (middle two panels), and rat brain microvascular endothelial cells (right two panels) stained with live (green)/Dead (red) viability/cytotoxicity indicator dyes 24 h after stimulation with control or 100 μ M glutamate, as indicated. Scale bar represents 100 μ m. **(B)** Average toxicity in neurons 24 h after treatment ($n = 8$ wells/group). A one-way ANOVA with *post hoc* Bonferroni was used for statistical comparisons; *indicates significant difference between the delineated bars. **(C)** Average toxicity of cells 24 h after acute treatment with both 0.5 μ M PPP and increasing concentrations of glutamate ($n = 4$ –8 wells/group). A two-way ANOVA was employed with glutamate concentrations and PPP-treatment serving as factors. No significant differences were observed. Graph color code: black circles = vehicle-treated neurons; green circles = PPP-treated neurons; red boxes = vehicle-treated astrocytes; dark gray boxes = PPP-treated astrocytes; blue triangles = vehicle-treated rat brain microvascular endothelial cells (RBMVEC); and light gray triangles = PPP-treated RBMVEC. All data are presented as mean \pm SEM.

RESULTS

Insulin-Like Growth Factor-1 Protects Against Excitotoxicity in Neurons

To determine if inhibition of IGF-1 signaling differentially affects the glutamate sensitivity of cells within the neuro-glio-vascular unit, cultures of neurons, astrocytes, and microvascular endothelial cells were established. Primary cerebral neurons were isolated from embryonic rat pups (male and female combined) and grown in culture for 8–10 days. Primary neurons at this stage of development are sensitive to excitotoxicity when exposed to high concentrations of glutamate and glycine, due to overactivation of ionotropic GluNR. As expected, acute treatment of neurons with 100 μ M glutamate/10 μ M glycine (termed glutamate stimulation throughout) resulted in a significant increase in cellular toxicity 24 h after stimulation (**Figures 1A,B**; $p < 0.001$ vs unstimulated). Co-administration of the GluNR antagonist MK-801 prevented excitotoxicity (**Figure 1B**; $p < 0.001$ vs glutamate stimulated control). Administration of exogenous 100 nM IGF-1 at the time of stimulation also prevented excitotoxicity ($p = 0.019$ vs glutamate stimulated control), consistent with previous reports.

No differences in the total number of neurons per field were noted following treatment (**Supplementary Figure 1A**).

The neuronal growth media contains a supplement (B27) that has a high concentration of insulin, which can cross-talk and activate IGFR. Thus, to better determine the protective effects of IGF-1 signaling, the excitotoxicity profile was again assessed without B27 present. No change in baseline levels of toxicity were observed between control neurons with and without B27, and glutamate stimulation still significantly increased toxicity in the absence of B27 (**Figure 1B**; $p = 0.002$ vs -B27 unstimulated control). Once more, exogenous IGF-1 prevented excitotoxicity (**Figure 1B**; $p = 0.003$ vs -B27 glutamate stimulated control).

To more specifically assess the impact of reducing IGF-1 signaling, the extent of excitotoxicity when IGFR was pharmacologically inhibited was then examined. For this, picropodophyllotoxin (PPP), a known small molecule inhibitor of the IGFR receptor kinase activity, was co-administered at the time of glutamate stimulation and toxicity was measured 24 h later. Surprisingly, PPP did not shift the concentration-dependence or maximal extent of glutamate toxicity in cultured neurons (**Figure 1C**). A two-way ANOVA revealed a significant effect of glutamate concentration ($F = 13.759$, $p < 0.001$), but no

significant effect of PPP treatment ($F = 2.598$, $p = 0.111$), and no significant interaction between the two factors ($F = 0.506$, $p = 0.827$).

Astrocytes and microvascular endothelial cells are not as sensitive to glutamate as neurons as no change in toxicity was observed across all glutamate concentrations (Figures 1A–C). Similarly, no increase in sensitivity was observed with IGFR inhibition at the time of glutamate stimulation (Figure 1C).

Prolonged IGFR Inhibition Increases Astrocytic Sensitivity to Glutamate

The lack of effect with pharmacological inhibition of IGFR in Figure 1C was inconsistent with the protective effects of exogenous IGF-1 in Figure 1B. However, co-administration of PPP at the time of glutamate could have temporal confounds that limit interpretation. Thus, we next administered PPP 24 h prior to glutamate stimulation to better mimic the prolonged loss of IGFR signaling observed in aging. Prolonged IGFR inhibition with 0.5 μ M PPP did not lead to difference in neuronal excitotoxicity ($F = 0.255$, $p = 0.615$), but did lead to differences in astrocyte viability ($F = 6.677$, $p = 0.011$; Figures 2A,B). Within group comparisons did not unveil a specific concentration of glutamate that PPP treatment significantly worsened, thus it is assumed that IGFR inhibition led to modest reductions in cell viability across all glutamate treatments.

No changes in endothelial cell viability were observed with prolonged IGFR inhibition; however, a pronounced reduction in total cell number was noted when microvascular endothelial cells were treated with 0.5 μ M PPP ($F = 117.942$, $p < 0.001$; Figure 2C). A significant difference within unstimulated controls with PPP pre-treatment was detected ($p < 0.001$), and this effect persisted across all glutamate concentrations. Thus, no main effect of, or interaction with, glutamate concentration was observed in the treated microvascular endothelial cells.

The change in endothelial cell number prompted further analyses of neuron and astrocyte counts with treatment. Similar reductions in astrocyte count were observed with 0.5 μ M PPP treatment ($F = 73.322$, $p < 0.001$; Figures 2A–C). Glutamate concentration did not alter astrocyte count nor interact with the PPP effect. Main effects of both glutamate concentration and PPP treatment on total neuron count were detected ($F = 3.309$, $p = 0.004$; $F = 7.835$, $p = 0.006$, respectively). *Post hoc* comparisons within groups revealed that the only significant difference within the PPP vs vehicle treatments was the drop in total cell number at 200 μ M glutamate ($p = 0.012$), likely an indication of dead cells washed away during the staining process at this high concentration of glutamate.

When the concentration of PPP was increased to 5 μ M, significant differences in neuron and astrocyte survival were observed (Figure 2D). Both glutamate concentration and PPP treatment had significant main effects on neurotoxicity ($F = 4.272$, $p < 0.001$; $F = 12.565$, $p < 0.001$, respectively). However, no interaction between the two factors was detected ($F = 0.732$; $p = 0.646$). Within group comparisons revealed significant increases in neurotoxicity when neurons pre-treated with 5 μ M PPP were stimulated with 50 and 200 μ M glutamate

($p = 0.048$ and $p = 0.006$, respectively). No baseline differences in toxicity amongst unstimulated controls treated with vehicle or PPP were observed ($p = 0.751$). A significant reduction in total neuron number was observed with 5 μ M PPP treatment and with glutamate ($F = 49.058$, $p < 0.001$; $F = 2.878$, $p = 0.009$, respectively; Figure 2E). No significant interactions between the factors was detected ($F = 0.932$, $p = 0.485$). Within factor comparisons showed decreased cell number within PPP-treated neuron cultures stimulated with ≥ 12.5 μ M glutamate, again suggesting potential loss of dead cells during the staining process since neurons are non-mitotic.

Prolonged IGFR inhibition in astrocytes also resulted in main effects of glutamate concentration and PPP treatment on the levels of cytotoxicity ($F = 157.321$, $p < 0.001$; $F = 2.209$, $p = 0.039$, respectively; Figure 2D). Moreover, a significant interaction between glutamate concentration and PPP treatment was also observed ($F = 2.743$, $p = 0.011$). Within factor comparisons revealed significant increases in PPP-pretreated astrocytes stimulated with ≥ 6 μ M glutamate, suggesting that while astrocytes display increased resistance to glutamate toxicity under control conditions, the loss of IGF-1 signaling predisposes astrocytes to excitotoxic stress. A significant reduction in astrocyte cell number was also observed with 5 μ M PPP treatment ($F = 257.893$, $p < 0.001$), across all glutamate concentrations. Main effect differences for glutamate concentration did not reach significance ($F = 1.882$, $p = 0.079$), nor did interactions between factors ($F = 2.071$, $p = 0.052$). Prolonged treatment with another IGFR inhibitor, NCP-ADW742, also increased susceptibility to glutamate excitotoxicity and reduced astrocyte cell number (Supplementary Figures 1B,C). Baseline toxicity levels and cell numbers were observed with administration of supplemental IGF-1 prior to glutamate stress (Supplementary Figures 1B–E), indicating that the prolonged loss of IGF-1 predisposes astrocytes to glutamate toxicity, despite their typical resistance to the stress.

Similar to the before no differences in endothelial cell toxicity were observed across any treatment groups, but 5 μ M PPP did reduce cell number across all glutamate concentrations ($F = 538.084$, $p < 0.001$; Figure 2E). Again, this suggests that while IGFR inhibition did not predispose endothelial cells to glutamate toxicity, it likely reduces cell division.

Astrocytes Fail to Protect Neurons When IGFR Is Inhibited

Cells within the neuro-glio-vascular unit coordinate to create an optimal microenvironment, by releasing growth mediators, buffering stressors, and regulating nutrient supply. Thus, we next assessed whether co-cultures of endothelial cells, neurons, and astrocytes would exhibit differences in glutamate sensitivity when IGFR is inhibited. As anticipated, 100 μ M glutamate did not significantly increase neurotoxicity in the presence of astrocytes and endothelial cells, nor did acute co-administration of 0.5 μ M PPP with 100 μ M glutamate ($p = 0.139$; Figure 3A). No differences in astrocyte or endothelial cell toxicity were observed in these acute treatments (Figures 3B,C).

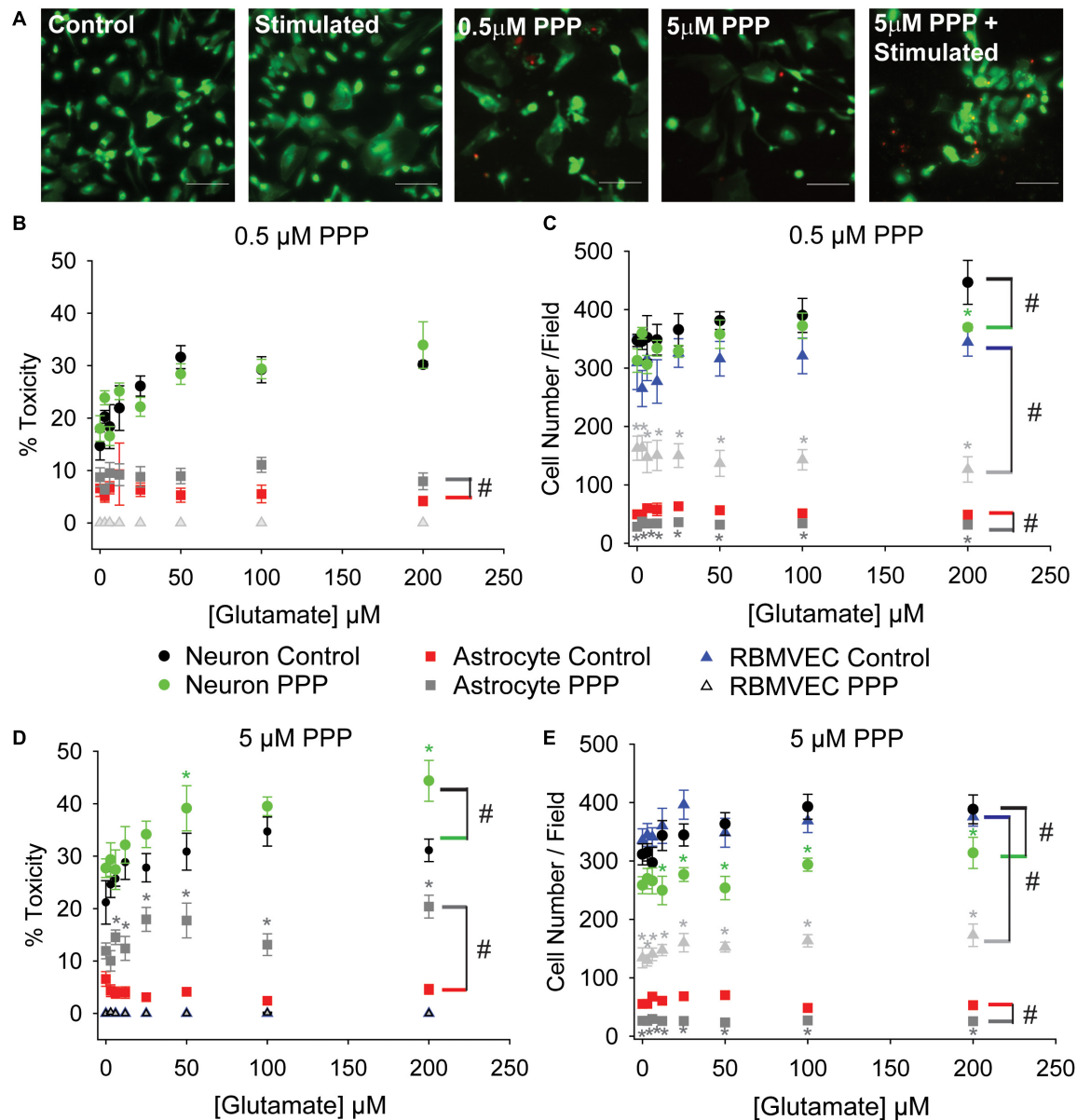


FIGURE 2 | Prolonged IGF1R inhibition increases astrocytic sensitivity to glutamate and reduces astrocyte and endothelial cell number. **(A)** Representative images of astrocytes stained with live (green)/Dead (red) viability/cytotoxicity indicator dyes. From left to right: unstimulated control, glutamate-stimulated vehicle-treated, unstimulated 0.5 μM PPP-treated, unstimulated 5 μM PPP, and glutamate-stimulated 5 μM PPP. Scale bar represents 100 μm. **(B)** Average toxicity and **(C)** average cell number per well of cells pre-treated with 0.5 μM PPP, and subsequently stimulated with increasing concentrations of glutamate, 24 h after stimulation. **(D)** Average toxicity and **(E)** average cell number per well of cells pre-treated with 5 μM PPP, and subsequently stimulated with increasing concentrations of glutamate, 24 h after stimulation. All data are presented as mean ± SEM, $n = 5-8$ wells/treatment group. A two-way ANOVA was employed with glutamate concentrations and PPP-treatment serving as factors. Cell types were each analyzed independently. # indicates a significant main effect within that cell type. * indicates a significant difference between PPP-treatment and vehicle-treatment within a given glutamate concentration. Graph color code: black circles = vehicle-treated neurons; green circles = PPP-treated neurons; red boxes = vehicle-treated astrocytes; dark gray boxes = PPP-treated astrocytes; blue triangles = vehicle-treated rat brain microvascular endothelial cells (RBMVEC); and light gray triangles = PPP-treated RBMVEC.

When astrocytes alone were pre-treated with 5 μM PPP and then combined with neurons and endothelial cells for glutamate stimulation, a significant increase in neurotoxicity was observed ($p = 0.007$ vs control) (**Figure 3D**). No differences in toxicity or cell number were noted in either the astrocytes or endothelial

cells within the triple cultures (**Figures 3E,F** and **Supplementary Figures 1D-F**). Together these data suggest that a reduction of IGF1R signaling in astrocytes impairs their ability to buffer glutamate and protect neurons from overexcitation, even when in the presence of endothelial cells.

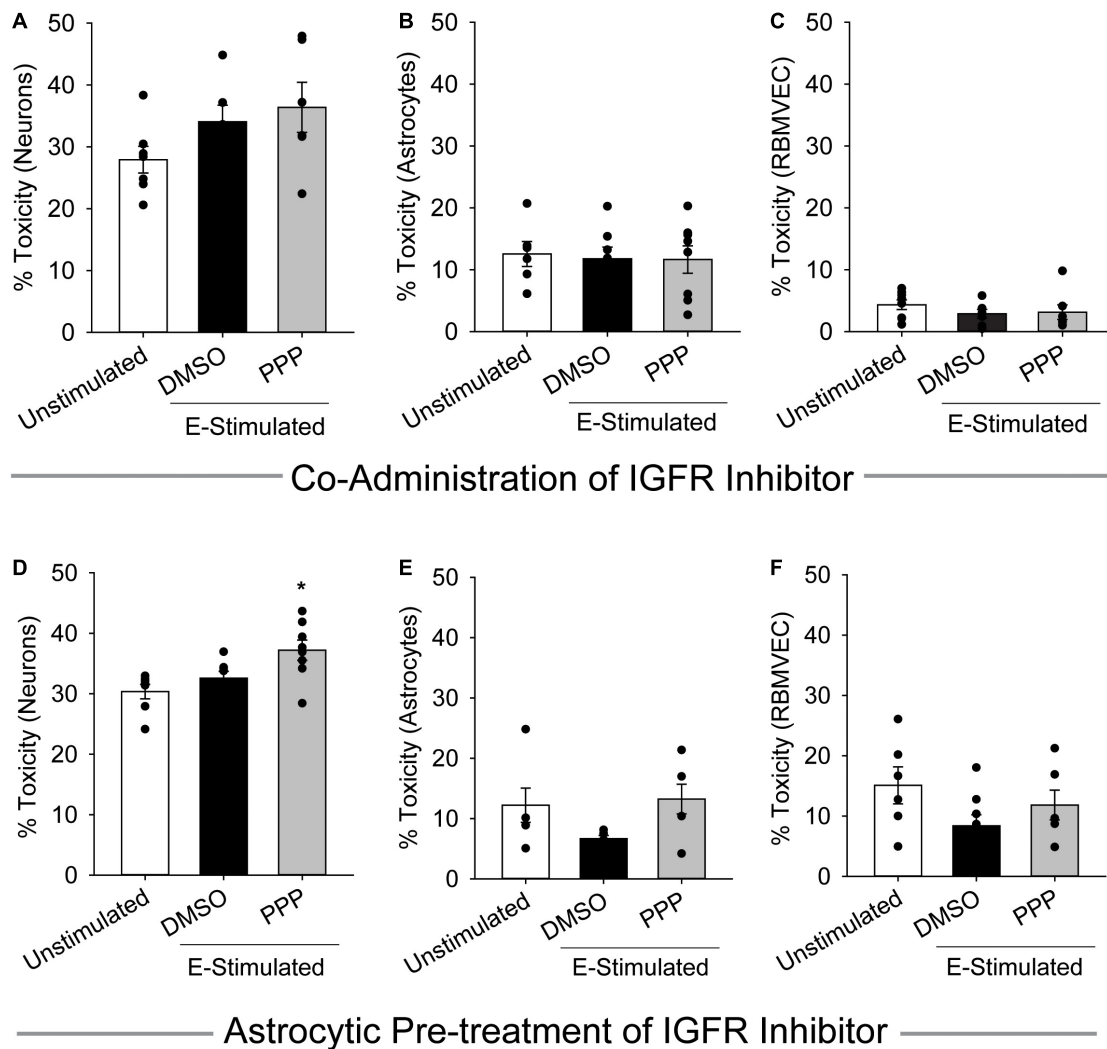


FIGURE 3 | Astrocytes fail to protect against excitotoxicity when IGFR is inhibited. Average cell death 24 h after triple cultures of neurons (A), astrocytes (B), and endothelial cells (C) were co-administered 0.5 μ M PPP and 100 μ M glutamate for 1 h ($n = 6-8$ wells/group). One-way ANOVA failed to detect differences in any of the treatment groups. (D-F) Astrocytes were pre-treated with 5 μ M PPP for 24 h prior to combining with endothelial cells and neurons for a triple culture system. Average cell death in the neurons (D), astrocytes (E), and endothelial cells (F) after stimulation with 100 μ M glutamate for 1 h ($n = 5-8$ wells/group). One-way ANOVAs with *post hoc* Bonferroni comparison was used for statistical analyses. *indicates significant difference between control and PPP-treated glutamate-stimulated neurons. All data are presented as mean \pm SEM, $n = 5-8$ wells/treatment group.

Astrocytic and Endothelial Reactive Oxygen Species Levels Are Elevated by IGFR Inhibition and Glutamate Stimulation

While acute co-administration of 0.5 μ M PPP and glutamate did not increase toxicity in astrocyte or endothelial cultures, alterations in metabolic function and oxidative stress may still occur with this stressor. Thus, astrocytes and endothelial cells were treated with 100 μ M glutamate alone, or in conjunction with 0.5 μ M PPP, and the production of ROS was quantified at various time points following treatment. Astrocytic ROS production increases in the hours following glutamate stimulation (Figure 4A). Significant effects of both

time and treatment were observed ($F = 185.57$, $p < 0.001$; $F = 3.187$, $p = 0.027$, respectively), as well as an interaction between both factors ($F = 4.586$, $p < 0.001$). 5 h post-treatment, vehicle treated and PPP-treated astrocytes stimulated with glutamate showed significant increases in ROS levels over unstimulated controls (both $p < 0.001$ vs unstimulated control). Moreover, co-administration of PPP and glutamate resulted in significantly increased ROS levels than glutamate stimulated controls ($p = 0.014$), suggesting that the extent of glutamate-induced ROS production was exacerbated by IGFR inhibition. 5 h after glutamate stimulation with another IGFR inhibitor, NVP-ADW742, also significantly increased ROS (Supplementary Figure 2A), indicating the effect is not limited to PPP. Additional analysis of mitochondrial bioenergetics at

this time point revealed that glutamate stimulation decreased maximal extracellular acidification rates, while PPP + glutamate decreased extracellular acidification and oxygen consumption rates, suggesting changes in both glycolysis and oxidative phosphorylation (**Supplementary Figure 2B**). No differences in basal respiration rates were noted with any of the treatments (**Supplementary Figure 2B**).

Similar to astrocytes, brain-derived microvascular endothelial cells showed increased ROS production 5 h after 0.5 μ M PPP was co-administered with 100 μ M glutamate (**Figure 4B**). A main effect treatment was observed ($F = 5.287$, $p = 0.004$). While glutamate alone did not increase ROS production in the endothelial cells ($p = 0.945$), a significant increase in ROS was seen in cells treated with PPP + glutamate vs unstimulated control ($p = 0.02$), and vs glutamate-stimulated control ($p = 0.004$).

Additional analyses of ROS levels 24 h following treatment revealed differential recovery from acute IGFR inhibition and glutamate stimulation. Astrocytes do not show increased ROS at 24 h with 100 μ M glutamate or 100 μ M glutamate + 0.5 μ M PPP treatment (**Figure 4C**). In fact, ROS levels were significantly reduced in glutamate-stimulated PPP-treated astrocytes vs unstimulated controls ($p = 0.024$) and PPP treatment was trending ($p = 0.053$ vs control). Endothelial cells continue to show increased ROS levels long after the acute exposure to 100 μ M glutamate + 0.5 μ M PPP ($p = 0.013$ vs unstimulated control; **Figure 4D**), suggesting that IGFR inhibition at the time of glutamate stimulation increases endothelial ROS levels long-term. Note, this was with acute PPP treatment, at a time point in which no differences in endothelial cell toxicity was observed (**Figure 1C**).

DISCUSSION

The age-related loss of IGF-1 has been linked to cognitive impairment, neurodegeneration, and increased susceptibility ischemic stroke and other neurovascular pathologies (Sonntag et al., 2013; Ashpole et al., 2015b). These deficits often come as a disconnect from the enhanced longevity observed in animal models of IGF-1 deficiency. While early-life reductions in circulating IGF-1 do lead to increased lifespan, reduced IGF-1 in advanced age has multiple consequences in animal models as it increases sarcopenia, bone frailty, learning and memory deficits, and cerebrovascular dysfunction (Ashpole et al., 2015a, 2016, 2017; Toth et al., 2015; Tarantini et al., 2016a, 2021a; Farias Quipildor et al., 2019). For example, IGF-1 deficiency in mice decreases stimulation-evoked cerebral blood flow, blood brain barrier integrity, and the strength and flexibility of cerebral arteries. IGF-1 deficiency also impairs the production and release of vasomediator eicosanoids from astrocytes, alters endothelial nitric oxide production, and increases susceptibility to hypertension-induced microhemorrhages (Toth et al., 2015; Tarantini et al., 2017, 2021a; Fulop et al., 2019). Considering this, it is not surprising that IGF-1 has long-been implicated in the risk and severity of ischemic stroke. As described earlier, clinical and preclinical evidence highlight inverse correlation (and causation

in animal models) between IGF-1 and the risk/outcome of ischemic stroke (Hayes et al., 2021).

