



Factors controlling permeability of the blood–brain barrier

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Received: 1 June 2015/Revised: 15 September 2015/Accepted: 17 September 2015/Published online: 24 September 2015
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Abstract As the primary protective barrier for neurons in the brain, the blood–brain barrier (BBB) exists between the blood microcirculation system and the brain parenchyma. The normal BBB integrity is essential in protecting the brain from systemic toxins and maintaining the necessary level of nutrients and ions for neuronal function. This integrity is mediated by structural BBB components, such as tight junction proteins, integrins, annexins, and agrin, of a multicellular system including endothelial cells, astrocytes, pericytes, etc. BBB dysfunction is a significant contributor to the pathogenesis of a variety of brain disorders. Many signaling factors have been identified to be able to control BBB permeability through regulating the structural components. Among those signaling factors are inflammatory mediators, free radicals, vascular endothelial growth factor, matrix metalloproteinases, microRNAs, etc. In this review, we provide a summary of recent progress regarding these structural components and signaling factors, relating to their roles in various brain disorders. Attention is also devoted to recent research regarding impact of pharmacological agents such as isoflurane on BBB permeability and how iron ion passes across BBB. Hopefully, a better understanding of the factors controlling

BBB permeability helps develop novel pharmacological interventions of BBB hyperpermeability under pathological conditions.

Keywords Inflammatory mediators · Integrins · Annexins · Agrin · MicroRNAs · Anesthetics

Introduction

The blood–brain barrier (BBB) refers to an interface between the brain parenchyma and the systemic circulation. It regulates the entry of nutrients, vitamins, ions, and other molecules into the brain, protects the brain from harm caused by toxins and pathogens, and is critical to normal brain functions [1]. BBB consists of a non-fenestrated layer of brain microvascular endothelial cells (BMECs). The BMECs are primary components of BBB and play a critical role in regulating BBB permeability by a complex tight junction (TJ) structure [2, 3]. A more recent and complete view of BBB is that it refers to a neurovascular unit of BMECs interacting with other brain cells, such as astrocytes, pericytes, neurons, and microglial cells [3, 4]. It was clearly demonstrated as early as in 1987 that astrocytes contribute to BMEC-mediated regulation of BBB permeability and maintain cerebrovascular integrity [5]. It has been shown that astrocytes release sonic hedgehog that up-regulates TJ proteins in BMECs [5, 6]. However, activated astrocytes seem to disrupt endothelial barrier integrity by releasing biologically active molecules that activate ubiquitin proteasome and degrade TJ proteins [7]. In addition, microglia, pericytes and neurons maintain BBB integrity and function via contributing to cerebral vessel stability and modulating signaling mediators [8–11]. Disruption of BBB integrity and functions is involved in a growing list of

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brain disorders (Table 1). Particularly, diminished BBB function is an early event of multiple sclerosis (MS) and amyotrophic lateral sclerosis [12–16], chronic inflammatory disorders of the central nervous system (CNS), and a hallmark of stroke [17–20]. In this review, factors controlling BBB permeability are discussed in three chapters, junctional proteins at the BBB, proteins of the basement membranes of the BBB, and signaling mediators, in line with their contributions to neurodegenerative diseases and neurological disorders (Table 1). In addition, the impact of pharmacological agents such as the volatile anesthetic isoflurane on BBB permeability is reviewed. Recently, how iron passes across the BBB has drawn much attention; progress regarding this issue is also briefly reviewed. Understanding the molecular mechanisms by which these factors induce disruption of BBB integrity and function may lead to develop safe and effective therapeutic approaches to protect or restore the BBB integrity in many brain disorders.

Junctional proteins at the BBB

TJ proteins

Brain microvascular endothelial cells have a highly specialized phenotype characterized by the presence of intercellular TJs with high P-face association [21], which restrict molecules from moving between the blood and the brain and are responsible for the severe restrictions on diffusion. Transmembrane TJs consist of three major integral proteins—claudins, occludin, and junction adhesion molecules. ZOs are cytoplasmic membrane-associated accessory proteins that connect the cytoplasmic tails of claudins and occludin to the actin cytoskeleton to maintain the TJ structure. Altering TJ-associated proteins changes BBB permeability [22]. Occludin, claudin-5, and ZO-1 are considered sensitive indicators of normal and disturbed functional states of the BBB. As described in the latter paragraphs, many signaling mediators and pharmacological

Table 1 Mechanisms underlying BBB disruption in brain disorders

Disorders	Mechanisms
Alzheimer's	↓ ZO-1 by MMPs [146, 154, 225], cytokines [172], and ROS [226, 227] Loss of pericytes by NF-κB [12] and APOE4 [154] ↓ Agrin and ↑ agrin insolubility [109, 111, 112]
Parkinson's	↓ Occludin and ZO-1 by MMP-3 [141] ↑ MMPs [140, 141]
Huntington's	Degradation of endothelial basal lamina and ↓ TJ proteins (claudin, occludin and ZO) by MMP-2/9 [142]
ALS	↓ TJ proteins expression (ZO-1, occludin, and claudin) by MMP-2/9 [15, 16]
Migraine	↓ TJ proteins expression (ZO-1, occludin, and claudin) by MMP-2 [143] and MMP-9 [144, 145]
Traumatic brain injury	↓ TJ proteins expression (occludin, claudin-5, and ZO-1) by iNOS [164] and MMP-9 [100, 129, 149, 153]
Stroke	↓ TJ proteins expression (occludin, claudin-5, and ZO-1) by MMP-2 [20], MMP-9 [73, 100, 129, 130, 148, 151, 156, 158, 163], ROS [19, 23, 130, 156, 158, 228], VEGF [181, 182], pericyte-derived VEGF [11], and astrocyte-derived VEGF [54–56, 183] ↓ TIMP-1/2 [73, 162, 163] ↓ Loss of α6β4 and