While recent studies are beginning to shed light on the impact of IGF-1 on neurovascular communication in advanced age, there remains a significant gap in our understanding of how individual cell types that normally coordinate signaling together within the neurovascular unit each respond to reduced IGF-1 signaling. In this study, we aimed to compare the effects of IGF-1 deficiency on the cellular responses to high levels of extracellular glutamate. As the predominant excitatory neurotransmitter in the brain, glutamate induces depolarization and initiates multiple calcium signaling cascades. During periods of ischemia or proteotoxic stress, neurons experience energy imbalances and ionic stressors that ultimately lead to terminal depolarization and the dumping of glutamate stores, inflammatory mediators, and oxidative stressors into the extracellular space, which expands the region of distress and malfunction [as recently reviewed (Choi, 2020)]. This excitotoxic cascade is one of the drivers of neuronal loss and glial activation in ischemic stroke and neurodegenerative diseases. Thus, understanding how the loss of IGF-1 alters the glutamate response of neurons, astrocytes, and endothelial cells within the neurovascular unit provides much-needed information on potential cellular mechanisms by which IGF-1 deficiency in advanced age influences health and function of the brain.

Recent evidence from our laboratory indicated that IGFR inhibition reduced the ability of astrocytes to buffer extracellular glutamate by decreasing glutamate transporter availability (Prabhu et al., 2019). Therefore, in this study, we hypothesized that maintenance of IGF-1 signaling in astrocytes is necessary for neuroprotection from glutamate excitotoxicity. We also hypothesized that maintenance of IGF-1 signaling in neurons was essential for limiting overexcitation, as other studies have shown exogenous IGF-1 can protect against excitotoxicity and oxidative stress (Wang et al., 2014; Li et al., 2017; Chen et al., 2019). Our results indicate that while exogenous IGF-1 can protect neurons from glutamate, acute and prolonged loss of IGFR signaling does not exacerbate neuronal death in cultures of pure neurons. Interestingly, inhibition of IGF-1R not only impairs the neuroprotective capabilities of astrocytes, it predisposes astrocytes to glutamate toxicity as well. It is unclear what the mechanism for this enhanced sensitivity may be. Perhaps the lack of glutamate transporter availability with IGF-1 signaling deficiency resulted in increased GluNR activation in the astrocytes as well, since astrocytes are known to express a variety of ionotropic and metabotropic GluNRs (Serrano et al., 2008; Lalo et al., 2011; Bradley and Challiss, 2012; Ceprian and Fulton, 2019). If this were the case, the astrocytes could be experiencing intracellular calcium imbalances and oxidative stress, which have both been shown to increase astrocyte toxicity in response to other stressors.

As mentioned, reductions in IGF-1 in the brain increase ROS levels and mitochondrial dysfunction. We observed significant increases in ROS production in both astrocytes and endothelial cells when IGFR was inhibited at the time of glutamate stimulation. We chose to focus our experimental design on these two cell types and this acute treatment because it did not lead to any changes in observed toxicity. Oxidative stress is expected in

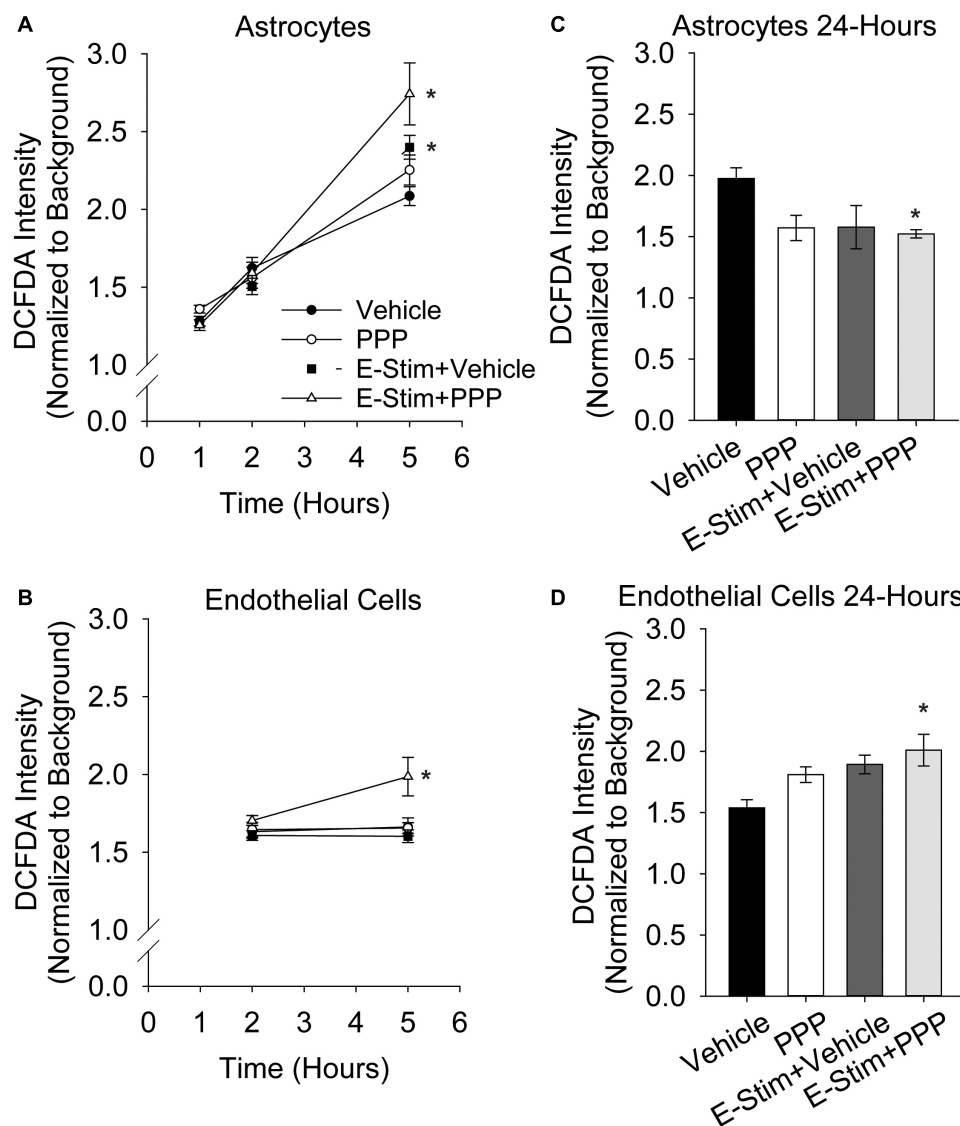


FIGURE 4 | Astrocytic and endothelial ROS levels are elevated by IGF1R inhibition and glutamate stimulation. Average ROS levels in pure cultures of astrocytes (**A**) or endothelial cells (**B**) co-treated with 0.5 μ M PPP and 100 μ M glutamate for 1 h, and subsequently measured 1, 2, and 5 h after treatment. Four cells were selected at random per field, with 3 fields per well, and 8–10 wells per treatment group. Background intensity within each image was measured and used to normalize. A two-way ANOVA was employed with both time and treatment as factors, and a *post hoc* Holm-Sidak was used to compare within groups. * indicates a significant effect of treatment at that time. Average ROS levels in pure cultures of astrocytes (**C**) or endothelial cells (**D**) co-treated with 0.5 μ M PPP and 100 μ M glutamate for 1 h, and subsequently measured 24 h after treatment. Four cells were selected at random per field, with 3 fields per well, and 4–8 wells per treatment group. A one-way ANOVA with *post hoc* Dunnett's test was used for statistical comparisons to control. * indicates a significant effect vs control. All data are presented as mean \pm SEM.

neurons that are dying from excitotoxicity, but neither astrocytes nor endothelial cells were sensitive to glutamate when IGF1R was acutely inhibited. The increased ROS levels in both cell types indicates that the cells were indeed undergoing a stress response with the combined treatment. While it could be inferred that the ability of astrocytes to return ROS levels back to baseline by 24 h may contribute to their increased resistance to excitotoxic stress, there is a disconnect in the endothelial cells that likely renders this to be more complicated. Perhaps an increase in endothelial cell toxicity would have been observed at later time points. Further

studies on the underlying mechanism(s) and consequence(s) of these observations are needed.

Combined triple cultures of neurons, astrocytes, and endothelial cells highlighted a few distinct responses to glutamate and IGF1R inhibition than the individual cultures of each cell type. First, the same concentration of glutamate that led to significant increases in neurotoxicity within pure neuron cultures failed to induce toxicity within the triple culture. This was an expected outcome, as astrocytes are known to buffer glutamate and co-cultures of neurons and astrocytes have

been shown to reduce the extent of neurotoxicity (Choi et al., 1987; Rosenberg et al., 1992). Second, and more interestingly, prolonged IGFR inhibition in astrocytes prior to glutamate stimulation did not increase the toxicity of astrocytes nor decrease the total cell number of astrocytes within the triple cultures. These effects were pronounced in pure astrocyte cultures, suggesting that co-culturing astrocytes with neurons and endothelial cells afforded them protection against the stressors of IGFR deficiency/glutamate stimulation. Additional studies examining ROS levels, inflammatory mediators, and growth factor production and release with triple cultures exposed to these same stressors may be warranted.

In our recent study examining glutamate uptake in astrocytes, we utilized similar concentrations of the pharmacological inhibitor PPP and did not observe a significant reduction in total cell number. Thus, we were surprised to see a reduction in astrocyte count here. However, in that study, cells were treated when at or close to confluence, which was evident in DAPI-stained microscopy images accompanying those data (Prabhu et al., 2019). In the current study, we aimed to treat at 75% confluence in order to avoid stress of over-growth, however, microscopic live/dead staining analysis showed that even the vehicle-treated controls were not at this level of confluence even 24 h after treatment. Thus, the reduction in cell number observed in pure cultures of astrocytes and endothelial cells was likely due to changes in cell division and growth with IGF-1 deficiency. Nevertheless, it is interesting that the reduction in astrocyte number with prolonged IGFR inhibition was restricted to the pure cultures, and not the triple cultures.

The complication of cell proliferation also limited our ability to utilize genetic approaches in our *in vitro* studies. IGF-1 is a primary growth factor with autocrine and paracrine functions, and genetic reductions that alter cell proliferation make it difficult to control studies of neuroendocrine hormones. We originally aimed to include siRNA and/or cell-specific knock-outs using our inducible cell-specific IGFR knock-out lines. However, due to obvious insufficient cell proliferation caused by siRNA transfection/transduction (both approaches were attempted) as well as Cre-recombinase induced knock-outs, we concluded that the significant reductions in cell number would confound results. Astrocyte count following knock-out with siRNA or with 4-OH-tamoxifen-induced recombination was less than 50% of that observed in the control-treated wells (siRNA vector transfected or vehicle-treated). Hence, pharmacological interventions were utilized to address our central question of how the loss of IGF-1 impacts the major cells of the neurogliovascular unit. While exogenous IGF-1 was added in a few studies here, the growth media of the cells contains insulin and IGF-1 at concentrations known to promote growth and development, thus pharmacological inhibitors were selected to more specifically reduce IGFR signaling. Alternative studies could alter the density of seeded cells to account for the loss in proliferation with knock-out, however, compensatory changes in other growth receptors (GH, InsR, etc.) and IGF-1 binding proteins will also need to be examined as they are known to occur with IGF-1 manipulations *in vivo* (Ashpole et al., 2015b).

Our comparative study is limited by not including additional cells found within the brain and blood-brain barrier. For example, pericytes not only serve as a protective layer of the blood brain barrier, they regulate blood capillary diameter within the brain by dilating and relaxing in response to glutamate (Kisler et al., 2017, 2020; Brown et al., 2019). Because of this, pericytes are important during ischemic reperfusion as they can assist with restoring flow of oxygen and glucose to the ischemic tissue (Yang et al., 2017; Khennouf et al., 2018; Sun J. et al., 2020). Little is known about the influence of IGF-1 on pericytes within the brain, and additional studies on how IGF-1 may influence pericytic glutamate response are needed.

Additionally, microglia are known to respond to excitotoxic insult, ischemia, and reperfusion. In fact, microglia have been implicated in the exacerbation of damage following glutamate excitotoxicity and ischemic stroke, due to increased production of ROS and pro-inflammatory mediators (Ma et al., 2017; Qin et al., 2019; Lian et al., 2020). At the same time, microglia also play an important protective role following tissue stroke damage, so the duality of their contribution to physiology/pathophysiology cannot be over-simplified to just a negative contribution to damage (Ma et al., 2017; Qin et al., 2019; Marino Lee et al., 2021). The age-related loss of circulating IGF-1 is associated with increased microglial activation, neuroinflammation, and susceptibility to a compromised blood-brain barrier. The relationship between IGF-1 and microglial activation is not limited to temporal correlations, as there are numerous studies that highlight IGF-1 directly regulates the structure and function of microglia. Exogenous IGF-1 alters phenotypic activation states and reduces microglial-associated production and release of tumor necrosis factor- α (TNF- α), interleukin-1 beta (IL-1 β), inducible nitric oxide synthase (iNOS), and ROS (Park et al., 2011; Grinberg et al., 2013). Administration of IGF-1 following ischemic and/or hemorrhagic stroke attenuates microglial activation, inflammation, and ROS (Serhan et al., 2020; Sun Z. et al., 2020). Considering this, we expect that reductions in IGF-1 signaling within our cell culture system would also promote pro-inflammatory and oxidative stress phenotypes, which would further neuronal death. However, one study highlights that microglial presence in tri-cultures are neuroprotective and reduces neurons loss and astrocyte hypertrophy even though the presence of microglia increased inflammatory cytokine production compared to cell culture without microglia (Goshi et al., 2020). Future studies of ischemic and excitotoxic stressors should take microglial contributions into account while controlling for the natural migratory phenotype microglia display following activation, as results may be confounded by increased basal levels of neuroinflammatory stress (which often show an activated phenotype in culture).

Transitioning to an *in vivo* system would circumvent concerns regarding the lack of including all cell types present in the cerebral tissue within our co-culture models; however, the intricacies of timed, cell-specific manipulations make *in vivo* models difficult, particularly when trying to assess effects without developmental compensation interference. Inducible transgenic animals for each cell type are needed to rigorously assess the effects of the loss

of IGF-1 in adulthood on each component of the neuro-glio-vascular unit. We previously established an inducible astrocyte-specific transgenic mouse model, and identified that glutamate handling machinery was reduced in the brains of these mice, which served as a rationale for studying glutamate excitotoxicity here (Prabhu et al., 2019). A recent study highlighted changes in neurovascular coupling when IGFR signaling was targeted in adult microvascular endothelial cells (Tarantini et al., 2021b). Thus, inducible models of astrocytic and endothelial cell IGFR knock-outs are available. Our singular and co-culture *in vitro* studies allow us to begin to delineate and compare the responses of specific cells without the confounds of other variables like paracrine compensation or pharmacological drug delivery or metabolism in one cell type over another. Further side-by-side comparisons of excitotoxic or ischemic damage in mice with inducible IGFR knock-out in neurons, astrocytes, endothelial cells, pericytes, and microglia would confirm the role for IGF-1 regulation of each cell, while also revealing other potential changes in physiology and pathophysiological responses in the intact brain.

Together, our study highlights that cell types within the neurovascular unit differentially respond to IGF-1 signaling. Despite differences in baseline susceptibility to glutamate, both neurons and astrocytes are both protected from excitotoxicity by IGF-1. Growth and division of endothelial cells and astrocytes are both influenced by IGF-1, however, endothelial cells remain resistant to glutamate toxicity even when IGF-1 signaling is reduced. The resistance of these supporting cells does not mean there are no consequences of reduced IGF-1 signaling in the time of glutamate stress, as both astrocytes and endothelial cells show signs of oxidative stress in the hours following exposure. The combination of neuron, astrocyte, and endothelial cells in culture afforded astrocytic protection from IGF-1 deficiency/glutamate stress but also highlighted a failure in the ability of astrocytes to protect the nearby neurons. Thus, the age-related loss of IGF-1 likely impairs the function and vitality of the entire neurovascular unit by differentially exerting stressors on neurons, endothelial cells, and astrocytes.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The animal study was reviewed and approved by Institutional Animal Use and Care Committee of the University of Mississippi.

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AUTHOR CONTRIBUTIONS

CH and NA were responsible for conceptualization, investigation, project administration, and formal statistical analyses. CH, JM, and NA developed and designed methodology. BA, AV, and NA performed blinded image analyses and curated data. CH prepared visualizations. CH, BA, and NA prepared the original draft. All authors reviewed and edited the manuscript.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnagi.2021.751304/full#supplementary-material>

Supplementary Figure 1 | Additional assessments of total cell number. **(A)** Average number of neurons per field 24 h after treatment. One-way ANOVA revealed no difference between means. **(B–D)** Astrocytes were pre-treated with 5 μ M PPP for 24 h prior to combining with endothelial cells and neurons for a triple culture system. Average number of cells per field in the neurons **(B)**, astrocytes **(C)**, and endothelial cells **(D)** cultures after stimulation with 100 μ M glutamate for 1 h ($n = 5–8$ wells/group). One-way ANOVA revealed no difference between means, and the p value of the ANOVA is listed for each. All data are presented as mean \pm SEM. **(E)** Average astrocyte viability 24 h after 100 μ M glutamate stimulation. Prior to stimulation, the cells were pre-treated with vehicle, 5 μ M NVP-ADW742, or 100 nM IGF-1 for 24 h ($n = 6–8$ wells/group). **(F)** Average astrocyte cell count following treatment with vehicle, 5 μ M NVP-ADW742, or 100 nM IGF-1 ($n = 6–8$ wells/group).

Supplementary Figure 2 | Astrocyte ROS and mitochondrial stress. **(A)** Average ROS levels 5 h after pure cultures of astrocytes treated with 0.5 μ M NVP-ADW742 and 100 μ M glutamate. Data were not normally distributed, so a Tukey's *post hoc* comparison was utilized following One-way ANOVA; * indicates a significant difference compared to vehicle ($n = 10$ wells/group). **(B)** Average oxygen consumption rate (y axis) and extracellular acidification rate (x axis) in astrocytes treated with vehicle control, 100 μ M glutamate, or 100 μ M glutamate + 0.5 μ M PPP for one hour. Measurements occurred 5 h after treatment. Maximal respiration is grouped in the top right 3 points, and basal respiration is grouped in the bottom left 3 points. A one-way ANOVA was used for each comparison of maximal OCR, maximal ECAR, basal OCR, and basal ECAR. *Post hoc* Dunnett's test vs vehicle control was used when relevant. * indicates significant difference in OCR, and # indicates significant difference in ECAR. All data are presented as mean \pm SEM, $n = 7–8$ wells/group.

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Treadmill Exercise During Cerebral Hypoperfusion Has Only Limited Effects on Cognitive Function in Middle-Aged Subcortical Ischemic Vascular Dementia Mice

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Clinical and basic research suggests that exercise is a safe behavioral intervention and is effective for improving cognitive function in cerebrovascular diseases, including subcortical ischemic vascular dementia (SIVD). However, most of the basic research uses young animals to assess the effects of exercise, although SIVD is an age-related disease. In this study, therefore, we used middle-aged mice to examine how treadmill exercise changes the cognitive function of SIVD mice. As a mouse model of SIVD, prolonged cerebral hypoperfusion was induced in 8-month-old male C57BL/6J mice by bilateral common carotid artery stenosis. A week later, the mice were randomly divided into two groups: a group that received 6-week treadmill exercise and a sedentary group for observation. After subjecting the mice to multiple behavioral tests (Y-maze, novel object recognition, and Morris water maze tests), the treadmill exercise training was shown to only be effective in ameliorating cognitive decline in the Y-maze test. We previously demonstrated that the same regimen of treadmill exercise was effective in young hypoperfused-SIVD mice for all three cognitive tests. Therefore, our study may indicate that treadmill exercise during cerebral hypoperfusion has only limited effects on cognitive function in aging populations.

Keywords: aging, behavior, cognitive function, mouse, subcortical ischemic vascular dementia, treadmill exercise

INTRODUCTION

Physical activity helps to promote and maintain brain health, including memory retention and cognitive performance. Research has shown that increased physical activity both prevents and ameliorates multiple brain diseases, including subcortical ischemic vascular dementia (SIVD). SIVD is the most common form of vascular cognitive impairment and dementia (VCID) (Erkinjuntti et al., 2000a,b). SIVD patients typically suffer from peri-ventricular white matter

degeneration that leads to stepwise development of neurological deficits and loss of executive function, such as difficulties with working memory (Roman et al., 2002; Roh and Lee, 2014; Prins and Scheltens, 2015; Wallin et al., 2018). Although SIVD is expected to become more prevalent as the population ages, to date, there are no clinically effective drugs.

There is an emerging body of evidence suggesting that patients with mild VCID perform better at cognitive functioning tests after aerobic exercise (Liu-Ambrose et al., 2016). However, there is a lack of basic data in experimental models that supports the efficacy of exercise for preventing SIVD/VCID progression. Recently, we reported that treadmill exercise improved cognitive function and increased the number of oligodendrocyte precursor cells (OPCs) in the white matter of hypoperfused-SIVD mice (Ohtomo et al., 2020). Nevertheless, an important question remains; although aging is a major risk factor for SIVD, and the majority of SIVD patients are elderly, no reports to date have examined the efficacy of exercise in aged SIVD mice. Therefore, in this study, we asked whether treadmill exercise alleviates cognitive decline by cerebral hypoperfusion in 8-months old mice.

MATERIALS AND METHODS

Overall Experimental Design

In this mouse model of SIVD, cognitive function (working memory, cognitive memory, and special learning) is known to decline in 4 weeks, presumably due to rarefaction of the white matter (Shibata et al., 2004, 2007; Ihara and Tomimoto, 2011; Yang et al., 2016; Ohtomo et al., 2020; Takase et al., 2021). Our intervention was a 6-week treadmill training which, according to our hypothesis, would slow cognitive decline and ameliorate the deterioration in performance over time. We hypothesized this because this treadmill protocol was shown to be effective in multiple CNS disease models, including young hypoperfused-SIVD mice (Ohtomo et al., 2020; Kinoshita et al., 2021). In this study, we conducted two experiments; Experiment 1 was for Y-maze test, and Experiment 2 was for novel object recognition test (NORT) and Morris water maze test (Figure 1). In both experiments, after the Sham or bilateral common carotid artery stenosis (BCAS) surgery on day 1, treadmill training occurred from day 8 to day 47 in the training group. Mice in the sedentary group were placed on the treadmill for an equivalent amount of time, but without running. In addition to the memory assessment, we also conducted immunohistological and biochemical analyses using mouse brain sections prepared from the mice in Experiment 1. For Experiment 1, 33 mice were used, and 6 mice (5 mice of the BCAS/Sedentary group and 1 mouse of the BCAS/Treadmill group) were excluded due to death or unhealthy conditions. For Experiment 2, 34 mice were used, and 6 mice were excluded due to death or unhealthy conditions.

Prolonged Cerebral Hypoperfusion Model (BCAS Mice)

All performed experiments followed an institutionally approved protocol in accordance with the National Institutes of Health

Guide for the Care and Use of Laboratory Animals and the law for the humane treatment and management of animals. Eight-month-old male C57BL/6J mice (Jackson Laboratory, United States) were housed in a specific pathogen-free conditioned 12-hour light/dark cycle room with free access to food and water throughout the experiment. After a week-long habituation period in our animal facility, mice were randomly selected to have a microcoil (0.18 mm diameter; Samini, Japan) applied to bilateral common carotid arteries for the induction of chronic cerebral hypoperfusion. The surgical procedure was performed as previously described (Shibata et al., 2004; Ohtomo et al., 2020). A control group received sham operation (after exposing bilateral common carotid arteries, the cervical incision was closed without coil application).

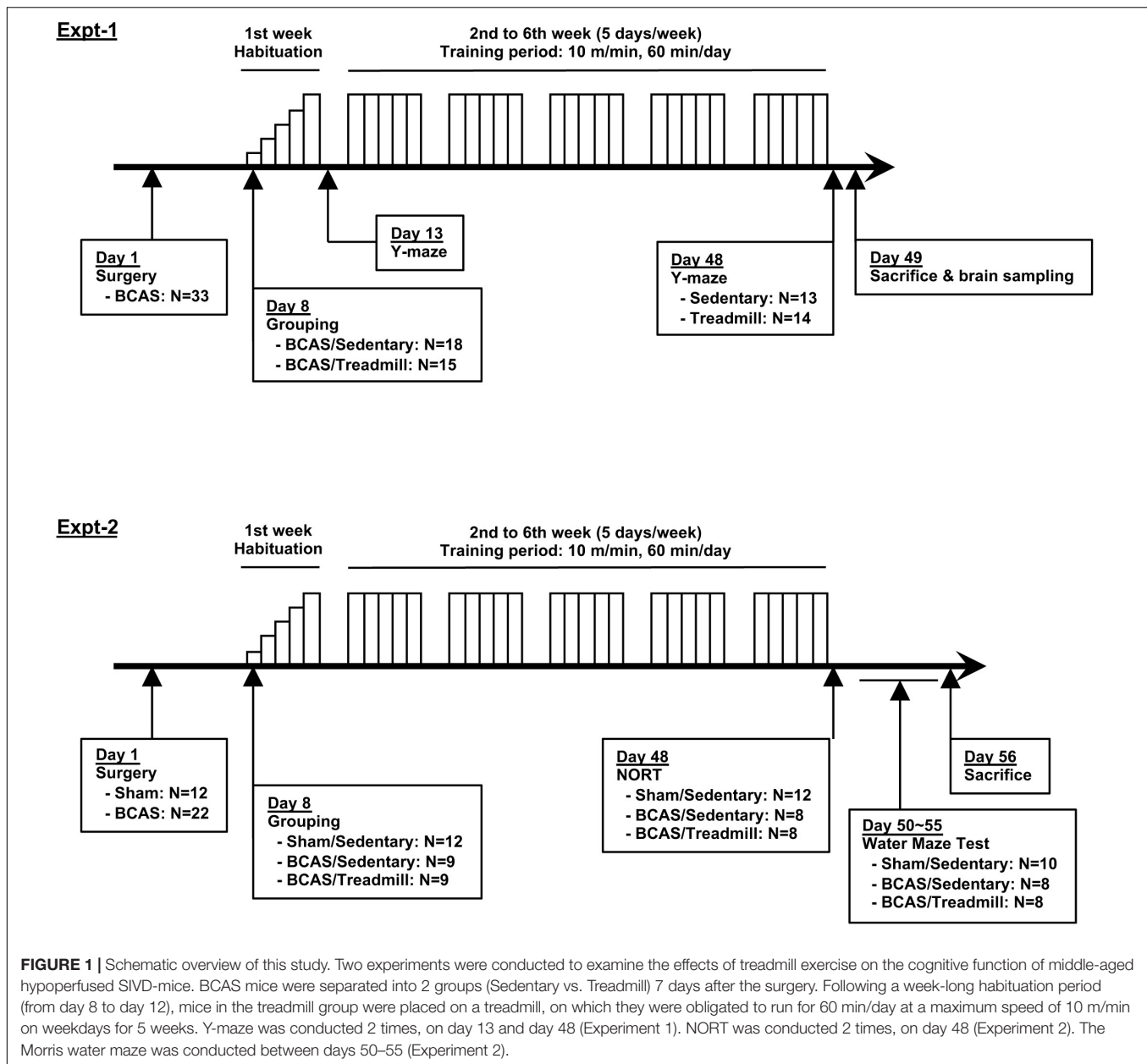
Treadmill Training Protocol

Sham or BCAS-operated mice were randomly divided into a sedentary group and a training group. The training group was forced to run on a treadmill device (Exer 3/6 Treadmill, Columbus Instruments, United States) for 6 weeks (in the early afternoon, 5 days a week on weekdays). Running speed suitable for exercise was determined from our previous studies that demonstrated protective effects against brain injury (Ohtomo et al., 2020; Kinoshita et al., 2021), beginning with a speed of 2 m/min, with an increase of 2 m/min every 2 min, until a maximum speed of 10 m/min was reached. The length of the total exercise was initially set at 20 min and increased daily by 10 min up to 60 min for familiarization, as shown in Figure 1. The sedentary group was placed on the device without running for the same amount of time as the training group. Mice were always placed in the same lane throughout the experiment.

Evaluation of the Muscle Fibers of Gastrocnemius Muscles

On day 49, the right gastrocnemius muscle was excised from randomly selected BCAS/Sedentary ($N = 10$) or BCAS/Treadmill ($N = 10$) mice from Experiment 1, under anesthesia with isoflurane before transcardial perfusion. Muscles perpendicularly embedded to Cryomold (Sakura Finetek USA, United States) with Tissue-Tek® (Sakura Finetek USA) were quickly frozen by liquid nitrogen and kept at -80°C until use. Five samples were randomly selected from each study group, and 10 μm -thick frozen sections were made using cryostat CM1520 (Leica, Germany). Sections were then fixed with -20°C methanol for 10 min and stained with Hematoxylin 2 (Thermo Fisher Scientific) and Eosin-Y (Thermo Fisher Scientific) according to the manufacturer's instructions. Stained sections were observed with ECLIPSE Ti-S (Nikon, Japan) and scanned with Retiga™ 2000R Fast 1394 Digital Camera (QImaging, Canada). Then, a cross-sectional area of 50 muscle fibers per mouse was measured by using ImageJ¹. To compare the area distribution of the muscle fibers between the groups, a histogram was drawn using PRISM® 7 (GraphPad Software, United States).

¹<https://imagej.nih.gov/>



Y-Maze

After the first and sixth week of exercise (day 13 and day 48), mice were tested for spontaneous alternation behavior with Y-maze between 6 AM and 9 AM. Our approach in conducting the Y-maze test multiple times in the same mouse was justified by the papers that the retention memory within the Y-maze task may not last longer than a few hours (Dellu et al., 2000; Fu et al., 2017). Each mouse was placed in the arm of symmetrical Y-maze apparatus to freely explore the maze for 8 min. The task was videotaped, and the sequence and the total number of arm entries were manually recorded later in a blinded manner. An arm entry was confirmed when bilateral hind paws were placed inside the arm. Arms were washed with 70% Ethanol between each test. Percentage of alternation was calculated as follows: number of

triads containing entries into 3 different arms/(total number of arms entered -2) $\times 100$.

NORT

Mice were tested for recognition memory by NORT between 8 AM and 1 PM on day 48. At first, the mice were habituated in an empty cage for 10 min before training. During training, they experienced 5-min exposure to 2 identical plastic blocks 2 times. After 30 min, they were then presented with 2 different plastic blocks (one of the original blocks replaced by a new block) for 5 min. The mouse behavior was videotaped and manually assessed in a blinded manner. Object recognition was scored by the total time spent either sniffing or touching the object. The performance of recognition memory was evaluated by the ratio

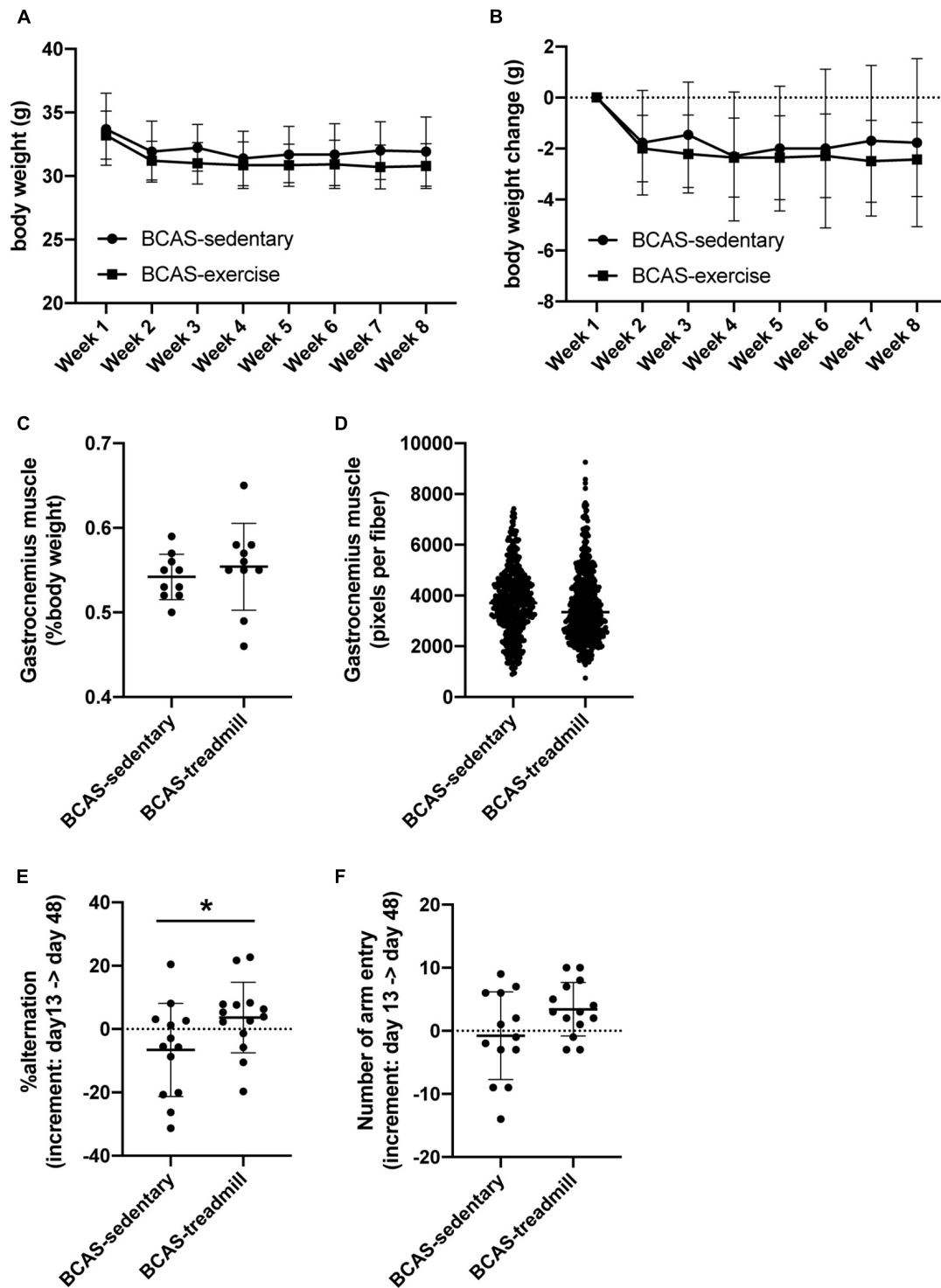


FIGURE 2 | Result of cognitive function test from Experiment 1. (A,B) Body weight transition of mice (A), and temporal body weight change of mice (B). There was no significant difference between the 2 groups (BCAS/Sedentary vs. BCAS/Treadmill). Data are expressed as mean \pm SD. (C,D) Weight of gastrocnemius muscle (percentage of body weight) (C) and distribution of the area of muscle fibers (D) at day 49. There was no significant difference between the 2 groups (BCAS/Sedentary vs. BCAS/Treadmill). Data are expressed as mean \pm SD. (E,F) Change in alternation (an index of working memory) in Y-maze test before and after the training period (E) and change in the number of arm entries in Y-maze test before and after the training period (F). There was a significant difference in the increment of alteration between the 2 groups (BCAS/Sedentary vs. BCAS/Treadmill) (* $P < 0.05$, Mann-Whitney U test). Data are mean \pm SD.

of the time spent on the new object to the total time spent on 2 objects (Ohtomo et al., 2020). Five mice were excluded from analysis due to lack of interest toward the objects (e.g., those mice did not sniff or touch the 2 objects).

Morris Water Maze Test

On days 50–55, mice were tested with Morris water maze to evaluate spatial learning and memory as previously described (Ohtomo et al., 2020). In brief, mice had to find a hidden (2 cm under the water surface) escape platform (diameter: 11.5 cm) which was placed in the northeastern quadrant of a circular pool (diameter: 110 cm) filled with opacified water kept at 25°C. In the acquisition phase, mice performed 4 trials per day from a random starting position for 5 consecutive days, with a maximum trial duration of 60 s. To test spatial memory, the platform was removed, and 60-second probe trials were performed on day 54 (4 h after the last acquisition test) and day 55 (4 h after the last acquisition test). Escape latency to find the hidden platform and number of entries into quadrants were examined using ANY-maze video tracking software (Stoelting Co., United States). Due to passive floating behavior in the water, 2 mice were excluded from analysis (one from the BCAS/Sedentary, and the other from the BCAS/Treadmill group). For the Sham/Sedentary group, we randomly picked 10 mice out of 12 Sham/Sedentary mice for the Morris water maze test due to time scheduling issues.

Western Blot Analysis

Mouse brains were removed following transcardial perfusion with 40 mL PBS (0.1 M), quickly frozen in dry ice, and kept at -80°C until use. Frozen brains were thawed to a semi-frozen state, and three 2 mm-thick coronal sections were obtained from the forebrain 1–7 mm anterior to the confluence of sinuses using the mouse brain matrix slicing tool. The brain sections were placed in ice-cold PBS, and the meninges were removed, after which the corpus callosum and cerebral cortex were carefully removed under a microscope with a disposable micro knife (Fine Science Tools, Canada). Four times the amount of PBS with protease inhibitor cocktail (Cytoskeleton, United States) was promptly added to the samples and dismembrated by sonic dismembrator (ThermoFisher Scientific, United States) on ice. After adding Triton X-100 (Sigma-Aldrich, United States) to a final concentration of 1%, samples were frozen at -80°C and thawed on ice. Then, they were centrifuged at 10000 × *g* for 15 min at 4°C to remove cellular debris. Protein concentration was determined by the BCA assay (Thermo Fisher, United States). Collected samples were heated with equal amounts of LDS sample buffer (ThermoFisher Scientific) and sample reducing agent (ThermoFisher Scientific) at 70°C for 10 min. Each sample was equally (30 µg) loaded onto 4–12% Bis-Tris gels (ThermoFisher Scientific) for electrophoresis, followed by transfer to a nitrocellulose membrane (Thermo Fisher Scientific). Membranes were blocked in 5% skim milk (LabScientific, United States) and incubated overnight at 4°C with primary antibodies against PDGFRα (1:500, Sigma-Aldrich, United States), MBP (1:1000, Thermo Fisher Scientific), and β-actin (1:10000, Sigma-Aldrich). Then, membranes were processed with peroxidase-conjugated

secondary antibodies [1:1000 for anti-rabbit antibody (GE Healthcare, United States), and 1:2000 for anti-mouse antibody (GE Healthcare, United States)] and visualized by enhanced chemiluminescence (Thermo Fisher Scientific). Visualized bands were semi-quantified with ImageJ.

Immunohistochemistry

Five samples were randomly selected from each study group, and 20 µm-thick coronal sections (corresponding to the area 0–1 mm anterior to bregma) were prepared for immunostaining using cryostat CM1520. Frozen sections were then fixed with -20°C methanol for 10 min. After being washed with PBS containing 0.1% Triton X-100 for 5 min 3 times, they were incubated in PBS/3% BSA solution for 1 h at room temperature. Then, sections were incubated in PBS/3% BSA solution containing primary antibodies anti-PDGFRα (1:100, R&D systems, United States) or anti-nestin (1:100, Abcam, United States) at 4°C overnight. After being washed with PBS 3 times, they were incubated with secondary antibodies (1:1000, Jackson ImmunoResearch Laboratories) for 1 h at room temperature. Finally, the sections were washed three times with PBS, and covered with VECTASHIELD® mounting medium with DAPI (Vector Laboratories, United States). Stained sections were observed with ECLIPSE Ti-S and scanned with Retiga™ 2000R Fast 1394 Digital Camera. All brain sections were blinded to the examiner before the evaluation of fluorescence intensity. Immunofluorescence intensity of PDGFRα-positive cells inside the subventricular zone (SVZ) was calculated by ImageJ. Bilateral sides of two coronal sections cut from the area +0.5 ~ +1.0 mm to the bregma were evaluated, and the average of 4 areas was calculated per mouse. The same procedure was conducted for evaluating the fluorescence intensity of nestin-positive cells.

Statistics

Statistical analysis was performed with R version 3.4.0² and PRISM® 7. For body weight, data were first tested with two-way analysis of variance (ANOVA), followed by *post hoc* Sidak's multiple comparisons test. For the acquisition phase of the Morris water maze test, data were first tested with two-way ANOVA, followed by *post hoc* Tukey's multiple comparisons test. Data from the probe test were first analyzed with Brown-Forsythe ANOVA, followed by *post hoc* Dunnett's T3 multiple comparisons test. NORT data were first analyzed with one-way ANOVA, followed by *post hoc* Tukey's multiple comparisons test. Mann-Whitney *U* test or Welch's *t*-test was used for other analyses. All values were expressed as mean ± SD. *P*-values less than 0.05 were considered statistically significant.

RESULTS

To examine the effects of treadmill exercise on the cognitive function of middle-aged hypoperfused-SIVD mice, we conducted two sets of experiments (Figure 1). Hypoperfused-SIVD mice were prepared by subjecting mice to the BCAS operation.

²<https://www.r-project.org/>

The baseline of cerebral blood flow is lower in aged mice (22 months old), compared to young mice (2–3 months old), thus the decrease of cerebral blood flow by BCAS is less impacted in aged mice (Baik et al., 2021). However, at least in our system, the BCAS operation caused prolonged cerebral blood flow reduction in middle-aged mice (8 months old) (**Supplementary Figure 1**). In the first set of experiments, we prepared two groups (hypoperfused-SIVD mice without exercise treadmill and hypoperfused-SIVD mice with exercise treadmill) to investigate whether exercise treadmill during cerebral hypoperfusion mitigated cognitive decline (e.g., from day 13 to day 48 after BCAS) using Y-maze test. In addition, after the completion of the Y-maze experiment, gastrocnemius muscles and brains were sampled for biochemical analyses. In the second set of experiments, we prepared three groups (sham mice without exercise treadmill, hypoperfused-SIVD mice without exercise treadmill, and hypoperfused-SIVD mice with exercise) to examine whether the cognitive function of hypoperfused-SIVD mice with treadmill exercise would be comparable with that of sham mice without treadmill exercise. For both experiments, male C57BL/6J (8 months old) mice were subjected to the 6-week exercise after sham or BCAS operation. One week was spent on habituation exercise, where running speed began at 2 m/min and was increased by 2 m/min every 2 min until a maximum speed of 10 m/min was reached; the duration of the daily exercise was initially set at 20 min and was increased daily by 10 min up to 60 min. In the following 5 weeks, mice were then subjected to running exercise at a speed of 10 m/min for 60 min/day.

In the first set of experiments, both groups (hypoperfused-SIVD mice without exercise and hypoperfused-SIVD mice with exercise) lost body weight after BCAS operation, and over time during the treadmill exercise training, they did not gain body weight regardless of treadmill exercise (**Figures 2A,B**). In addition, there was no significant difference in the right gastrocnemius muscles between the two groups (**Figure 2C** for the percentage of gastrocnemius muscle weight and **Figure 2D** for the distribution of muscle fiber thickness). However, in the Y-maze tests, while hypoperfused-SIVD mice without treadmill exercise showed a decline in cognitive function from day 13 to day 48 after BCAS operation, hypoperfused-SIVD mice with treadmill exercise did not (**Figures 2E,F**).

In the second set of experiments, we assessed cognitive function with the Morris water maze test and NORT. In the Morris water maze test, there was no significant difference in the memory acquisition between the three groups (sham mice without exercise vs. hypoperfused-SIVD mice without exercise vs. hypoperfused-SIVD mice with exercise) (**Figure 3A**). In addition, in the probe tests at 4 and 24 h after the last session of acquisition trial, there were no differences in the memory retention performance between the groups (**Figures 3B–D**). In NORT, as expected, sham mice without exercise showed a preference for the novel object; however, hypoperfused-SIVD mice without exercise showed no preference between the familiar and novel objects (**Figure 3E**). On the other hand, there was no significant difference in the preference between sham mice without exercise and hypoperfused-SIVD mice with exercise (**Figure 3F**).

Finally, we examined whether white matter pathology would be affected by treadmill exercise, because white matter damage is one of the major characteristics of SIVD. Fluoro-myelin staining showed that there was no difference in myelin density in the corpus callosum between hypoperfused-SIVD mice with or without treadmill exercise (**Figure 4A**). Western blotting also confirmed that treadmill exercise did not change the myelin density of corpus callosum, assessed by the level of MBP, which is a major component of the myelin sheath (**Figure 4B**). After white matter damage, compensatory responses would be activated, and OPCs play an important role in increasing the number of myelin-producing oligodendrocytes. However, the number of OPCs was not affected by the treadmill exercise in middle-aged hypoperfused-SIVD mice (**Figures 4B,C**). In addition, there was no significant difference in the Nestin-positive cells, which can differentiate into both neurons and OPCs in the adult brain after brain injury (**Figure 4C**). And finally, we checked if the middle-aged SIVD mice had a significant infarction in the cortical area. It is now well accepted that prolonged cerebral hypoperfusion by BCAS does not cause infarcts in young mice, but little is known about this point in middle-aged mice. But we confirmed no ischemic infarction in the cortex by hematoxylin-eosin staining with randomly selected 10 brain samples from the SIVD mice (**Supplementary Figure 2**).

DISCUSSION

Our current study demonstrated that in middle-aged mice with prolonged cerebral hypoperfusion, treadmill exercise ameliorates the decline of working memory in the Y-maze test. On the other hand, treadmill exercise was limited in its supportive effects on cognitive function in the Morris water maze test and NORT. Previously, we showed that the same regimen of treadmill exercise suppressed cognitive decline, assessed by the three cognitive function tests (Y-maze, Morris water maze test, and NORT), in young mice with cerebral hypoperfusion (Ohtomo et al., 2020). Our previous study also showed that the treadmill exercise during cerebral hypoperfusion increased the number of OPCs within the SVZ region of young hypoperfused-SIVD mice (Ohtomo et al., 2020), although we did not confirm the positive effects of treadmill exercise on OPC function in this study. Therefore, together with our previous data (Ohtomo et al., 2020), our current study may suggest that the efficacy of treadmill exercise in cognitive function would decrease with age, but treadmill exercise still seems effective in supporting cognitive function even in aging populations. This is consistent with past studies that treadmill exercise supported cognitive function in both young and aged AD mice, but the efficacy was smaller in the aged groups (Choi et al., 2021).

An important aspect of our current study is the finding that treadmill exercise is less effective in middle-aged SIVD model mice. To the best of our knowledge, except for our previous report (Ohtomo et al., 2020), there are only two previous studies that evaluate the effectiveness of exercise in preventing/ameliorating cognitive decline caused by chronic cerebral hypoperfusion (Jiang et al., 2017; Lee et al., 2017). These two studies showed

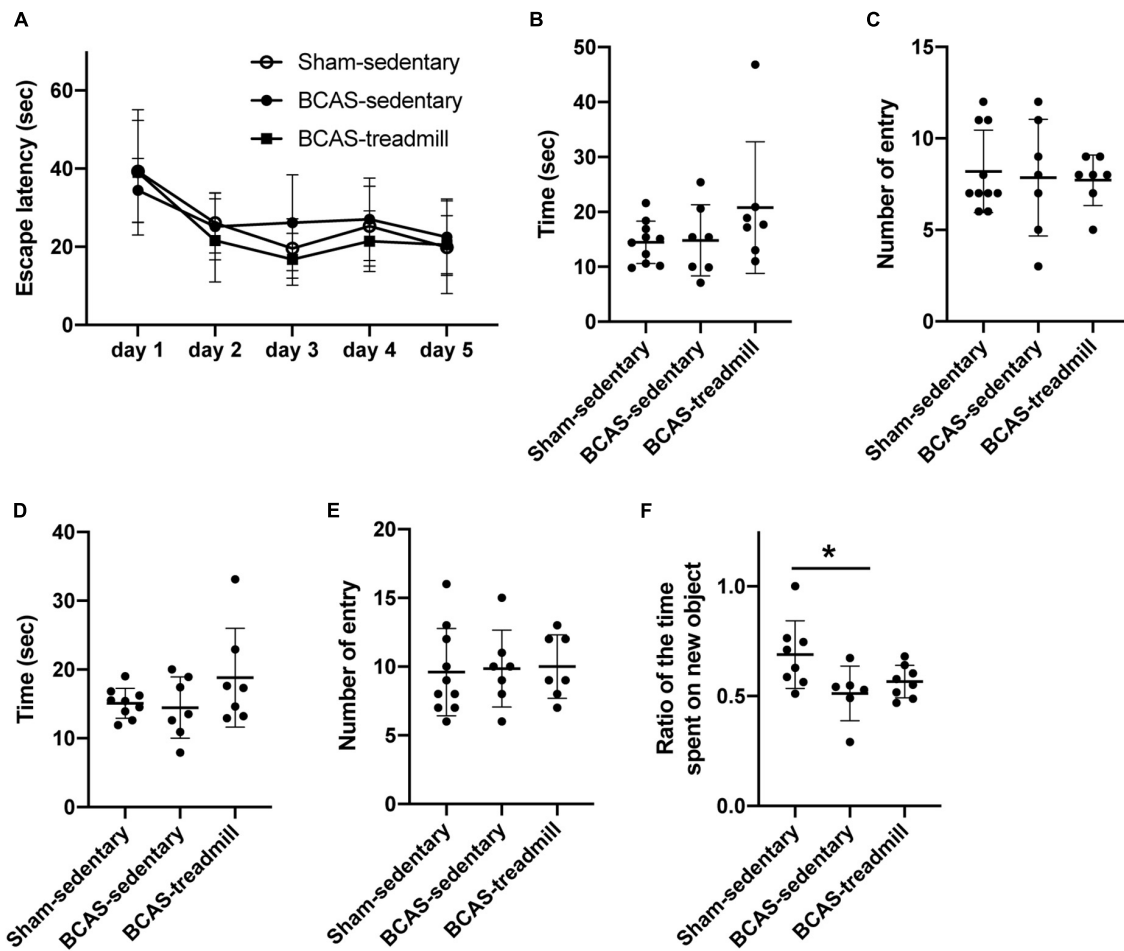


FIGURE 3 | Result of cognitive function test from Experiment 2. **(A)** During the acquisition phase of Morris water maze, no difference was observed in latencies to locate the hidden platform between the 3 groups (Sham/Sedentary vs. BCAS/Sedentary vs. BCAS/Treadmill). Data are mean \pm SD. **(B,C)** Result of the probe trial (4 h after the last acquisition session) in the Morris water maze test. There were no significant differences in both **(B)** number of entries into and **(C)** time spent in the quadrant which formerly contained a hidden platform in the acquisition phase. Data are mean \pm SD. **(D,E)** Result of the probe trial (24 h after the last acquisition session) in the Morris water maze test. There were no significant differences in both **(D)** number of entries into and **(E)** time spent in the quadrant which formerly contained a hidden platform in the acquisition phase. Data are mean \pm SD. **(F)** Compared to the Sham/Sedentary group, the BCAS/Sedentary group showed less preference to the novel object (* $p < 0.05$, one-way ANOVA followed by *post hoc* Tukey's test). However, there was no significant difference between Sham/Sedentary and BCAS/Treadmill groups. Data are mean \pm SD.

that treadmill exercise is effective in ameliorating cognitive dysfunction using young SIVD model rats. Our findings further add to this existing literature. The efficacy of treadmill exercise during cerebral hypoperfusion in cognitive function might be less effective in aging populations. This may be partly explained by the fact that compensative responses after brain injury, such as neurogenesis and oligodendrogenesis, are dampened by age (Miyamoto et al., 2013; Liang et al., 2016). While the treadmill exercise increased the number of OPCs within the SVZ region in young SIVD mice (Ohtomo et al., 2020), our current study did not confirm this effect in middle-aged mice. In addition, the body weight loss after cerebral hypoperfusion was not recovered by treadmill exercise in middle-aged mice (Figures 2A,B), although the same protocol of treadmill exercise helped young SIVD mice recover their body weight (Ohtomo et al., 2020). In stroke mice,

the body weight recovery was associated with an improvement of motor function (Desgeorges et al., 2017; Kinoshita et al., 2021), which may also explain the difference in the efficacy of treadmill exercise between young and middle-aged mice.

Although we have demonstrated that treadmill exercise ameliorates cognitive decline caused by cerebral hypoperfusion to some extent in middle-aged mice, there are some limitations and caveats in this study. First, we used only one protocol of treadmill exercise, which was shown to be effective for young mice (Ohtomo et al., 2020; Kinoshita et al., 2021). In general, in aged populations, the capability of movement and the total volume of muscle is lower compared to younger populations, so the exercise protocol for young mice may not be suitable for middle-aged mice. The optimization of the conditions of treadmill exercise for middle-aged (or even aged) mice would be necessary for

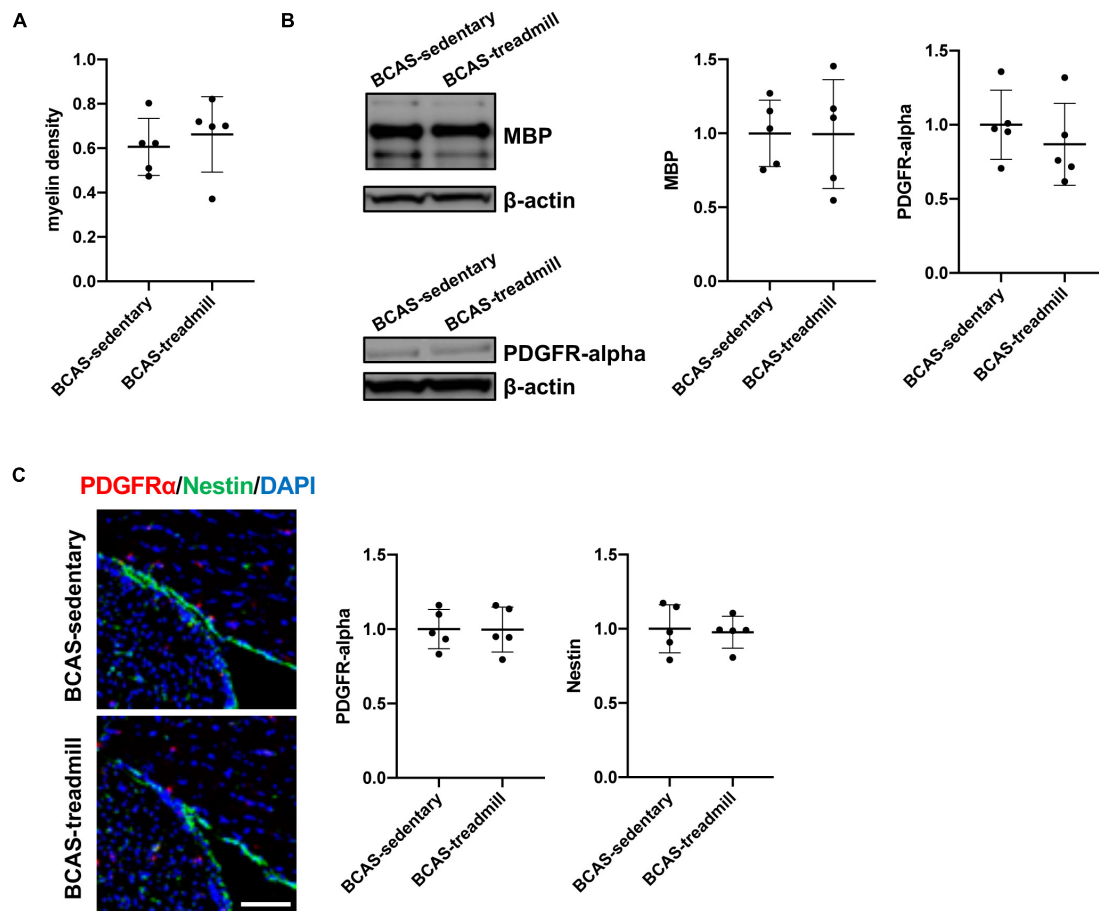


FIGURE 4 | (A) Result of myelin density of the corpus callosum, assessed by fluoromyelin staining. Myelin density was calculated based on the intensity of age-matched male C57/BL6J mice (average of 5 mice). There was no significant difference in myelin density between the 2 groups (BCAS/Sedentary vs. BCAS/Treadmill). Data are expressed as mean \pm SD. **(B)** Result of MBP and PDGFR α western blot using samples from the corpus callosum region. Data are expressed as mean \pm SD. **(C)** Immunohistochemistry showed that there were no significant differences in both the intensity of PDGFR α -positive cells and the intensity of Nestin-positive cells within the SVZ region. Data are expressed as mean \pm SD.

future studies to pursue exercise as a therapeutic option for SIVD and other CNS disease patients. Second, because we focused on the effect of treadmill exercise on middle-aged SIVD mice, our current study did not include a sham-exercise group in our experiments. However, considering that treadmill exercise improves cognitive function in older mice (Maejima et al., 2018), further studies would be needed to answer the question of whether the effect of treadmill exercise is ineffective in middle-aged mice or only in middle-aged SIVD mice in our system. Careful comparison of the effect of treadmill exercise between these two groups may also help us to find a therapeutic target for SIVD. Third, our current study does not address the potential mechanism behind the effects of exercise on cognitive function. In young mice, the activation of compensatory responses, such as an increase of OPC number within the SVZ region, may partially contribute (Ohtomo et al., 2020); however, no OPC activation was observed in middle-aged mice. The hippocampal region plays an important role in cognitive function, including spatial learning and memory, and our pilot study indicated that in the

SGZ region of hippocampus, treadmill exercise may increase the number of stem cells (data not shown). Because accumulating evidence now suggests the close relationship between exercise and the neuronal responses in the hippocampus (Moon et al., 2016; Liu and Nusslock, 2018; Lourenco et al., 2019; Baik et al., 2021), it would be helpful to examine the cellular and molecular signaling in the hippocampus after treadmill exercise in middle-aged SIVD mice in future studies. The final caveat of our study is that we used only male mice for our experiment. Since the sex-difference may alter the dynamics of systemic humoral factors as mentioned in the published literature (Herson et al., 2013; McMullan et al., 2016; Stanford et al., 2017), it will be important to determine if our results can be replicated in middle-aged female mice.

In conclusion, we have demonstrated that treadmill exercise has some effect in reducing cognitive decline in middle-aged mice with prolonged cerebral hypoperfusion that mimics the pathophysiology of SIVD. Exercise is a safe behavioral intervention and has the potential to be a non-pharmacological

therapy for several CNS diseases, including SIVD. Because polypharmacy among elderly patients has become a serious social issue around the world (Fried et al., 2014), future studies are warranted to pursue the therapeutic option of exercise as a non-pharmacological approach to decrease the cognitive decline in SIVD or other dementia patients.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author/s.

ETHICS STATEMENT

The animal study was reviewed and approved by the Massachusetts General Hospital Institutional Animal Care and Use Committee.

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AUTHOR CONTRIBUTIONS

RO, HI, KK, GH, and HT: collection of data. RO, KK, HI, and GO: data analysis. RO, KC, JL, EL, and KA: manuscript writing. RO, KC, GO, CW, HK, AI, JL, EL, and KA: interpretation. RO, GO, EL, and KA: conception and design. EL and KA: funding acquisition. All authors: final approval of manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

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Gradual Not Sudden Change: Multiple Sites of Functional Transition Across the Microvascular Bed

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In understanding the role of the neurovascular unit as both a biomarker and target for disease interventions, it is vital to appreciate how the function of different components of this unit change along the vascular tree. The cells of the neurovascular unit together perform an array of vital functions, protecting the brain from circulating toxins and infection, while providing nutrients and clearing away waste products. To do so, the brain's microvasculature dilates to direct energy substrates to active neurons, regulates access to circulating immune cells, and promotes angiogenesis in response to decreased blood supply, as well as pulsating to help clear waste products and maintain the oxygen supply. Different parts of the cerebrovascular tree contribute differently to various aspects of these functions, and previously, it has been assumed that there are discrete types of vessel along the vascular network that mediate different functions. Another option, however, is that the multiple transitions in function that occur across the vascular network do so at many locations, such that vascular function changes gradually, rather than in sharp steps between clearly distinct vessel types. Here, by reference to new data as well as by reviewing historical and recent literature, we argue that this latter scenario is likely the case and that vascular function gradually changes across the network without clear transition points between arteriole, precapillary arteriole and capillary. This is because classically localized functions are in fact performed by wide swathes of the vasculature, and different functional markers start and stop being expressed at different points along the vascular tree. Furthermore, vascular branch points show alterations in their mural cell morphology that suggest functional specializations irrespective of their position within the network. Together this work emphasizes the need for studies to consider where transitions of different functions occur, and the importance of defining these locations, in order to better understand the vascular network and how to target it to treat disease.

Keywords: neurovascular, pericyte, brain, mural cell, arteriole, capillary, endothelial cell

INTRODUCTION

The human brain has an extensive vascular network essential for supplying the brain's rich energy demands. Oxygen, glucose and other nutrients and signaling molecules are sent to the brain *via* finely regulated cerebral blood flow which is delivered through arteries, arterioles and capillaries. Deoxygenated blood and waste products are then removed *via* capillaries, venules and veins. In the neocortex, large pial vessels run along the surface then dive and penetrate the brain, before branching into smaller vessels to form the dense capillary network (Duvernoy et al., 1981). Pial and parenchymal arterioles and capillaries respond to neuronal nutrient and oxygen demand by dilating and contracting (Iadecola, 2017) to alter blood flow locally, a process called neurovascular coupling (NVC). NVC is controlled by the neurovascular unit (NVU), formed by vascular cells (mural cells - smooth muscle cells and pericytes, and endothelial cells), astrocytes and neurons. In addition to modulating contractile tone and blood flow, the NVU is also fundamental in regulating blood brain barrier (BBB) permeability and nutrient delivery (Hawkins and Davis, 2005; Iadecola, 2017; Sweeney et al., 2018), as well as helping to coordinate the brain's immune response by restricting leukocyte invasion into the tissue.

Different parts of the vascular tree contribute differentially to these various functions, but in many cases, it remains unclear how function changes along the vessel bed. This is partly because of variability in nomenclature and definitions across studies, and partly because of an understandable, but ultimately misleading, tendency to oversimplify the manner in which these functional transitions occur. It is important to better understand these functional transitions, however, in order to identify which cells and vessels need to be targeted to manipulate these various processes therapeutically in conditions such as Alzheimer's disease or stroke.

Here we use existing literature and novel data to consider which components of the vasculature mediate different functions, to better understand functional transitions across the vascular network. We suggest that evidence indicates multiple transition points at different positions in the vascular network, and therefore to gradual, heterogeneous changes in vascular function, rather than sudden transition points between distinct vessel types.

Transitions in Anatomy Across the Cerebral Microvascular Bed

Across the vascular bed, from arteriole to capillary to venule, the diameter of vessels first decreases then increases, and the morphology of their mural cells (smooth muscle cells or pericytes) abluminal to the endothelial tube changes. Penetrating arteries and arterioles are typically between 15 and 40 μm in diameter and are composed of an inner layer of endothelial cells with abundant caveolae. They contain an internal elastic lamina and therefore express elastin in the vessel wall, which can be detected from its binding to the dye Alexa 633 hydrazide (Shen et al., 2012). At the surface, they often have 2-3 outer layers of closely packed ring-shaped smooth muscle cells (SMCs), which are electrically coupled *via* gap

junctions (Chow et al., 2020; Garcia and Longden, 2020) and strongly express smooth muscle alpha actin (αSMA). At the pial surface, the blood vessel is separated from the pial membrane by Virchow-Robin space, which narrows after the arteriole enters the brain through a "pial funnel" (Gao et al., 2015). Downstream from this region, SMCs form only one layer before transitioning to a pericyte morphology. Grant et al. (2019) report the first cells with "bump on a log" morphology as always occurring beyond the first branch from the arteriole, and most reports consider the arterioles to be covered with annular SMCs (e.g., Hamilton et al., 2010; Grant et al., 2019; Kisler et al., 2017). However, even near the pial surface, mural cells' morphology changes from a simple annulus to gaining a distinct soma and processes. While these cells are often still termed smooth muscle cells, historically this transitional form would be called a pericyte (Zimmermann, 1923; Attwell et al., 2016).

At the first branch point from penetrating arterioles are precapillary sphincters: contractile mural cells encircling a narrow section of vessel between the penetrating arteriole and first order capillary (Chambers and Zweifach, 1946; Grubb et al., 2020). The first branches off the penetrating arteriole are often termed precapillary arterioles or first order capillaries (where penetrating arterioles are branch order 0). These vessels tend to have a diameter of between 5-15 μm , and are enwrapped with mural cells that express αSMA and have a protruding, distinct soma, and processes that cover the vessel to a large degree, recently termed "ensheathing pericytes" (Hartmann et al., 2015; Grant et al., 2019; but see Hill et al., 2015; Attwell et al., 2016). These first branches are often termed the precapillary arteriole or transitional zone, though these terms sometimes are used to mean only the first vessel branching off the penetrating arteriole (e.g., Thakore et al., 2021), and other times include the 2nd and 3rd branches with larger pericyte coverage and αSMA actin expression (Gonzales et al., 2020).

Downstream of these vessels, after 1-3 branches from the diving arteriole (Grant et al., 2019) pericytes become less dense and their processes cover less of the vessel as branch order increases and vessel diameter decreases (range 3-9 μm). These capillary pericytes have been reported to have low (Bandopadhyay et al., 2001) or absent (Nehls and Drenckhahn, 1991; Grant et al., 2019) expression of αSMA and have been termed mesh and thin strand pericytes (Hartmann et al., 2015). Latterly these have been combined into a single category (capillary pericytes), due to the difficulty in distinguishing a transition point between these two morphologies (Grant et al., 2019). These capillaries with low or absent αSMA expression and low pericyte coverage are often termed mid-capillaries.

In post-capillary venules, pericytes have a stellate morphology, becoming less extended with thicker, more radial processes (Joyce et al., 1985; Hartmann et al., 2015). These post-capillary venules branch to form large venules ($> 50 \mu\text{m}$), which have weak αSMA labeling, indicating limited contractile potential (Hill et al., 2015). On larger venules, SMCs are circumferential but are less compact and show a less complete coverage of the vasculature and a leaf-like instead of banded appearance compared to arteriolar SMCs (Hill et al., 2015).

Therefore, the anatomy of the vessels and mural cells of the cerebral vasculature, as commonly reported, is used to define the following distinct vascular segments: pial arterioles, penetrating or diving arterioles, precapillary sphincters, pre-capillaries, also termed transitional or low branch order capillaries, mid-capillaries, post-capillaries and venules. These have been recently condensed into 4 categories, arterioles, a transitional zone, capillaries and venules, with distinct mural cells on each (Hartmann et al., 2021a). In some ways this categorization is a helpful way to discuss features of different parts of the vascular bed, as the recent Hartmann et al. review does in a clear and informative manner. Our argument, however, is that it is important to recognize that these categories are overly simplistic because they are superimposed on more gradually changing anatomy and function across the vascular bed. For example, existing literature shows that mural cell morphologies transition gradually not abruptly along the network. This concept was first discussed by Zimmermann (1923) and is evident from more recent work which shows that there is a continuum of cell lengths seen from SMCs through ensheathing to mid-capillary pericytes (Grant et al., 2019). Furthermore, the shift in vessel coverage of pericyte processes argued to mark a boundary between ensheathing and mesh pericytes has a different distribution across branch orders than that of α SMA expression (Grant et al., 2019), i.e., α SMA transitions occur at largely different locations than do transitions in pericyte coverage of the vessel.

Neurovascular Coupling Differences Across the Cerebral Microvascular Bed

This heterogeneity of anatomy across the vascular tree presumably reflects the varying functional roles performed by different vessels. Anatomical transitions have been best mapped onto functional changes in terms of dilatory capacity and involvement in neurovascular coupling (i.e., dilation in response to increased neuronal activity). Disease affects different parts of the vasculature in specific ways, making it important to understand how and where such functional transitions occur to be able to intervene successfully to target pathological mechanisms.

Classically, neuronal activity causes local pial and penetrating arterioles to dilate, causing an increase in blood flow to active brain regions (Attwell et al., 2010). However, different components of the vasculature are now known to play distinct roles in generating this response. Mid-capillary regions, being closer on average than any other vessel segment to most neurons, likely first detect neuronal activity, either *via* the accompanying increase in extracellular potassium concentration (Longden et al., 2017) and/or *via* production of vasoactive signaling molecules such as prostaglandin E2 (Hall et al., 2014; Mishra et al., 2016) or EETs (Mishra et al., 2016; Zhang et al., 2021). This signal then spreads upstream from the capillary bed, causing dilation of upstream low branching order capillaries and arterioles (Hall et al., 2014; Rungta et al., 2018), *via* spread of hyperpolarizing currents carried by inward rectifying potassium channels (Zhang et al., 2011; Longden and Nelson, 2015). Whether mid-capillaries constrict and dilate in response to modulation of neuronal

activity remains somewhat controversial. Some studies report that they do not (Hill et al., 2015), but a growing body of evidence supports their contractility (Hall et al., 2014; Kisler et al., 2017; Hartmann et al., 2021b; Shaw et al., 2021), though this may be *via* a different mechanism than in arterioles and lower branching order capillaries (Hartmann et al., 2021b).

The size, frequency and timing of responses to neuronal activity are often reported to vary across vascular compartments though not always in a consistent manner. Arteriole response magnitudes vary along their length, with surface sections of penetrating arterioles dilating more than deeper sections, probably as a result of the mechanical restriction imposed by brain tissue (Gao et al., 2015), though deeper arteriole sections have also been found to dilate more rapidly than shallower sections (Tian et al., 2010). In alpha-chloralose anaesthetized somatosensory cortex, Hall and colleagues found response frequency decreased as branching order increased (i.e., capillaries responded less frequently than arterioles; Hall et al., 2014), though first and second branch order capillaries showed larger and faster dilations than arterioles, a finding subsequently supported by other studies (Zhang et al., 2021). In the olfactory bulb, the region around the branch off the arteriole also dilated first, with slower responses up and downstream (Rungta et al., 2021). However, the same group found much more variable timings in neocortex of ketamine-medetomidine anaesthetized or awake mice, with capillaries or arterioles each being faster on some occasions (Rungta et al., 2021), and in both these reports, mid-capillary dilations were very small (though these data conflate responders and non-responders, unlike some other studies, Hall et al., 2014; Shaw et al., 2021). Precapillary sphincters, where studied, have often shown larger dilations, as a proportion of their diameter, than adjacent arterioles and capillaries (Grubb et al., 2020; Zambach et al., 2021).

In short, there are functional differences between neurovascular coupling responses along the vascular network, with mid-capillaries responding less frequently than upstream vessels to increases in local neuronal activity, and the fastest responses often, but not exclusively, observed in the first and second order capillaries. The physiological mechanisms underlying this heterogeneity in contractile responses are not wholly clear. Pericytes expressing α SMA seem likely to mediate more reliable, often larger and faster changes in vascular tone (Hill et al., 2015; Hartmann et al., 2021b), but as α SMA is commonly expressed in vessels up to and including the 3rd branch from the penetrating arteriole (Grant et al., 2019), variations in α SMA expression alone cannot explain the observed pattern of responses.

Indeed, several vasoactive pathways and mechanisms modulating the diameter of different vascular segments have been found to vary across the vascular network. Vessels beyond the penetrating arteriole were found to dilate *via* ATP-mediated increases in astrocytic calcium and prostaglandin action on EP4 receptors, while diving arterioles dilated *via* NMDA receptor mediated NO production (Mishra et al., 2016). Another functional transition point at the same location was also recently observed, as capillary endothelial cells were found to express MFSD2A, inhibiting caveolae formation,

while arteriolar endothelial cells showed much lower MFSD2A expression, instead having abundant caveolae that were necessary for intact neurovascular coupling (Andreone et al., 2017). Because MFSD2A-mediated reduction in caveolae formation reduces transcytosis to sustain BBB integrity (Ben-Zvi et al., 2014), this suggests a possible transition in both BBB function and neurovascular coupling mechanisms at the same arteriole-to-capillary point.

It is not just the mechanism of vasodilation that is sensitive to the position in the microvasculature: The integration of vascular signals across the vascular network is an area of increasing focus, with many recent papers shedding new insights on how vasoactive signals are propagated upstream of their detection in the mid-capillary bed. Potassium currents (Longden et al., 2017), nitric oxide (Kovacs-Oller et al., 2020), calcium and ATP signals (Thakore et al., 2021) have all recently been shown to be important in the transmission of vasodilatory signals and, by their nature, these signals span vascular segments and may be influenced by functional variations across these sections. While many studies have not explicitly investigated how different vascular segments affect signal transmission, Thakore et al. (2021) recently showed that ATP application or TRPA1 channel activation in the capillary bed produces slowly-propagating calcium signals through capillary endothelial cells that depend on ATP release from Panx1 channels, but are converted, by IK and SK channels in a transitional segment, into rapidly-propagating electrical signals that dilate upstream arterioles. A similarly propagating signal has been shown to be initiated in the capillary bed by PGE2 (Rosehart et al., 2021; this issue). In these papers, the transitional zone was defined as the first branch off the penetrating arteriole, and did not include lower branching order vessels that would be expected to express α SMA, and it is not clear whether these higher branching order vessels also show rapid IK or SK-mediated propagation of dilation.

Perhaps unsurprisingly given the varying vasoactive properties of different vascular segments, their function can be differentially affected by disease. In Alzheimer's disease, capillaries may mediate functionally significant decreases in cerebral blood flow, as capillaries but not arterioles were found to be constricted due to endogenous A β -mediated endothelin signaling to pericytes (Nortley et al., 2019), and to be plugged by neutrophils (Cruz Hernández et al., 2019). Similar effects may be seen after stroke, with capillaries constricting in brain slices during ischemia (Hall et al., 2014) and showing stalled flow and constrictions after reperfusion *in vivo* (Yemisci et al., 2009), though the largest constrictions may be seen in α SMA-expressing vessels rather than mid-capillary regions (Hill et al., 2015). Conversely, in mice carrying the main genetic risk factor for Alzheimer's disease, APOE4, capillary function was unaffected, but the large pial vessels were dysfunctional, with vasomotion and dilation frequencies reduced compared to APOE3-expressing controls (Bonnar et al., 2021). However, none of these studies define the precise location of these transitions in disease susceptibility.

To summarize, the contractile properties of the vascular tree, in terms of both capacity and timing of dilations, mechanisms mediating neurovascular signaling and integration, and the sensitivity of these processes to disease states all vary

depending on the position in the vascular network. In some cases (e.g., response frequency) function seems to gradually change along the network, while in others (e.g., mechanism of neurovascular coupling), there are more abrupt transitions of function, and in many cases the location of functional transitions remains unknown.

Oxygen Supply

The major function of the brain's blood supply is to provide it with nutrients. These include the energy substrates, oxygen and glucose, as well as nucleosides and amino acids needed for mRNA and protein synthesis. Classically, the dense capillary bed has been thought of as the site of nutrient exchange, but actually measuring where nutrient delivery occurs is not straightforward.

Improved two-photon phosphorescence lifetime imaging of oxygen probes has recently provided insights into the concentration gradients of oxygen across the microvasculature in mouse cortex, and therefore where most oxygen delivery likely takes place. In anaesthetized cortex, arterioles were found to be responsible for 50% of the extracted oxygen (Sakadžić et al., 2014), with capillaries (here defined as vessels two branches downstream of the penetrating arteriole) appearing to provide a reserve for oxygen delivery capacity, taking on a larger role during hypercapnia and, as suggested by modeling, when oxygen consumption rates increased. Indeed, in more recent studies in awake mice (therefore with a larger basal oxygen consumption rate), less oxygen was found to be delivered by the arterioles (34%), with the majority of the remainder expected to be delivered by the capillary bed (Li et al., 2019).

The factors driving oxygen delivery by these different vessels include red blood cell linear density, speed, local oxygen concentration gradients, and intravascular resistance to oxygen diffusion. These differ between cortical layers (Lyons et al., 2016; Schmid et al., 2017; Li et al., 2019) and are differently variable within these layers. For example, red blood cell (RBC) flow was lowest and more homogenous between capillaries and, correspondingly, oxygen extraction was largest in deep cortical layers, with faster flow and less oxygen extraction in superficial capillaries (Schmid et al., 2017; Li et al., 2019). This was in part because blood pressure dropped more in the diving arteriole when feeding deeper layers (Schmid et al., 2017). Furthermore, from arterial to venous capillaries, oxygenation becomes increasingly variable as an increasingly clear relationship between RBC flow and pO₂ emerges [more O₂ having been extracted from capillaries with slowly moving RBCs (Li et al., 2019)]. Thus, position in the vascular network in terms of both tissue depth and branching order affects how blood delivers oxygen to the tissue.

Nutrient Supply and the BBB

Unlike oxygen, which can diffuse through cell membranes, other nutrients are supplied to the brain *via* specialist transporters which allow their passage across the BBB. The conventional view is that arterioles and arteries control the blood supply to the tissue but do not participate in the exchange of nutrients in the brain, which is instead mediated by capillaries and post-capillary venules (Yuan and Rigor, 2010). The BBB strictly controls the

influx and efflux of molecules across its endothelial layer largely by active transport. In addition to being vital for neuronal function, these proteins can be targeted for drug development to allow drugs to access the brain.

Little work has studied where in the vascular network these transporters are most active, though a recent RNA Seq study showed them to be predominantly expressed in capillary and venule mural cells, in contrast to transcription factors which were overrepresented at the arterial end of the vasculature (Vanlandewijck et al., 2018). A key question to be addressed in future, is whether this myriad of different transporters and transcription factors transition in similar or different locations across the vascular network.

The BBB and Immune Regulation

The BBB is also a key site of regulation of immune access to the brain, which again varies across the vascular tree. Unlike arteriolar and capillary endothelial cells, post-capillary venular endothelial cells are connected by adherens rather than tight junctions, meaning they are more leaky than the capillaries which feed them (Rous and Smith, 1931; Poher and Sessa, 2014). The basement membrane that surrounds them also has a different structure, being lamellar instead of homogenous (Braverman, 1989). Leukocyte infiltration into the surrounding tissue primarily happens in this region, leukocytes migrating through gaps between pericytes and thus being regulated by pericyte morphology (Proebstl et al., 2012). Pericyte contractility may feed into this process, as pericyte relaxation (not contraction) widens the gaps between cells, facilitating leukocyte infiltration (Wang et al., 2012). Once in the parenchyma, leukocytes migrate along NG2-expressing pericytes, which release factors to support their migration (Stark et al., 2013). Thus, an important transition in vascular immune function appears to occur between NG2-positive pericytes on capillaries and the NG2-negative pericytes on veins, though whether this happens to the same extent in brain as in peripheral tissues remains unknown. In addition to regulation of immune cell entry, the vasculature can also transport cytokines into the brain, as well as respond to circulating factors with parenchymal production of cytokines and chemokines (Banks et al., 2009; Rustenhoven et al., 2017), but how this varies across the vascular network is also unknown.

More studies into the site of immune regulation in the brain would be valuable, however, as, perhaps unsurprisingly, these pathways of immune regulation by pericytes and endothelial cells are important for understanding and treating disease. Pericyte damage and increased permeability of the brain vasculature to plasma proteins and immune cells are hallmarks of Alzheimer's disease, multiple sclerosis, and stroke (Daneman, 2012), where infiltrating immune cells have also been shown to induce vascular dysfunction, demyelination, axonal damage, and neurodegeneration (Hochmeister et al., 2006; Ryu et al., 2015). Understanding where in the network structural and functional contributions of the vasculature to the regulation of neural inflammation occur may thus enable us to understand which cells should be best targeted therapeutically.

Vasomotion

As well as nutrients being delivered across the BBB, the vasculature also plays an important role in the removal of potentially neurotoxic substances from the surrounding brain tissue. Vasomotion, a phenomenon first reported in bat wing veins (Jones, 1853), is a spontaneous low frequency oscillation in blood vessel tone (typically centered near 0.1 Hz; Mayhew et al., 1996); and independent of heartbeat or respiration), which is present in vessels (particularly arteries) throughout the body (Aalkjær et al., 2011). This rhythmic pulsing of vessels is thought to contribute substantially to paravascular clearance (Aldea et al., 2019; van Veluw et al., 2020) and tissue oxygenation (Tsai and Intaglietta, 1989; Aalkjær et al., 2011; Thorn et al., 2011; Mateo et al., 2017). In arterioles, oscillations are generated within the vascular wall as a result of local SMC dilations and contractions, possibly *via* phospholipase C and phospholipase A₂ mediated cyclical release of calcium from IP₃-sensitive stores (Haddock et al., 2002). Capillaries have not been reported to show these low frequency oscillations in diameter, though RBC flow does show similar fluctuations which may be due to upstream diameter fluctuations (Colantuoni et al., 1994; Biswal and Hudetz, 1996). However, a functional transition in vasomotion of vessel diameters has not yet been pinpointed.

Given the importance of vasomotion as a driving force for paravascular tissue clearance, it is not surprisingly affected by vascular-degenerating diseases. In a mouse model of Alzheimer's disease, arterioles surrounded with A β showed impaired vasomotion when driven by neuronal activity, and showed reduced dextran clearance from the parenchyma (van Veluw et al., 2020). Mice carrying the main genetic risk factor for Alzheimer's disease, APOE4, also showed a reduction in pial arteriole vasomotion compared to APOE3 controls (Bonnar et al., 2021). Thus, vasomotion is important for regulating supply and clearance of substances to the brain, and is affected by disease. However, we do not yet know where in the vascular tree the transition point falls in which the vessels cease to show vasomotion.

Summary

Vascular anatomy, contractility, nutrient delivery, immune regulation and clearance all differ across the vascular bed. To a large extent, functions of the vascular network have been assumed to transition broadly with branching anatomy, from vasomotion pial and diving arterioles that show reliable, large neurovascular coupling responses, through fast responding pre-capillaries/low branching order capillaries, to less reliably dilating mid-capillaries. These mid-capillaries are thought to have a stronger role than upstream vessels in BBB regulation and nutrient transport, while further downstream, post-capillary venules regulate immune cell infiltration. While this framework no doubt captures the broad changes that occur over the vascular network, it is by no means clear that the vasculature can be usefully divided into such discrete divisions. Indeed some of the research discussed above points to the existence of additional functional transitions within vascular segments, including the decreasing size of dilations in superficial and deeper diving

arterioles (Gao et al., 2015), and pericyte coverage of vessels continuing to decrease into the capillary bed (Grant et al., 2019). However, most focus on localizing functional changes across the vasculature has fallen on contractile function or mural cell anatomy, so it is unclear whether other functions transition at the same, or different, locations. If different functions show the same transition points as contractile function, we could reasonably consider the vascular segments separated by these transition points as different vessel types. If, however, different vascular functions transition at different locations, it becomes somewhat meaningless to use these shifts in contractile function to define vascular segments.

To test the principle of whether there are clear locations within the vascular network at which transitions occur between multiple functions of the vasculature, we used immunohistochemistry to label different functional markers in brain slices expressing DsRed under the control of the NG2 promoter, as well as *in vivo* two-photon imaging of vascular responses to visual stimuli in awake mice to study the contractile characteristics of the vascular bed at the different functional transition points. There was no single point in the vasculature where the function transitioned from “arteriole-like” to “capillary-like.” Classic pericyte markers, NG2 and PDGFR β were in fact expressed throughout the vascular network, while transitions in markers for contractility, arteriolar compliance, neurovascular coupling and angiogenic potential occurred at different points in the vascular bed. We observed no sudden transitions in mural cell morphology across the vascular network, or at termination points of functional markers, though branch points showed distinct changes in mural cell morphology throughout the vasculature studied (arteriole-capillary). Finally, we also observed transitions of neurovascular and vasomotion function down the depth of arterioles, highlighting that arterioles become more capillary-like further from the pial surface.

MATERIALS AND METHODS

Animals

All experimental procedures were performed in accordance with the 1986 Animal (Scientific Procedures) Act and approved by the United Kingdom Home Office and the University of Sussex or University College London animal welfare ethical review boards. Experiments were performed on mice aged 1–8 months of both sexes (3–8 months for *in vivo* experiments). Mice were all on a C57BL/6J background which were either wild-type, expressed DsRed under the control of the NG2 promoter (Zhu et al., 2008), or for data collected from the 186 vessels recorded for the *in vivo* experiments expressed GCaMP6f under the control of the Thy1 promoter (Dana et al., 2014) or were SST-Cre crossed with floxed GCaMP6f. Food and water were available *ad libitum*, and mice were housed at 22°C in a 12 h light/dark cycle (which was reversed for the mice used in *in vivo* experiments).

Ex vivo Imaging

Each immunohistochemistry experiment used tissue from a minimum of 3 animals, on at least 3 different experimental days.

Methods for slicing, fixing and labelling were as described previously (Mishra et al., 2014; Boyd et al., 2021), as follows:

Slicing

NG2 DsRed mice were sacrificed by Schedule 1 approved methods (cervical dislocation followed by decapitation). The brain was then removed from the cranium and anterior and posterior coronal sections were removed, producing a block containing the central portion of the cerebral hemispheres. This block was then mounted onto a chilled slicing block using cyanoacrylate glue, with the inferior surface of the brain facing an agarose block. Using a vibratome, 200 μ m coronal brain slices were prepared in ice-cold slicing solution containing (mM): NaCl (124), NaHCO₃ (26), glucose (10), KCl (2.5), MgCl₂ (2), CaCl₂ (2), NaH₂PO₄ (1), kynurenic acid (1), bubbled with 95% oxygen and 5% CO₂. Slices were incubated in a slice storage container containing slicing solution bubbled with 95% oxygen and 5% CO₂ at room temperature to recover for > 30 min before fixation. Some slices were incubated in oxygenated slicing solution containing the fluorescent dye AlexaFluor 633 hydrazide (20 μ M) to label elastin (Shen et al., 2012).

Fixation, Blocking and Permeabilization

Slices were fixed in 4% paraformaldehyde solution for 30 min in a 24-well plate on a rotary shaker. Slices were then washed (3 \times 10 min) in 0.1 M phosphate buffered saline (PBS) on a rotary shaker at room temperature. In preparation for antibody labeling the slices were incubated for 3 h in 0.2% Triton X-100, 10% normal goat serum and 0.2M glycine in 0.1M PBS (blocking and permeabilizing solution) at room temperature to block generalized secondary antibody staining and to permeabilize the tissue.

Immunohistochemical Labeling

The following primary antibodies were used, all at 1:200 dilution in 0.1M PBS: rabbit anti- α SMA (Abcam), chicken anti-nestin (Abcam), rabbit anti-PDGFR β (Santa Cruz Biotechnologies), rabbit anti-GLUT-1 (Abcam), and chicken anti-GFAP (Abcam). Slices were incubated in primary antibody in PBS overnight (12–16 h) on a rotary shaker at room temperature, washed (3 \times 10 min) in 0.1 M PBS, followed by a 6–8 h fluorescently labeled appropriate secondary antibody incubation in 0.1M PBS at room temperature (AlexaFluor 647 goat anti-rabbit IgG or AlexaFluor 488 donkey anti-rabbit IgG - both Life Technologies; Goat anti-chicken CF488 IgY - Sigma). A no-primary control experiment was used for each antibody to check for non-specific binding of secondary antibodies. After labeling, slices were mounted onto microscope slides with Vectashield hard-set mounting medium including DAPI (Vector Laboratories) and covered with a glass cover slip. Cover slip borders were then sealed with nail varnish.

Confocal Imaging

SMCs and pericytes on large and small blood vessels, respectively, were readily identifiable through DsRed expression under the control of the NG2 promoter. Fluorescent labeling of the proteins described previously allowed visualization of the expression of proteins involved in the various functions of the cerebral

vasculature. Fixed slices were imaged using a Leica SP8 or Zeiss LSM780 confocal laser-scanning microscope. Imaging with multiple wavelengths was performed as sequential scans at each wavelength, to minimize “bleed-through” of fluorescence between channels. Z stacks of penetrating arterioles and their downstream capillary bed were obtained using a 20x air objective. These stacks were then projected as ‘maximum-intensity projections’ and stitched together using the MosaicJ plugin of ImageJ software (NIH, Bethesda, MD, United States). High-power Z-stacks of regions of interest, including the transition point of the various markers, were acquired using a 63x oil immersion objective.

Image Analysis

Image analysis was conducted using FIJI/ImageJ. Intersoma distances and diameters were calculated manually using the measurement tool in ImageJ. Measurements were made in one z-plane by drawing a line from the mid-point of the pericyte soma nearest the transition in labeling, to the mid-point of the closest proximal vascular mural cell on the same vessel. Somata were identified by DsRed expression and confirmed by DAPI stained cell nuclei. Diameters were measured as the distance between the innermost DsRed labeling on either side of the vessel lumen (i.e., the diameters include endothelial cell thickness). Branch points of the vessels were identified from the penetrating arteriole, termed branch order 0, incrementing by 1 at each subsequent branch point.

In vivo Imaging

Surgery and Two-Photon Imaging

Cranial window surgery to insert an optical window over V1 suitable for chronic, awake two-photon imaging was performed on 11 mice (6 female) under isoflurane anesthesia (see Shaw et al., 2021 for full methods). Following a one-week minimum post-surgery recovery period, mice were habituated to head fixation over multiple sessions. The experimental set-up consisted of a cylinder fitted with a Kuebler rotary encoder placed under a two-photon microscope with red and green filters (Scientifica), and in front of two computer screens for visual stimulus presentation. The objective used for two-photon imaging was a water-based 16x aperture (LWD, Nikon) with tissue excited at 940 nm (Chameleon Vision II Ti:Sapphire laser, Coherent). The visual stimulus was a drifting grating (PsychoPy) that varied either by contrast (25, 63, or 100%, all at 315° orientation), size (20° small circular stimulus or 220° full screen stimulus, both at 100% contrast), or spatial frequency (0.04 or 0.2 cycles per degree). As our data selected only the trials where the mouse was not running, chi-square tests were conducted to ensure all vessel segments were subject to the same distribution of stimulus conditions across trials (contrast $p = 0.94$; size $p = 0.99$; spatial frequency $p = 0.99$). Imaging sessions recorded fluorescent blood vessels and calcium activity (in excitatory or somatostatin cells), however, for the purposes of this study only data recorded from vessels was analyzed. To visualize blood vessels (penetrating arterioles and their downstream capillaries) during darkness or visual stimulation whilst the animal was running or resting, mice were injected with 2.5% (w/v) Texas Red Dextran dissolved in

saline (70 kDa *via* tail vein or 3 kDa subcutaneously, Fisher Scientific). For the stimulus-dependent vessel dilation data, we looked only at the dilations occurring during rest trials (i.e., to remove locomotion confounds). Imaging sessions were recorded in SciScan software (SciScan v1.2.1, Scientifica), where the imaged vessels ranged between 0 and 729.6 μm in depth (mean: 182.55 μm , SD: 132.65 μm), had an average pixel size of 0.1958 μm (range: 0.1484 - 0.4431 μm , SD: 0.08 μm) and were acquired at speeds of 7.6 Hz.

Vessel Classification

Vessel diameter, vessel depth and branch order were measured for all vessel classifications (see **Table 1**). Because we wanted to look at responses down the penetrating arteriole, in these experiments we separately classified the penetrating arteriole and capillary branching orders. This was therefore different to that used for immunohistochemical analysis, where the penetrating arteriole was always 0, and branch orders always referred to the position in the capillary bed. Here, for penetrating arterioles, branch order started at 0 at the pial surface, and increased by 1 down the length of the diving vessel after a branch offshoot was encountered. For capillaries, the first offshoot protruding off the penetrating arteriole was always given a branch order of 1 and vessel branch order increased by 1 for each bifurcation encountered (see **Supplementary Figure 1**).

Data Extraction

To improve image quality, vessel recordings were subject to several preprocessing steps prior to analysis. Using ImageJ: image type was set at 8 bit, images were despeckled, and light artifacts from the stimulus minimized using the ‘stack contrast adjustment’ plugin. To correct for motion artifacts (mainly arising from locomotion) vessel recordings were also image registered using Suite2P (Pachitariu et al., 2016). A custom MATLAB script was used to extract vessel diameter along the full width at half maximum (see Shaw et al., 2021). Diameter measurements were averaged along the entire vessel length, leaving an averaged (per frame) continuous diameter trace over time.

Data Analysis

Stimulus-dependent responses: Continuous vessel diameter traces were cut into 30 s trials around the stimulus presentations (stimulation occurred between 5-10 s). Data trials were then normalized to the 5 s pre-stimulus baseline period ($\Delta D/D = (D - D_{\text{min}})/(D_{\text{max}} - D_{\text{min}})$), and multiplied by 100 (to present data as a% increase from baseline). Only ‘rest’ trials were included, meaning there was no significant locomotion occurring in the 2 s prior to or during the stimulus. Locomotion events had a duration > 1/3 s and were distanced from other locomotion epochs by at least 1 s. The locomotion-free stimulus-centered vessel trials were categorized as responsive or non-responsive to stimulation, with responsive trials being those in which the maximum dilation during the stimulus event exceeded $0.5 \times$ the standard deviation of the 5 s baseline period. The maximum dilation was calculated by finding the peak of the trace during the stimulus. The peak at the cessation of the stimulus was taken

TABLE 1 | Vessel characteristics/responses separated by vascular segment.

Vessel Label	Branch Order	Sample size	Num Trials	Diameter (μm)	Depth (μm)	Pixel size (μm)	Response rate (%)	Peak dilation (%)	Peak dilation (μm)	Power at 0.1Hz
Penetrating Arteriole	0	21 vessels (20 for Welch's)	5-19 (mean: 12.86)	10.14 \pm 4.66	74.40 \pm 48.31	0.2438 \pm 0.096	35.51 \pm 25.36	All: 5.59 \pm 5.20 R: 12.50 \pm 17.80	All: 0.47 \pm 0.05 R: 0.66 \pm 0.11	0.040 \pm 0.05
Penetrating Arteriole	1	22 vessels	2-17 (mean: 11.23)	9.62 \pm 4.91	133.39 \pm 78.07	0.2351 \pm 0.078	26.99 \pm 26.95	All: 4.69 \pm 3.29 R: 7.90 \pm 4.56	All: 0.35 \pm 0.03 R: 0.34 \pm 0.06	0.025 \pm 0.03
Penetrating Arteriole	2	12 vessels	3-17 (mean: 12.33)	8.60 \pm 2.35	130.03 \pm 90.50	0.2025 \pm 0.053	19.66 \pm 14.22	All: 3.14 \pm 1.48 R: 5.25 \pm 2.86	All: 0.43 \pm 0.04 R: 0.40 \pm 0.07	0.017 \pm 0.03
Penetrating Arteriole	3 +	9 vessels (8 for Welch's)	11-17 (mean: 13.78)	11.31 \pm 4.65	234.14 \pm 62.05	0.2528 \pm 0.089	13.87 \pm 11.35	All: 3.03 \pm 1.90 R: 3.32 \pm 3.42	All: 0.75 \pm 0.1 R: 0.86 \pm 0.15	0.017 \pm 0.02
Capillary	1	50 vessels (49 for Welch's)	2-17 (mean: 11.24)	6.72 \pm 3.36	161.51 \pm 139.57	0.1603 \pm 0.026	18.89 \pm 18.28	All: 5.47 \pm 6.69 R: 9.29 \pm 8.99	All: 0.37 \pm 0.03 R: 0.42 \pm 0.07	0.025 \pm 0.06
Capillary	2	29 vessels	3-19 (mean: 12.41)	5.84 \pm 2.02	161.86 \pm 139.19	0.1595 \pm 0.022	17.10 \pm 16.23	All: 5.22 \pm 3.84 R: 7.90 \pm 8.79	All: 0.34 \pm 0.03 R: 0.36 \pm 0.07	0.027 \pm 0.05
Capillary	3	22 vessels	4-19 (mean: 12.14)	5.75 \pm 1.81	182.96 \pm 100.9	0.1541 \pm 0.017	25.14 \pm 27.40	All: 4.28 \pm 2.67 R: 7.00 \pm 4.98	All: 0.28 \pm 0.04 R: 0.24 \pm 0.06	0.011 \pm 0.02
Capillary	4 +	21 vessels	2-18 (mean: 11.43)	4.84 \pm 1.62	219.14 \pm 92.02	0.1564 \pm 0.020	18.53 \pm 22.92	All: 9.04 \pm 11.79 R: 7.14 \pm 4.56	All: 0.22 \pm 0.03 R: 0.14 \pm 0.03	0.009 \pm 0.01

Values are calculated per individual vessel (presented as mean \pm STD between vessels). All: averaged across all trials; R: averaged across responsive trials only. Note that some sample sizes are lower for the Welch's power spectrum comparisons (specified in brackets) as 3 vessels were removed as outliers (see methods for outlier details).

by measuring the value of the vessel dilation trace at 10 s. The onset time of dilation responses was calculated as the time to 10% of the maximum peak (during the stimulation period). Four noisy data trials (from 2222 total trials) were removed from the diameter traces [3 trials from vessel 77 (C5) and 1 trial from vessel 167 (C5)] because dilation shifts were very noisy (high frequency shifts reflective of motion artifacts/signal loss) and peaks exceeded 10x the average (mean dilation peak across all responsive vessels during visual stimulation: 8.43 vs 101.3%, 123.2, 214.3, and 92.4%).

Power Spectra Responses

Welch's power spectral density estimates were computed across all continuous vessel diameter traces. These recordings could be during rest, visual stimulus and/or locomotion, and we included the entire continuous trace in the power analysis as (a) the stimulus presentation occurred for 5 s every 30 s, meaning the power spectrum peak corresponding to stimulus presentation would be at 0.03 Hz (outside our 0.05-0.15 Hz range of interest); and (b) there was no significant difference in the amount of time spent in locomotion between the different vessel categories (mean time spent in locomotion: 13.07%, standard deviation: 6.39%, $p = 0.45$). All continuous data was detrended by subtracting the baseline (the 8th percentile calculated over a 15 s time window; see Bonnar et al., 2021). The MATLAB function "pwelch" was used to conduct discrete Fourier transforms across 60 s time windows. Data plots display raw power spectra and power spectra corrected for 1/f (pink) noise (1/f corrected trace = Welch power spectrum trace * corresponding Welch frequency trace). All data underwent outlier removal based on the maximum value detected between 0-1 Hz (all traces containing peaks greater than 3 * the standard deviation of all the maximum values were removed). This resulted in the removal of 3/186 vessel diameter traces (from categories: PA BO0, PA BO3, and Cap BO1). For comparing power spectra, the power value at 0.1 Hz was extracted. The number of vessels which showed high power at 0.1 Hz was assessed by separating vessel diameter traces by those above and below a set threshold (standard deviation across all detected power values at 0.1 Hz * 0.5).

Statistics

Statistical analyses were conducted in RStudio. Data in graphs are presented as mean \pm SEM, and individual dots on bar graphs represent data from individual blood vessels, whereas violin plots were used to represent individual stimulus-presentation trials. For comparing the distribution of vessel responsivity rates or stimulus condition, data was taken from individual stimulus-presentation trials, and tested using a 3D Cochran-Mantel-Haenszel test with Fisher's *post hoc* comparison. For comparing the sizes of vessel dilations or the power in the vasomotion frequency range, where data was non-normal, data was averaged across vessels and a Kruskal-Wallis test with *post hoc* pairwise Wilcoxon rank sum tests were utilized. Branch order and/or vessel categorization (penetrating arteriole/capillary) were set as the independent variable, dilation peak (for all trials during stimulus period or at stimulus cessation), time to 10% of dilation peak, power (value at 0.1 Hz or AUC between 0.05-0.15 Hz)

or locomotion frequency (% time spent in locomotion) as the dependent variable. For comparing the individual vessel diameters and depths, where data was normally distributed, one-way ANOVAs with Tukey's *post hoc* comparisons were used. For *post hoc* tests, for clarity only significant ($p < 0.05$) or trend ($p < 0.1$) level comparisons are shown, but these tests were always carried out, so absence of a displayed p value indicates a non-significant ($p > 0.1$) result.

RESULTS

“Pericyte Markers” Are Expressed Throughout the Microvascular Network

NG2, a chondroitin sulfate proteoglycan, is considered a pericyte marker and has been reported at low levels in smooth muscle cells (Kumar et al., 2017). However, NG2-controlled DsRed labeling is found throughout the microvascular tree, including in smooth muscle cells with a clear banded morphology (Figure 1). Immunohistochemical labeling for another pericyte marker, PDGFR β (Winkler et al., 2010), also reveals protein expression throughout the microvascular tree, including in SMCs on pial vessels as well as penetrating arterioles and capillaries (Figure 1).

Markers of Contractility and Elasticity Terminate at Different Positions of the Vascular Tree

As discussed above, a much studied transition across the vascular bed has been that between vascular mural cells expressing high levels of α SMA and those expressing low or no levels of α SMA, which broadly occurs between ensheathing and mesh pericytes, though the distribution of α SMA termination positions is different from that of the change in vessel coverage (Grant et al., 2019). As discussed above, this switch in α SMA expression may coincide with the faster responses observed in low branching order capillaries (Rungta et al., 2018), though contractility of pericytes may extend beyond this transition in α SMA expression, as mid-capillary pericytes also dilate (Hall et al., 2014; Kisler et al., 2017; Hartmann et al., 2021b), presumably using different contractile machinery (Hartmann et al., 2021b). This latter study also found these mid-capillary pericytes to show slower constriction to optogenetic stimulation than did α SMA expressing-pericytes.

Vessels also show a transition in expression of elastin across the vascular bed, representing another flow regulation functional transition. In mammals, the number of layers of elastin expressed in the vessel wall scales with blood pressure and arterial size, being highest in the largest arteries (Cocciolone et al., 2018), but with only one layer, just abluminal to endothelial cells, for most intracerebral arterioles (Shinaoka et al., 2013). Elastin expression terminates before arterioles branch successively to become capillaries, presumably reflecting the lower blood pressures to which these smaller vessels are exposed. Elastin's main role is to increase vessel distensibility, storing energy in the vessel wall during systole, and releasing it during diastole, thus smoothing blood flow across the heart beat and dampening

pressure waves, protecting downstream vascular beds from large fluctuations in pressure (Shinaoka et al., 2013). In addition to this biomechanical role, it may also act as a signaling molecule, regulating and attracting smooth muscle and immune cells (Cocciolone et al., 2018).

Elastin can be readily labeled by i.v. injection or incubation of tissue with AlexaFluor 633 hydrazide (Shen et al., 2012), but its role in microvascular regulation remains understudied. Recently, however, it has been used to identify intracortical arterioles in which endothelial caveolae were found to be critical players in neurovascular coupling (Chow et al., 2020), and was found around precapillary sphincters branching off penetrating arterioles that regulate blood flow and pressure into the cortical capillary network (Grubb et al., 2020; Zambach et al., 2021).

To test whether the transitions between α SMA and elastin expression occur at the same location, thus defining two different vessel “types,” we used immunolabeling for α SMA combined with AlexaFluor 633 hydrazide labeling of elastin. We found that elastin terminated on vessels of large and small diameters (Figures 2B,D) but always upstream from the point of termination of α SMA (Figures 2A-D,F; 4/4 double-labeled branches showed non-overlapping termination points).

Nestin Expression in the Capillary Bed Extends Beyond α SMA Labeling

Nestin is an intermediate filament protein, commonly thought to be expressed in proliferating cells (Suzuki et al., 2010), and is associated with cardiovascular remodeling (Calderone, 2018). Recently it has been shown to be expressed in quiescent as well as proliferating endothelial cells and, indeed, may exert an inhibitory effect on endothelial cell proliferation (Dusart et al., 2018). Nestin expression by endothelial cells may therefore not necessarily reflect ongoing angiogenesis, but rather indicate the angiogenic potential of vessels. Double labeling of nestin and α SMA or elastin in NG2-DsRed mice revealed the capillary network to be nestin-positive (Figure 3). Nestin labeling extended from the capillary bed into lower branching order capillaries that were also α SMA positive (Figures 3B-D), but never overlapped with elastin-labeled vessels (Figures 3F-H).

To quantify the expression patterns of nestin, α SMA and elastin across the microvascular network, we next categorized the termination points of each label according to vessel lumen diameter, branching order from the penetrating arteriole (where the penetrating arteriole is 0th order; Figure 4A), and the distance between mural cells, or inter-soma distance (ISD). This latter measurement serves as an indicator of the morphology of these cells, as they transition from banded SMCs that are adjacent to each other, to pericytes with distinct soma and processes that become progressively longer along the vascular bed into the capillary network (Grant et al., 2019). Termination points of the different markers occurred at vessels of similar diameters (Figure 4B), but at different branch orders (Figure 4A) or ISD values (Figure 4C), indicative of different positions in the vascular tree. Specifically, elastin termination points were on vessels of significantly lower branch orders than termination points for α SMA or nestin, and mural cells at elastin and

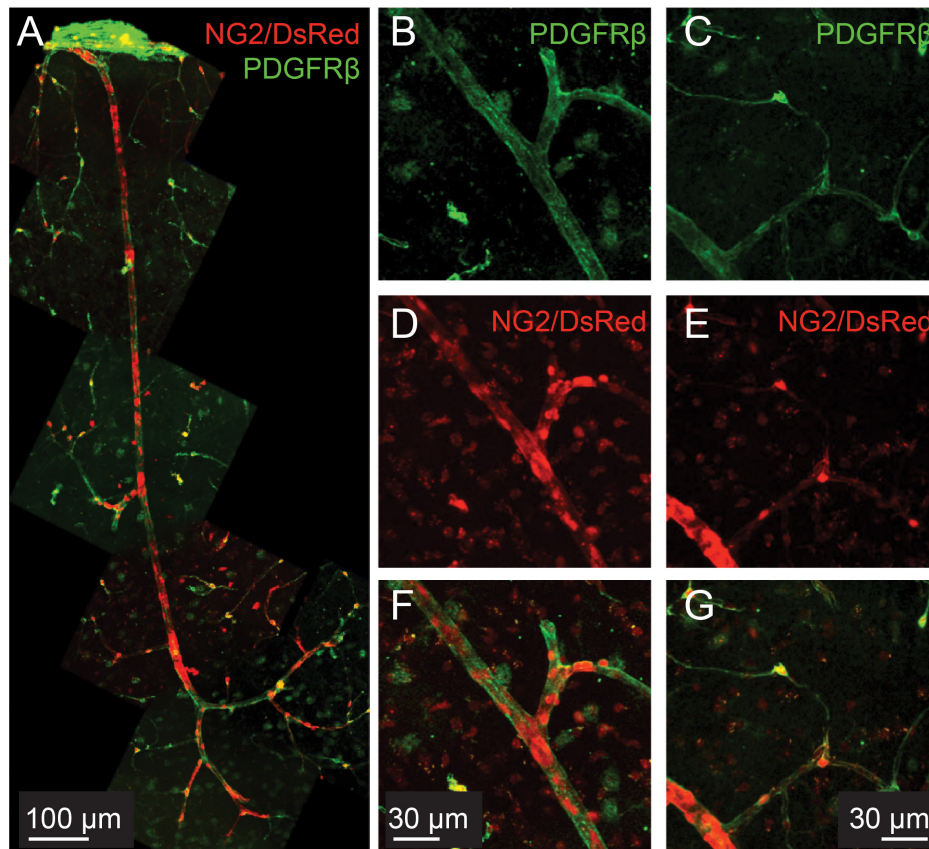


FIGURE 1 | PDGFR β is expressed in smooth muscle cells as well as pericytes in the cortical microvasculature. Green: immunohistochemical labeling for PDGFR β . Red: DsRed expressed under the control of the NG2 control promoter. **(A)** Example vascular bed from pial surface (top) to layers V/VI of cortex (bottom). **(B–G):** Two different example vessels at higher magnification, to show capillary pericyte labeling of both markers.

nestin termination points were significantly closer together than at termination points for α SMA. Furthermore, consistent with our observations from double labeling of elastin and α SMA (**Figure 2**), the ISD at which elastin terminated was always smaller than that where α SMA terminated (**Figure 4D**). Thus, each functional transition occurred at a different point in the vascular tree, as defined by branch order and/or ISD. This means that in addition to the functional transitions between elastin presence and absence, and α SMA presence and absence, there is an additional functional transition point, where nestin terminates, between these two positions.

nNOS Association With Blood Vessels Is Strongest in Low Branching Order Vessels

The vasodilatory signaling molecule nitric oxide (NO) is released by sub-populations of interneurons during synaptic activation, and can mediate or modulate neurovascular coupling (Attwell et al., 2010). In neocortex, if not cerebellum, it seems predominantly involved in regulating arteriole but not capillary diameter, a transition appearing to occur between the diving arteriole and first capillary branch (Hall et al., 2014;

Mishra et al., 2016). To investigate whether this reflects neuronal NO sources for vasodilation differing along the vascular tree, we immunohistochemically labeled brain slices for neuronal nitric oxide synthase (nNOS or NOS1) while labeling the vasculature with Alexa647-conjugated isolectin B4, which binds to the basement membrane (Peters and Goldstein, 1979). Consistent with previous studies (Vlasenko et al., 2007), some arterioles showed clear nNOS labeling around (but not within) vessels which extended into the capillary bed, as well as parenchymal signal (**Figures 5A–F**). To assess whether nNOS is preferentially expressed around particular elements of the vascular network, we measured the intensity of labeling immediately around vessels of different branching orders, and at increasing distances from these vessels (**Figure 5G**). Linear mixed modeling (with distance from vessel and branch order as fixed factors and vessel as a random factor) showed nNOS labeling to be significantly more intense at the vessel compared to the parenchyma ($F = 7.20$, d.f. = 3,30, $p = 0.0009$), but there was no difference in nNOS labeling around different branches ($F = 0.87$, d.f. = 3, 30, $p = 0.47$), nor did branch order affect the drop-off in signal away from the vessel ($F = 0.25$, d.f. = 9, 30, $p = 0.98$). When just the labeling at the vessel was considered, branch order was borderline-significant ($F = 4.1$, d.f. = 3,6, $p = 0.068$). Together these results suggest

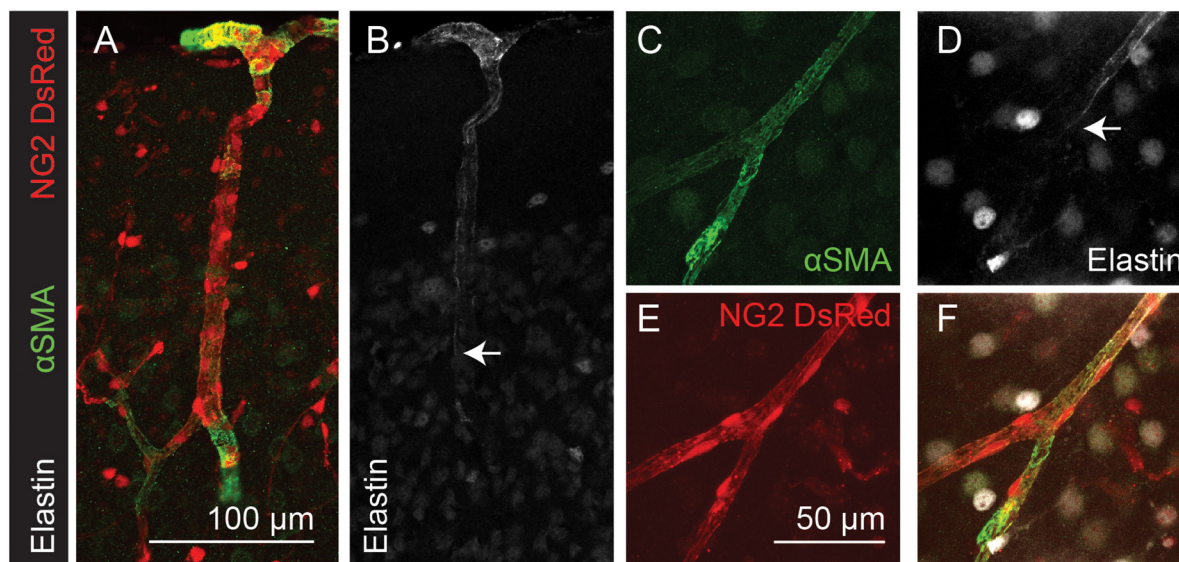


FIGURE 2 | Elastin labeling (white) terminates before α SMA labeling (green) in the cortical microvasculature (arrows). Red: DsRed expressed in NG2-positive mural cells. Panels (A–F) show arteriole and capillary example images. Arrows indicate termination point of elastin labeling. Images are representative of 9 (α SMA) and 12 (elastin) vascular trees, which are quantified in **Figure 4**.

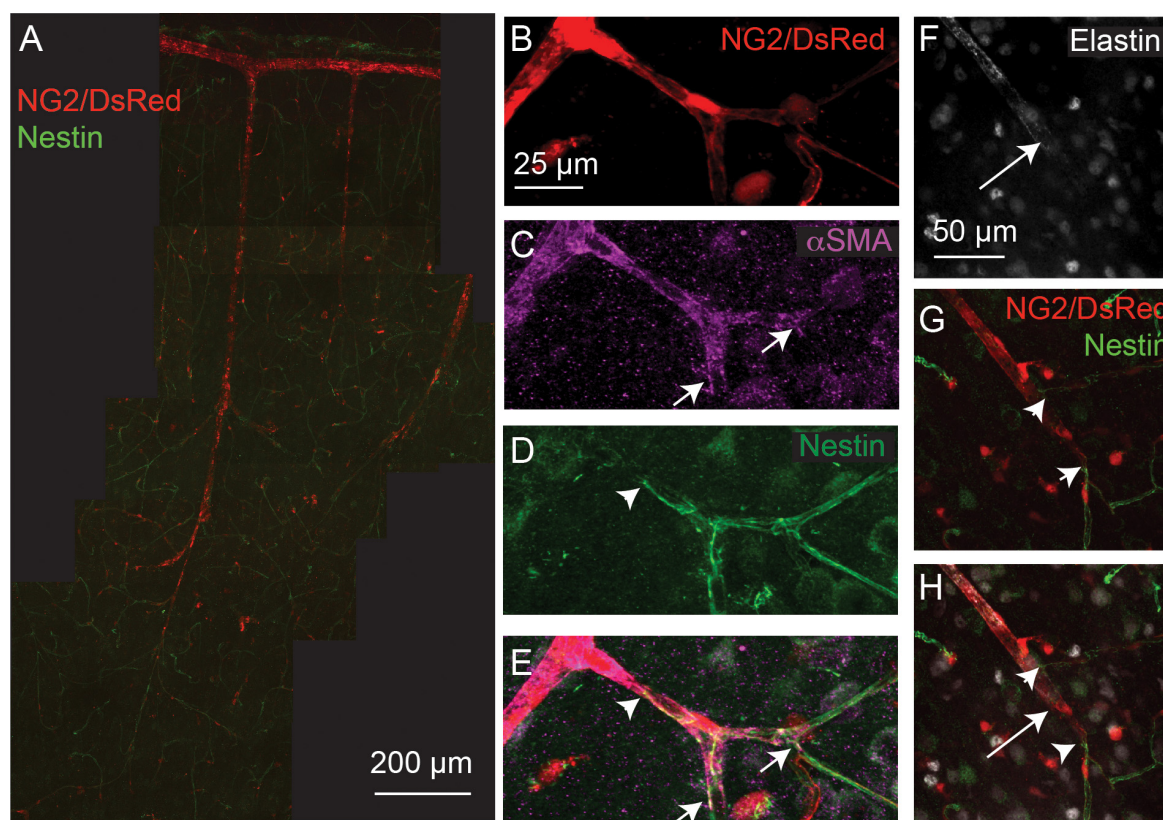
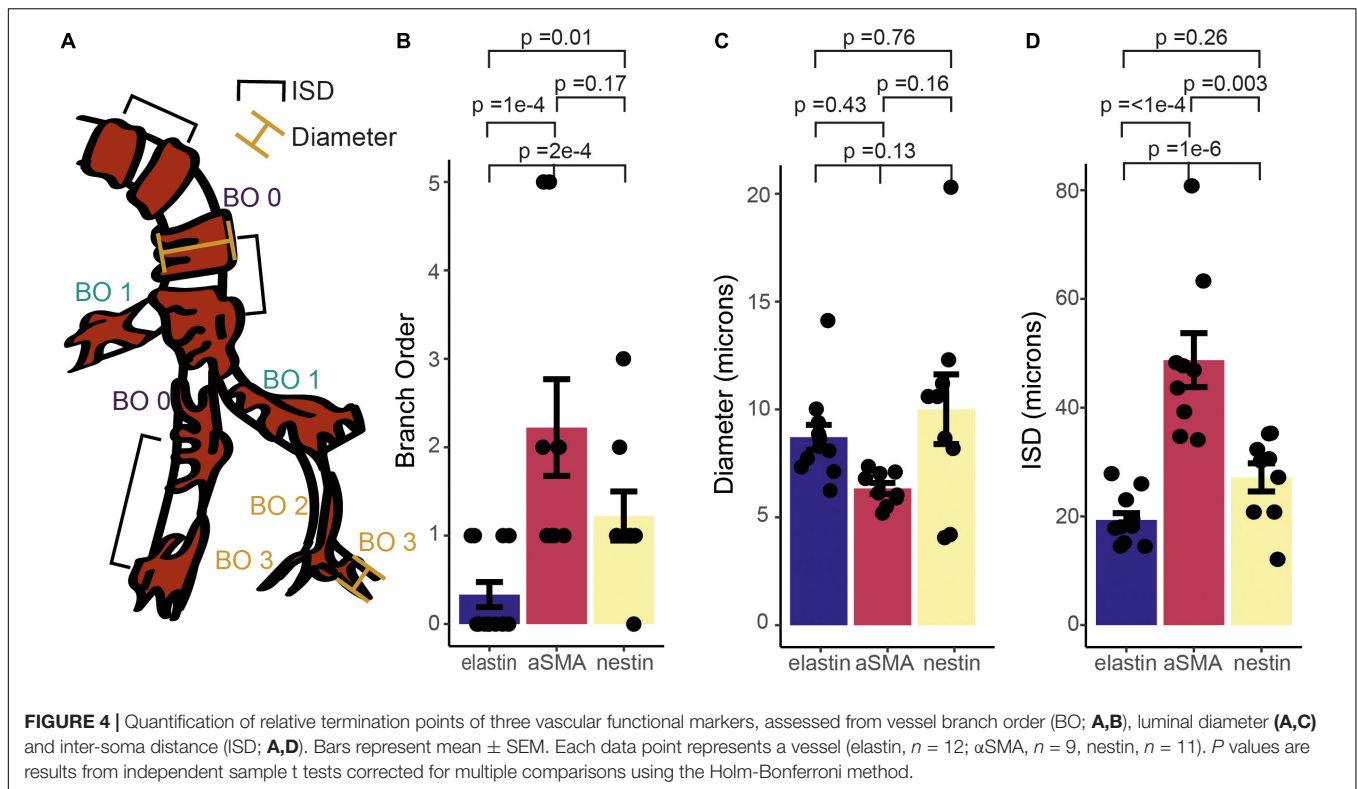


FIGURE 3 | Nestin expression (green) extends from the capillary bed to terminate (arrowheads) on vessels that express α SMA (magenta; termination point showed with small arrows), but does not extend as far as the termination point of elastin (long arrows). Red: DsRed-NG2 positive mural cells. (A) Example vascular bed from pia (top) to layer V/VI of cortex. (B–E) and (F–H) are two other example vessels at higher magnification. Images are representative of 11 vascular trees, which are quantified in **Figure 4**.



that nNOS interneurons do target blood vessels, but that the degree of association is similar between penetrating arterioles and the first few capillary branches from the arteriole. The tendency for marginally stronger labeling at arterioles may contribute to the different dependence of neurovascular coupling reported in capillaries and arterioles (Mishra et al., 2016), but is unlikely to explain it entirely. Interestingly, recent work has shown that NO can modulate the propagation of vasodilation through the vascular network (Kovacs-Oller et al., 2020). It is an open, and important, question if nNOS-derived NO could contribute to neurovascular coupling by modulating the integration of vascular signals across the network in addition to its direct effect on vasodilation, and how such NO release would interact with endothelial (eNOS)-derived NO, which is also an important modulator of neurovascular function (e.g., Toth et al., 2015; Chow et al., 2020). Studying the pattern of targets of NO signaling (e.g., soluble guanylyl cyclase, or cytochrome P450 ω hydroxylase (Sun et al., 2000) should be informative in identifying how different vessels can respond to this perivascular NO production.

Shifts in Functional Expression Are Not Accompanied by Sudden Shifts in Vascular Cell Length

Previous work suggests that mural cell morphology changes in some respects near to the point at which α SMA expression terminates, with α SMA-expressing ensheathing pericytes covering the underlying vessel to a greater degree than mesh pericytes immediately downstream of the α SMA termination point (though the distribution of changes in coverage is shifted

to higher branch order vessels than the termination of α SMA; Grant et al., 2019). However, the degree of vessel coverage by processes is not the only morphological change that pericytes undergo across the vascular network, as they change cell length (indicated by ISD), as well as soma orientation and shape, number and orientation of processes and many other descriptors (Zimmermann, 1923). ISD is different across pericyte categories as defined by vessel coverage, with ensheathing pericytes having a lower ISD than mesh pericytes (Grant et al., 2019; Shaw et al., 2021), so we wondered whether any abrupt shifts in ISD would be observed at functional marker transitions, suggestive of a major change in vessel type at this point. We therefore plotted the ISD immediately before and after the termination points of α SMA and elastin, normalized to the ISD at the termination point (Figures 6A,B). There was no significant change in ISD at the termination point of either functional marker. We had measured more ISD values on the elastin vessels, so also calculated the ISD two further cells away from the termination point, which also showed no differences compared to the cells nearer the transition point, further emphasizing the lack of an abrupt change (Figure 6C).

Mural Cells Are More Densely Spaced at Branch Points Compared With Non-branch Regions

Pericytes are often found at vessel bifurcations (Hartmann et al., 2015), but we also noticed that they appeared to have a different morphology at branch points, cells being clustered with a shorter ISD and greater vessel coverage (Figure 7A). Indeed, both

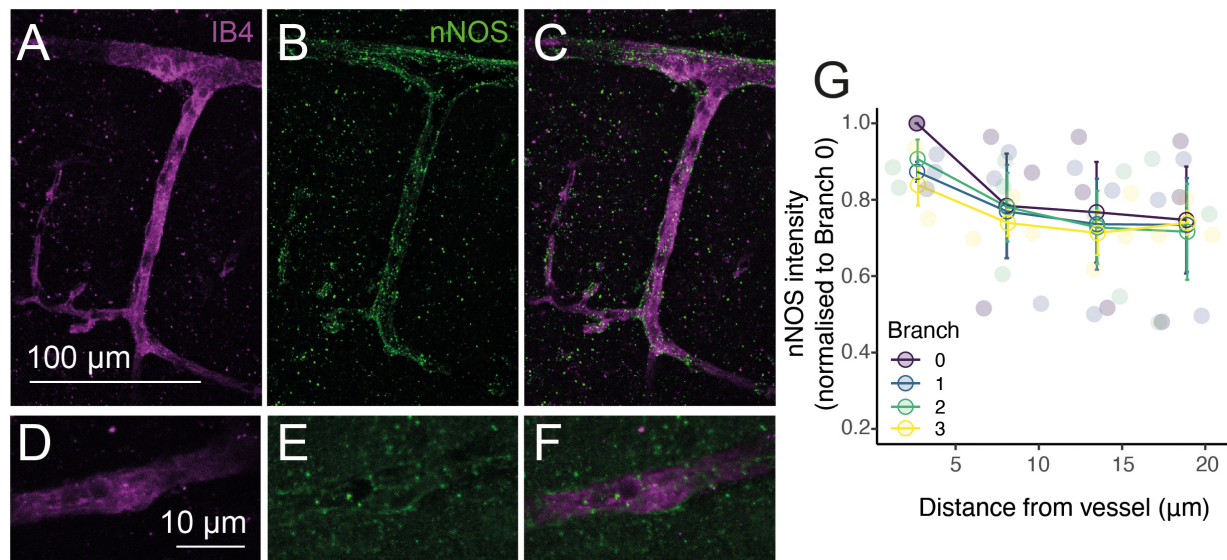


FIGURE 5 | NO production occurs preferentially near vessels. Alexa 647-conjugated Isolectin B4 (IB4; **A,C,D,F**; magenta) and immunohistochemical labeling of nNOS (**B,C,E,F**; green) show increased nNOS expression around vessels. (**G**): Quantification of nNOS labeling around different vessel branches, expressed by normalizing the intensity of labeling around the vessel to that at the penetrating arteriole (0th order vessel). Data represents mean \pm SEM. Individual points show different vessels ($n = 3$).

vessel diameter and ISD were greater at branch points than expected based on the average of the upstream and downstream values of these parameters (**Figure 7B**). Furthermore, while the increase in diameter was larger at branch points on larger vessels (**Figure 7C**), the change in morphology, indicated by the relative change in ISD, occurred on all vessels irrespective of the vessel diameter or ISD at the branch point (across a range of vessels from 5–20 μm in diameter; **Figures 7D,E**). Because ISD generally increases as vessel diameter decreases into the vascular bed (**Figure 7F**), mural cells at bifurcations therefore, at least in terms of ISD, have a morphology more like larger upstream vessels. This suggests the vascular network shows functional specialization at bifurcations, in addition to classic arteriole-capillary transitions of function. Pericytes at branch points can generate different calcium signals and constriction of different downstream branches, but downstream branch points were less responsive to applied vasoactive agents than upstream vessels (Gonzales et al., 2020). Our data suggest that these downstream bifurcations may still be specialized in some manner, compared to adjacent non-branch capillaries. Pericytes are known to occur frequently at bifurcations (Hartmann et al., 2015). Our analysis reveals that these pericytes at or near branch points are also shorter than expected from the ISD of non-branch pericytes immediately up or downstream.

In vivo Vascular Responses

Functional markers and anatomical changes in mural cell density indicate, therefore, that vascular function changes gradually across the vascular network, with multiple functional transition points corresponding to changes in vascular distensibility, proliferative capacity and contractility. These changes are

superimposed on gradually changing mural cell properties that do not show clear alterations (at least in some features) at these transition points, but which do show specializations at branch points that likely reflect functional changes. Furthermore, inspection of the images of DsRed-positive mural cells on arterioles (in **Figures 1A, 2A, 3A, 7A** and **Supplementary Figure 1**) show that, unlike commonly assumed, mural cells on arterioles often lose their annular shape as the arteriole dives into the cortex, forming intermediate pericyte morphologies with a distinct soma and processes in deeper regions.

We wanted to test, where possible, how these transitions reflected alterations in the physiological responses of different components of the vascular bed, so studied three properties of visual cortical microvasculature: frequency and size/timing of dilations in response to visual stimulation, and low frequency oscillations in the vasomotion range. We compared responses of the penetrating arteriole before and after smaller branches had come off the main vessel, as well as comparing responses of the arteriole to increasing branch orders of capillaries (see **Supplementary Figure 2D** for how different vessels' branch order was defined in order to separate vascular responses along the length of the penetrating arteriole). All data is summarized in **Table 1** (and data demographics [cortical depth, diameter] shown in **Supplementary Figure 3**).

Arterioles Dilate the Most Near the Cortical Surface but Deeper Sections Behave More Like Capillaries

Vessel responsiveness was assessed across the microvasculature by testing if visual stimulation led to an increase in vessel diameter (of > 0.5 standard deviations of the baseline). Sections

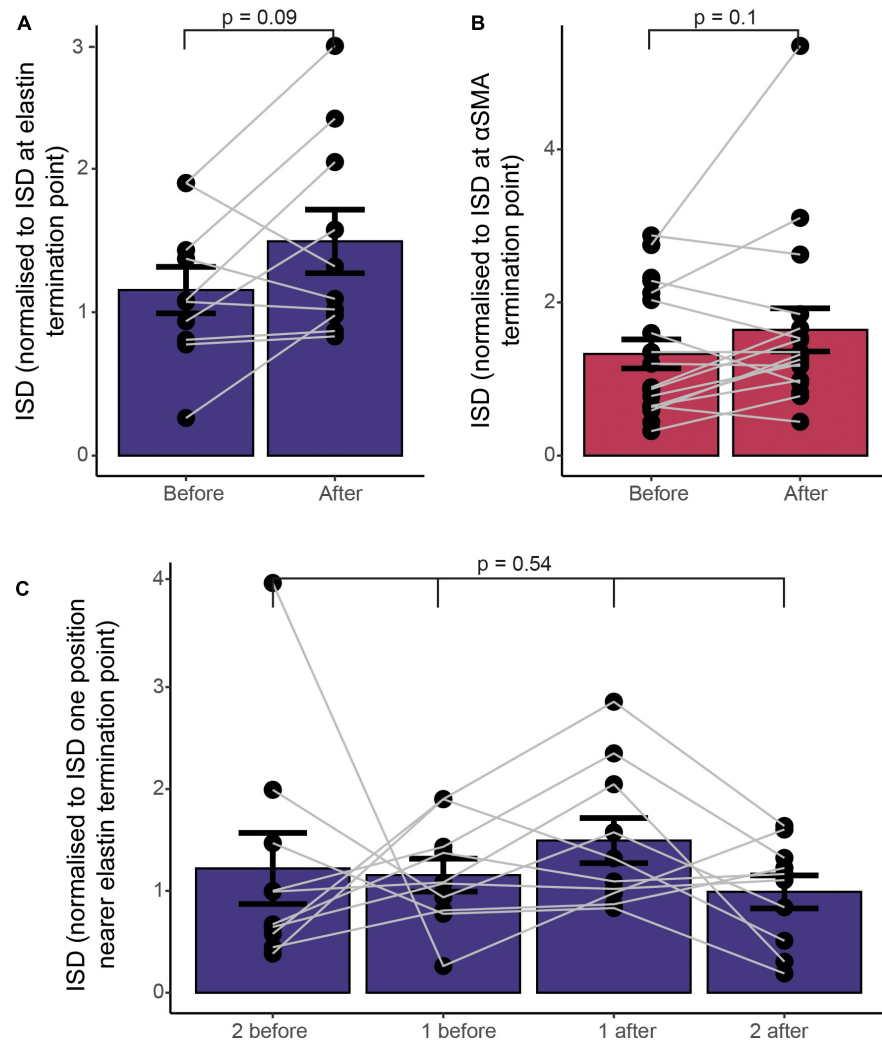


FIGURE 6 | Functional markers terminate at different points along the vascular tree, without a concomitant change in vascular mural cell morphology, as assessed by ISD. **(A)** ISD before or after the termination point normalized to the ISD at the termination point of elastin **(A)** or α SMA **(B)**. Bars show mean \pm SEM, with connected data points representing individual vessels ($N = 9$ for α SMA and 11 for elastin). P values are from paired t tests **(A,B)** or a repeated measures ANOVA **(C)**.

of arterioles near the cortical surface dilated significantly more frequently than either downstream arteriole sections (\geq BO2) or downstream capillaries (Figure 8 and Supplementary Figures 4–6). Specifically, descending the penetrating arteriole, dilations occurred with a similar frequency, and responses when they occurred were of a similar size in the first two sections of the vessel (before the first capillary branch and between the 1st and 2nd branches off the penetrating vessel), but these superficial responses were more frequent and tended to be larger than responses of the deepest sections of the penetrating vessel (Figure 8).

Across the capillary bed (i.e., any vessel downstream of the penetrating arteriole, populated by high α SMA ensheathing pericytes or low α SMA mid-capillary pericytes), dilation responses were of a similar frequency and size, with response frequencies being similar to the deep sections of the penetrating vessel (i.e., PA2 + ; Figure 8). There were no differences between

the speed of dilations across the different vascular segments (Supplementary Figures 4–6).

More Arterioles Near the Cortical Surface Show Higher Power in the Vasomotion Range Compared to in Deeper Sections and the Capillary Bed

We next measured vasomotion of the vascular diameter of surface and deep sections of penetrating arterioles and different branches of the capillary bed, finding that vasomotion (power at 0.1Hz) was similar between arteriole sections and between capillaries (Figures 9A–F). When we compared arterioles near the cortical surface (PA0–1), to deeper arteriole sections (PA2 +) and capillaries (C1 +), there was some evidence that penetrating arterioles show more vasomotion near the pial surface, as power at 0.1Hz was highest in superficial penetrating arterioles

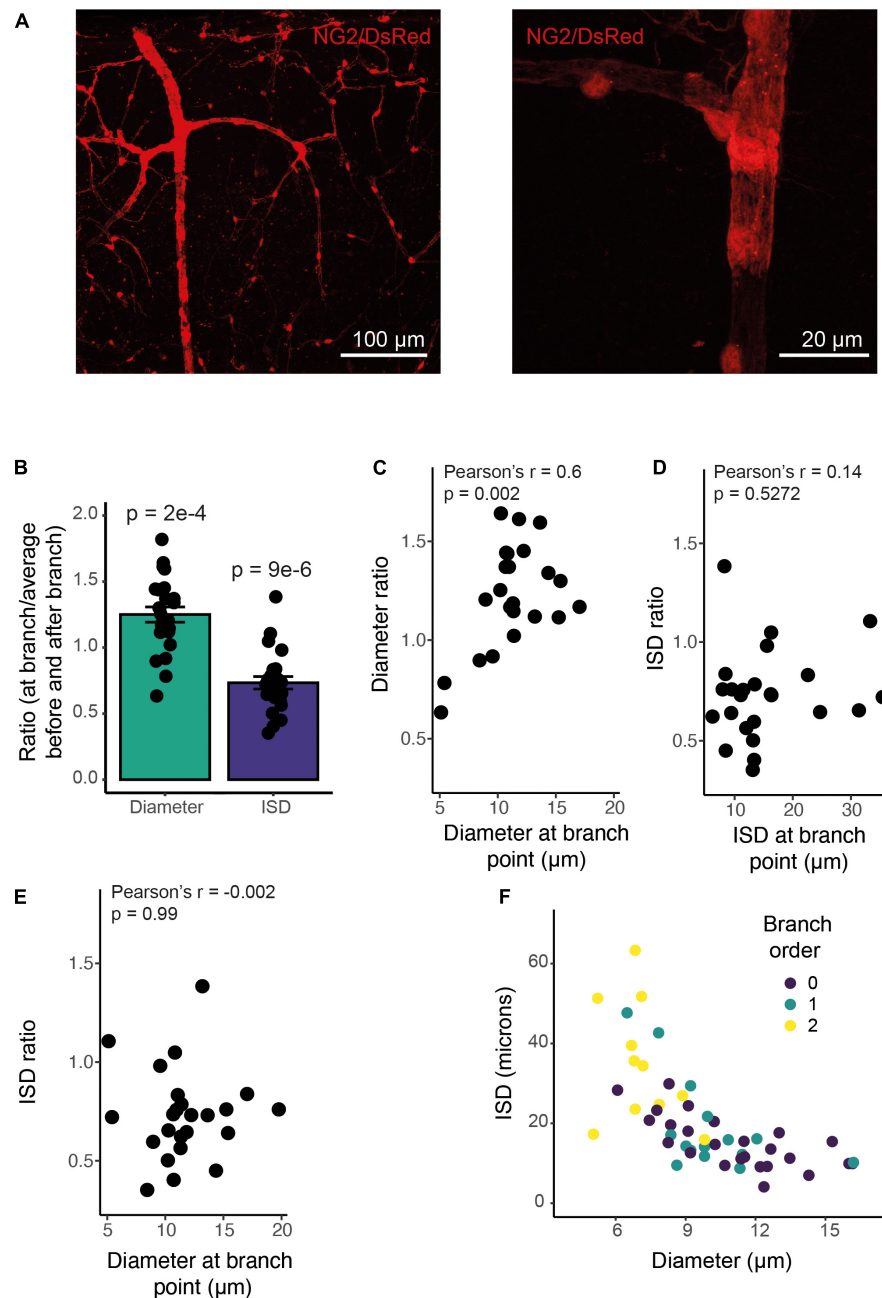


FIGURE 7 | ISD increases along the vessel but is smaller at branch points of any size. **(A)** Example NG2/DsRed labeled cortical vasculature, showing a whole penetrating arteriole and branches (left), and a high-resolution image of a single branch point, with clustered pericytes (right). This whole vascular tree is shown in **Supplementary Figure 1**. **(B)** Branch points have a larger diameter and more densely spaced pericytes than surrounding non-branch regions: The ratio of diameter at a branch point to the average of values immediately up and downstream of that branch point is larger than 1, and the ratio of ISD at a branch point to the average of ISD up and downstream of that branch point is less than one. $N = 24$ branch points. P values are from one sample t tests compared to 1. **(C)** This diameter ratio correlates significantly with the vessel diameter: smaller vessels show a relatively smaller increase in diameter at branch points. ISD ratio (calculated as in B) is uncorrelated with either ISD **(D)** or diameter **(E)**. Thus, smaller vessels show the same proportional increase in pericyte density at branch points as do large vessels. Bars represent mean \pm SEM, data points are individual branches from 16 vessels. **(F)** Mural cell ISD increases with decreasing vessel diameter. Dots represent individual ISD values from 20 vessels.

(Figure 9G), and significantly more upstream arteriole sections (PA0-1: 33%) showed an increase in power in the vasomotion range compared to capillaries (C1 + : 13%) (0.5 SD above baseline; Figure 9G,H).

DISCUSSION

Our results demonstrate the existence of an intermediate transition point in vascular function between the termination

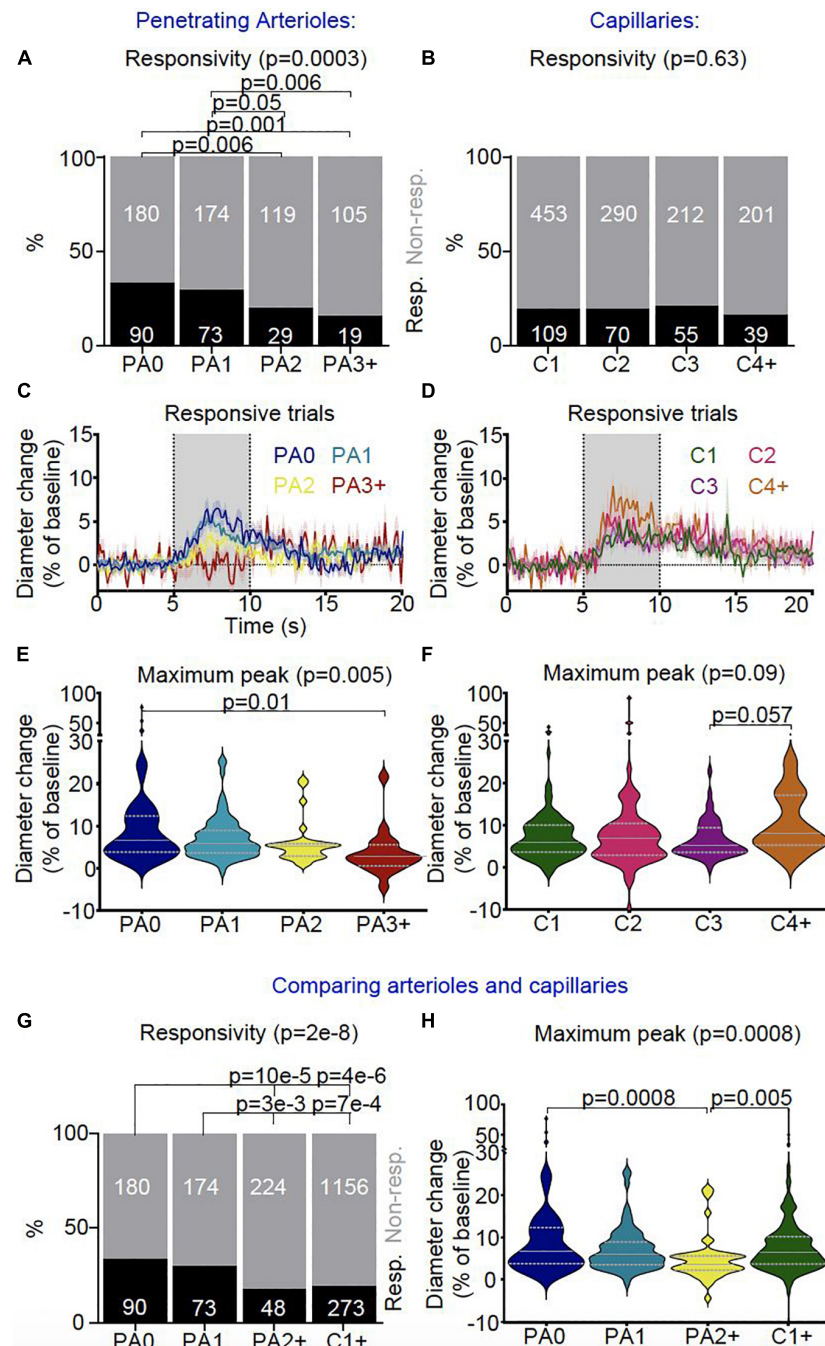


FIGURE 8 | *In vivo* vascular stimulus-dependent responses separated by vessel segment (responsive trials only). Stimulus-induced vascular dilations were classified as responsive (black) or non-responsive across the vascular segments for **(A)** penetrating arterioles (PA0 nTrials = 270, nVessels = 21; PA1 nTrials = 247, nVessels = 22; PA2 nTrials = 148, nVessels = 12; PA3 + nTrials = 124, nVessels = 9) and **(B)** capillaries (C1 nTrials = 562, nVessels = 50; C2 nTrials = 360, nVessels = 29; C3 nTrials = 267, nVessels = 22; C4 + nTrials = 240, nVessels = 21). Lower order (PA0 and PA1) penetrating arterioles were more likely to dilate during stimulus presentation than higher order diving arterioles (PA2, PA3 +), whereas no differences were found in the response rates between capillaries. Vessel responses were plotted for the **(C)** penetrating arteriole (PA0 nTrials = 90, nVessels = 18; PA1 nTrials = 73, nVessels = 15; PA2 nTrials = 29, nVessels = 9; PA3 + nTrials = 19, nVessels = 7) and **(D)** capillary (C1 nTrials = 109, nVessels = 36; C2 nTrials = 70, nVessels = 21; C3 nTrials = 55, nVessels = 17; C4 + nTrials = 39, nVessels = 14) classifications during the stimulus for responsive trials only (traces represent mean \pm SEM across trials; data from all trials are shown in **Supplementary Figure 4**). Gray bar (5–10 s) shows when the stimulus was presented. Responsive trials, averaged across individual vessels, and the maximum dilation during the stimulus presentation was compared between branch orders for **(E)** penetrating arterioles and **(F)** capillaries. There were differences in the maximum dilation during stimulus presentation across sections of penetrating arterioles ($p = 0.04$; Kruskal-Wallis test as data is highly skewed), with the largest dilations in the most superficial section (PA0) ($p = 0.06$ vs PA3 +, pairwise comparison with Wilcoxon Rank Sum test), whereas stimulus-induced dilations were not significantly different across the capillary bed. We then compared vessel responses between the superficial and deep sections of penetrating arterioles (PA0 nTrials = 270, nVessels = 21; PA1 nTrials = 247, nVessels = 22; PA2 + nTrials = 272, nVessels = 21) and the capillary network (C1 + nTrials = 1429, nVessels = 122). **(G)** Lower order (more superficial) penetrating

(Continued)

FIGURE 8 | arterioles (PA0-1) were more likely to dilate to visual stimulus than higher order penetrating arterioles (PA2+) and capillaries (C1+) ($p = 2e-8$).

(H) Stimulus-dependent vascular dilation responses were then averaged across each vessel and superficial and deep penetrating arterioles were compared with the capillary bed (PA0 nTrials = 90, nVessels = 18; PA1 nTrials = 73, nVessels = 15; PA2+ nTrials = 48, nVessels = 16; C1+ nTrials = 277, nVessels = 90; data including non-responsive trials is shown in **Supplementary Figure 6**). The maximum dilation differed across vessel categories ($p = 0.05$; Kruskal-Wallis test), being borderline significantly smaller in deep PA sections than either superficial arteriolar segments or the capillary bed ($p = 0.06$ vs PA0-1 and C1+). Horizontal gray lines on violin plots show median (solid line) and interquartile range (dotted lines), and statistical comparisons of vessel responsivity rates were made using a Chi-square test with Fisher's *post hoc* comparison, and of dilation peaks using a Kruskal-Wallis test with *post hoc* Wilcoxon rank sum tests to assess pairwise comparisons.

points for elastin and α SMA – a transition in nestin expression. Furthermore, pericyte morphology as assessed from ISD does not alter abruptly around these transition points, suggesting that gradual changes in morphology along the vascular network are superimposed upon multiple sharp changes in protein expression levels. In both large and small vessels, branch points are also functionally specialized, having denser pericyte coverage than adjacent up and downstream vessels. Conversely, we did not find any transitions in expression of classic pericyte markers PDGFR β and NG2, or perivascular nNOS levels, which were expressed from the pia to the mid-capillary bed. Finally, *in vivo*, below their second branch, penetrating arterioles showed similar neurovascular coupling and vasomotion to capillaries, having a lower frequency of dilation or vasomotion compared to the first two segments of the penetrating arterioles.

Thus overall, our data support a view of the vascular network whereby sharp distinctions between different vascular segments do not exist, but rather different functions transition at different positions within the vascular tree, within classic vascular categories such as “arterioles” and “capillaries” as well as between them. This means that vascular function overall changes gradually across the vascular network, including along classically defined single “vessel types” such as diving arterioles.

Expression of Pericyte Markers Throughout the Vascular Network

We found PDGFR β and NG2 to be expressed throughout the arteriole-capillary vascular network. Both are classically considered to be pericyte markers (Winkler et al., 2010; Armulik et al., 2011), but our results are consistent with the described effects on smooth muscle cells of PDGFR β gain-of-function mutations, which increase leukocyte accumulation in the aorta (He et al., 2015). Single cell RNA seq analyses also support the more widespread expression of both PDGFR β and NG2, with mRNA transcripts found in smooth muscle cells as well as pericytes (Vanlandewijck et al., 2018), albeit at moderately lower levels.

The function of these two “marker” proteins may be different depending on their vascular location. NG2 is known to be important for neovascularization and stabilization of newly formed vessels, *via* the interaction of NG2 with integrins and growth factor receptors on the same and other cells (Stallcup, 2018). In capillary pericytes of the mature vasculature, however, it promotes the formation of new capillaries through angiogenesis, whereas in larger vessels it instead may promote arteriogenic remodeling of vessel diameter (Rundek and Della-Morte, 2015). Such remodeling can occur after decreased tissue oxygen and, consistent with widespread NG2 expression, cells in surface and penetrating arterioles, and the capillary bed, have been

found to proliferate after cerebral ischemia (Wei et al., 2001). PDGFR β also functions differently in arteries compared to smaller vessels. In culture, pericytes but not smooth muscle cells shed PDGFR β in response to stress (Sagare et al., 2015) and mutations that block PDGFR β signaling reduce pericyte number and increase capillary leakiness, without affecting smooth muscle cells (Nikolakopoulou et al., 2017). Conversely, inhibition of PDGFR β pharmacologically or using siRNA preserves cerebral arterial smooth muscle cells and arterial vascular tone after sub-arachnoid hemorrhage (Shiba et al., 2012; Wan et al., 2019), highlighting a potential pathophysiological contribution of PDGFR β signaling to arterial smooth muscle cells.

Thus both NG2 and PDGFR β are expressed throughout the cerebral microvasculature, but differ functionally depending on their location, presumably due to differences in expression levels of other proteins that are localized to different parts of the vascular network.

nNOS Is Expressed Around Arterioles and Capillaries

Given the involvement of nNOS-derived NO in arteriole but not capillary neurovascular coupling (Mishra et al., 2016), we expected to observe differential expression of nNOS along the vascular network. However, while nNOS was expressed at greater levels around vessels than in the parenchyma, this occurred to a similar degree for all vessels studied (up to 3rd branch order) and not just the diving arterioles. NO can control the electrical coupling of pericytes in the retina (Kovacs-Oller et al., 2020) as well as the production of other vasoactive molecules such as 20-HETE (Liu et al., 2008; Hall et al., 2014), so neuronally derived NO may be released onto all vessels but play a different role at arterioles than capillaries, generating a dilation in the former and modulating the response in the latter. A gradient of expression of the other constitutive NOS isoform, endothelial NOS, has not been reported, and RNA Seq data suggests it is expressed at similar levels in arterial, capillary and venous endothelial cells (Vanlandewijck et al., 2018).

Correspondence of Elastin and α SMA Labeling to Transitions in Physiological Responses

As previously described, elastin and α SMA labeling in the vascular wall both label arterioles (Shen et al., 2012; Grant et al., 2019) with α SMA labeling extending into the capillary bed (Grant et al., 2019; Chow et al., 2020; Thakore et al., 2021). Here we show these termination points are distinct and non-overlapping, occurring at significantly different branch orders and with different pericyte morphologies as assessed by ISD. The location of elastin labeling quite well matches locations

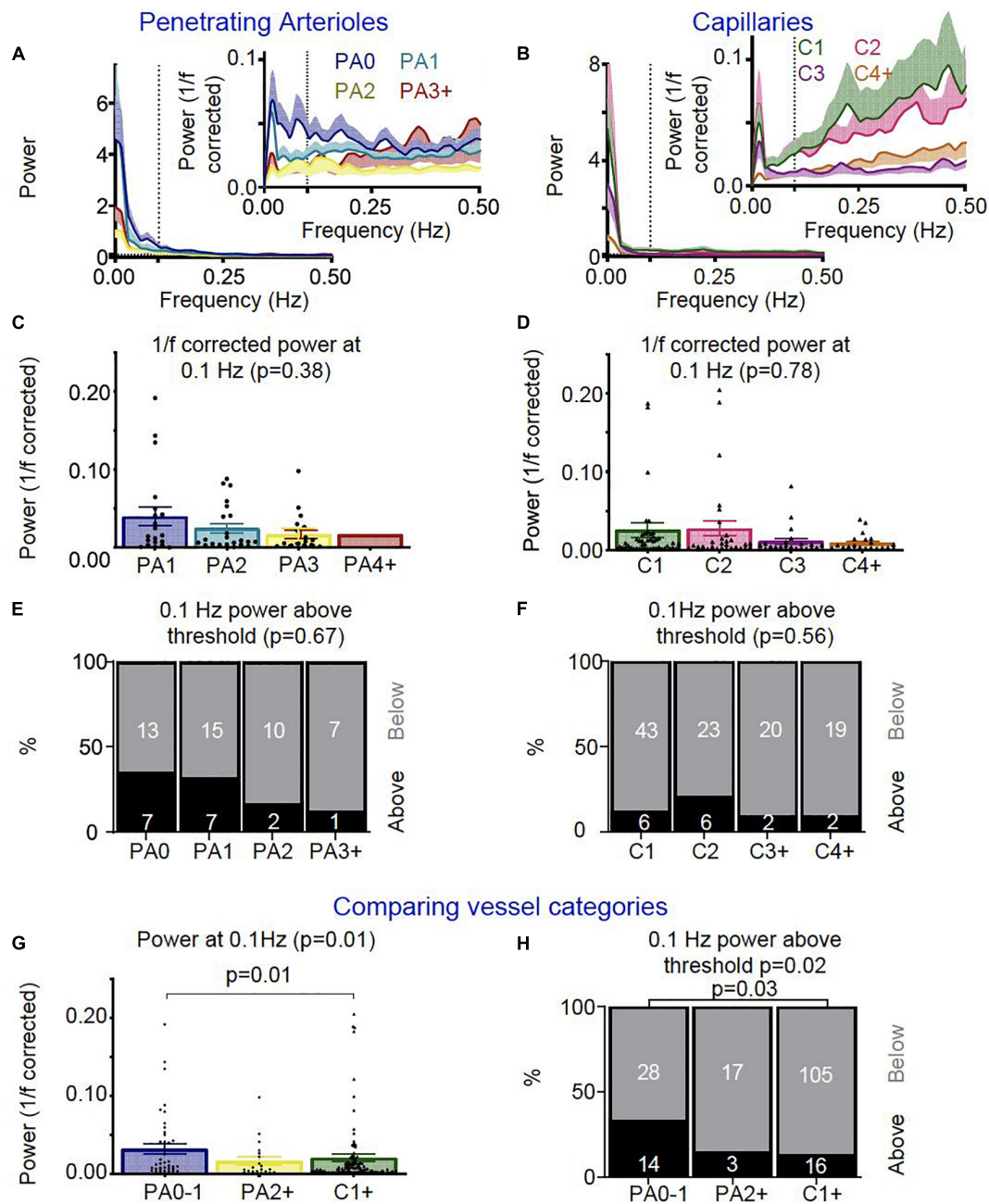


FIGURE 9 | *In vivo* vascular vasomotion responses separated by vessel segment. Average power spectra of **(A)** penetrating arteriole (PA0 nVessels = 20, PA1 nVessels = 22, PA2 nVessels = 12, PA3 + nVessels = 8) and **(B)** capillaries (C1 nVessels = 49, C2 nVessels = 29, C3 nVessels = 22, C4 + nVessels = 21) were separated by branch order for raw traces (left) and 1/f corrected traces (right insert). No significant differences were seen in the 1/f corrected power at 0.1 Hz between **(C)** penetrating arteriole or **(D)** capillary segments. Error bars represent mean \pm SEM, and power at 0.1 Hz was compared between individual vessels using a Kruskal Wallis test. **(A)** threshold was set for assessing the number of vessels which showed high 1/f corrected power at 0.1 Hz (threshold: $0.5 \times$ standard deviation across all vessels' 0.1 Hz 1/f corrected power values), and no significant differences were seen between **(E)** penetrating arteriole or **(F)** capillary vascular segments in the ratio of vessels with higher 1/f corrected power at the vasomotion frequency (numbers in bars represent individual vessels). We then compared vessel responses between the penetrating arterioles (PA0-1 nVessels = 42, PA2 + nVessels = 20) and capillary network (nVessels = 120). As for neurovascular coupling responses, superficial (PA0-1) and deep (PA2 +) arterioles were compared with the capillary bed (C1 +). **(G)** Average 1/f corrected power at 0.1 Hz was different across these vessel groups, with superficial arterioles showing significantly more power at 0.1 Hz than capillaries ($p = 0.01$; Wilcoxon rank sum pairwise comparisons). **(H)** The lowest order penetrating arterioles (PA0-1) also had more vessels with higher 1/f corrected power at 0.1 Hz than other categories ($p = 0.01$), specifically when compared to the capillaries ($p = 0.03$). Statistical comparisons of the number of vessels with higher power in the vasomotion range were made using a Chi-square test with Fisher's *post hoc* comparison, and of power at 0.1 Hz using a Kruskal Wallis test with Wilcoxon rank sum pairwise *post hoc* tests.

where we see transitions in physiological responses - the second branch off the diving arteriole - below which responses seem more like those in the capillary bed. Of our 12 elastin-labeled vessels, 8 terminated on the penetrating arteriole (75%), of which 5 (42% of the total) terminated before the first branch from the arteriole, i.e., upstream of where we observe a change in response frequencies. Thus, arteriole elastin labeling broadly, but not tightly, corresponds with the superficial part of the diving arteriole where neurovascular dilations and vasomotion were most frequently observed.

Though associated with contractile ability, the termination point of α SMA does not, however, correlate very well with the size or frequency of neurovascular coupling responses or vasomotion, we observed *in vivo*. α SMA universally terminated beyond the penetrating arteriole, but we found no differences in neurovascular response frequency, dilation size or vasomotion between different capillary branching orders (1-4 +), though many fewer of these smaller vessels express α SMA. Furthermore, responses were equally frequent in α SMA-expressing deep sections of the penetrating arteriole as in the capillary bed, and capillary dilations were actually larger than these deep arteriole dilations. This is at odds with previous findings, where dilations in ≥ 4 th order vessels were substantially smaller than higher order vessels in whisker barrel cortex of awake mice (Rungta et al., 2021). The reasons for this are unclear. Firstly, we compare both response frequency and response sizes, whereas these two measures are conflated in Rungta et al.'s paper. However, as we also saw similar response frequencies and sizes across the capillary bed, this cannot explain why we do not see smaller responses in ≥ 4 th order vessels. The cortical area is different (visual vs. somatosensory), and the degree of neuronal stimulation might be different (whole field drifting gratings vs. a single whisker deflection), which could perhaps have an impact on neurovascular coupling. Another potential cause of these differences is the method of detecting vascular diameter. We used xy images from which we calculate the diameter perpendicular to every point of a small length of a vessel's axis, thus averaging across space (12-109 pixels, or 2.4-22 microns), while Rungta et al. used line scans of vessels to measure vascular diameter at a single position. The spatial smoothing we used in this paper is likely to give us a higher sensitivity to small deflections in diameter.

Previous work in anesthetized animals, has also found different vascular segments to show different time courses of dilation, with deeper sections of arterioles or first order capillaries responding faster than superficial arteriolar segments (Tian et al., 2010; Hall et al., 2014; Rungta et al., 2018). Our data from awake mice did not show this, with similar response kinetics between vascular segments.

Transitional Segment?

The vascular segment between the penetrating arteriole, or end of elastin labeling, and the end of α SMA labeling has often been termed a "precapillary arteriole" or a "transitional segment" (e.g., Rungta et al., 2021), representing a region where vascular function transitions between arteriole and capillary. However, our *in vivo* data suggests that this segment, corresponding

roughly to branch orders 1 to 3, is not (in our hands) where transitions of contractile behavior occur. Other transitions do occur in this zone: We found endothelial expression of the intermediate filament protein nestin extends out of the capillary bed to a position between the termination points of elastin and α SMA. In the retina, calponin, filamentous microtubules, α SMA, filamentous actin and myosin heavy chain were all also found to change expression levels across this section (Gonzales et al., 2020). However, these all transitioned at different positions, calponin terminating on the arteriole, microtubules on branch 1 and α SMA on branch 2, with filamentous actin and myosin heavy chain gradually decreasing in expression levels from branch orders 0 to 4 (beyond which vessels were not studied). These progressive changes in function across a number of markers fit with the gradual changes in mural cell morphology or ISD we observed before and after the termination points of elastin and α SMA, at roughly branch orders 0 and 3, respectively. Thus this "transitional zone" is not uniform, with a single type of vascular cell and, crucially, is not the only region where such transitions of function are occurring, as similar transitions in vasomotion, neurovascular coupling, and mural cell morphology also occur when descending the penetrating arteriole: Neurovascular coupling responses and vasomotion were observed more frequently in superficial segments of diving arterioles than in either downstream capillaries or deep sections of the arterioles, which were similar to each other in response characteristics. Correspondingly, mural cells lost their annular smooth muscle cell morphology to gain a distinct soma and processes at lower reaches of the diving arterioles. This suggests arteriolar mural cells can be pericytes, unlike has been argued (Hartmann et al., 2021a).

This data suggests that the division of the vascular network into four functional segments: arterioles with SMCs, a transitional zone with ensheathing pericytes, capillaries with capillary pericytes and venules with venular SMCs (Hartmann et al., 2021a), while helpful in discussing broad functional changes across the network, is overly simplistic and neglects the gradual transition in functions that occurs. Indeed, multiple other functional transitions occur at other positions in the vascular tree, including between the pial and penetrating arterioles: Penetrating arterioles exhibit higher contractile tone at low intravascular pressures than pial arterioles (Longden et al., 2016). Neurovascular response sizes are also smaller, possibly because of the lower external pressure on the surface vessel compared to penetrating arterioles which are surrounded by brain tissue (Gao et al., 2015).

Branch Points May Be Functionally Specialized

Our data suggest that pericytes exist at a higher density at branch points than on surrounding vessel lengths, and this clustering occurs to a similar degree on small and large microvessels. This suggests some functional specialization at branch points. 90% of branch points in the transitional segment were previously found to have a pericyte at that location

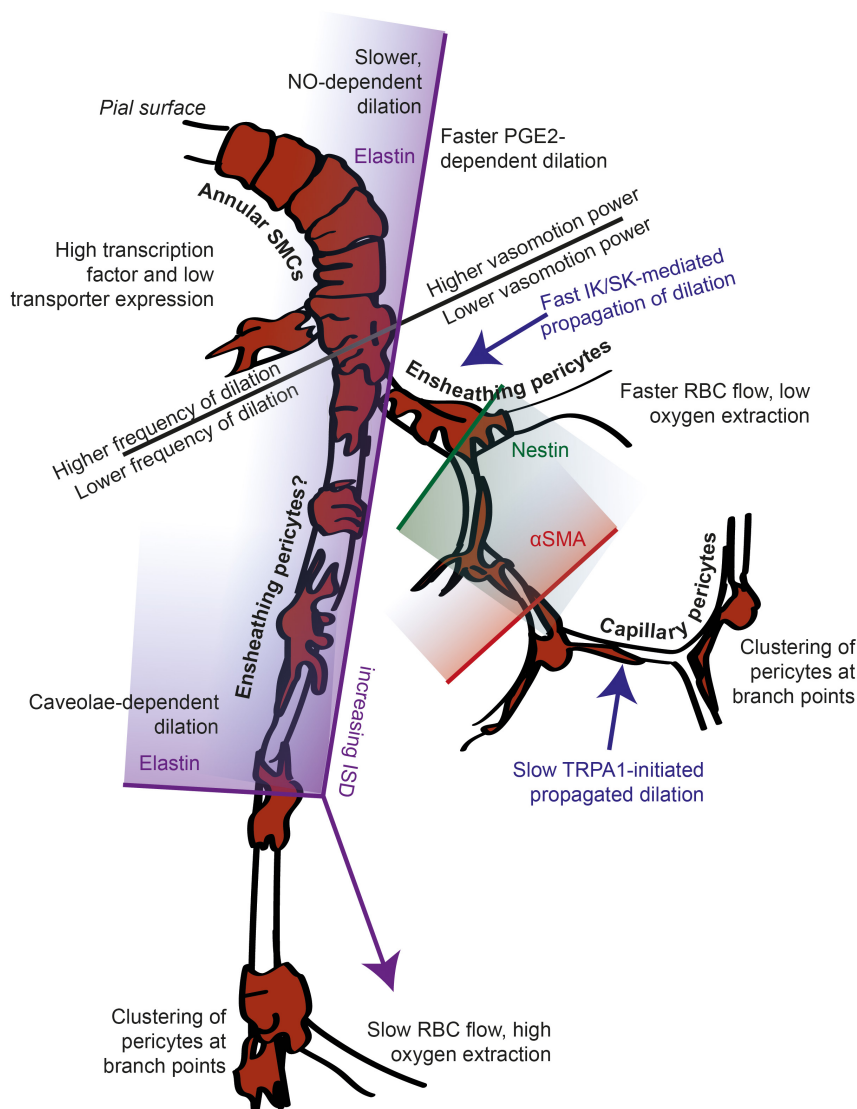


FIGURE 10 | Summary of known functional transition points across the arteriole-capillary axis. The location of many functional transitions remains unknown, indicated by label placement in a general zone, without an arrow (e.g., between slow and fast propagated dilation, and expression levels of transporters vs. transcription factors). Gradient shading shows that the labeled marker is expressed in the vascular zone to the shaded side of the termination point indicated. Current nomenclature for mural cells is indicated, with mural cells on the arteriole suggested to be ensheathing pericytes rather than smooth muscle cells due to their pericyte morphology. This is backed up by the similar behavior of deep arteriole sections compared to the capillary bed, though there are other differences (e.g., elastin labeling, caveolae) as indicated.

compared to only 45% of more distal branch points (Gonzales et al., 2020). This corresponds with the increase in pericyte ISD we report here, but our data suggest that even the distal branch points are functionally specialized, as they have a shorter ISD than surrounding non-branch point regions. The pericytes at proximal branch points had calcium sparks that corresponded with selective constriction of individual branches, suggesting branch points serve to direct blood flow to active neurons (Gonzales et al., 2020). Distal branch points (> 4th branch order) were not found to be contractile, and calcium changes in these pericytes were not reported. As our data suggest these distal capillaries do dilate to a similar degree

as the proximal branches, it would be valuable to study whether these branch points' pericyte calcium changes also correspond to changes in vascular diameter of the different downstream branches.

Transcriptomic Gradients Could Illuminate Transitions in Whole Range of Functions

Our data and the wider literature currently support multiple transitions of function at different positions of the vascular bed, including down diving arterioles, and along increasingly

branching capillaries. Contractile function and neurovascular coupling are the functions most widely studied, but transitions at different locations are also seen in oxygen supply as well as expression of transcription factors, transporters, regulators of angiogenesis and immune regulators. However, these studies are all limited by a low capacity to look at different functions. Rather than focusing on individual vascular functions, single cell RNA Seq has the potential to illuminate the whole range of gene expression differences that exist across the vascular network (He et al., 2018). Clustering of vascular endothelial and mural cells has revealed that endothelial gene expression changes gradually, suggesting that there are likely not simultaneous transitions in expression of many genes at the same point on the vessel. More abrupt transitions between smooth muscle cell and pericytes have been reported (Vanlandewijck et al., 2018), which may fit with the sharp transitions observed in some vascular features (e.g., the shift in vessel coverage from ensheathing to mesh pericytes). However, this study also shows large numbers of genes that are expressed in wider zones of the vasculature, e.g., pericytes and arteriolar but not arterial SMCs, supporting the existence of multiple transition points. Indeed, another RNA Seq study identifies multiple pericyte clusters suggestive of multiple functional transition points (Zeisel et al., 2018). Future studies could link single cell RNA seq data to specific spatial locations within the vascular network, to illuminate this issue further. Such data would allow us to better understand the vascular regions to target with pharmacological interventions, given Alzheimer's disease and stroke are vasculature-degenerating conditions that differentially affect the various parts of the vascular network.

CONCLUSION

Our data, and the literature, support the existence of multiple transition points in vascular function, different proteins being expressed at overlapping sections of the vascular network, and properties of neurovascular and vasomotion response rates, sizes and timings varying at different places in the network (summarized in **Figure 10**). These various functional transitions are superimposed on gradually changing mural cell morphologies across the vascular tree, which show specializations at branch points. Thus, while categorization of vessels or mural cells may be a useful simplification in some circumstances, it is important to remember that, for example, an upstream ensheathing pericyte is not identical to a downstream ensheathing pericyte, nor to one on a branch point. Understanding where and how different vascular functions (e.g., oxygen and nutrient supply, waste clearance, immune regulation) are supported across the vascular tree is vital to understand how different sections are impacted, and could be targeted, during disease, but will likely require approaches such as spatially-localized RNA Seq to identify how the transcriptome as a whole alters across the cerebral microvascular network.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found at Figshare.com, doi: 10.25377/sussex.17840939.

ETHICS STATEMENT

All experiments were carried out in compliance with the UK Animal Experiments (Scientific Procedures) Act 1986 after approval of the local University of Sussex or UCL Local Ethics Committees and under project or personal licences granted by the UK Home Office.

AUTHOR CONTRIBUTIONS

KB, MH-H, DA, and CH collected the data. KS, KB, MH-H, DA, and CH analyzed the data. KS, SA, OB, and CH wrote the manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnagi.2021.779823/full#supplementary-material>

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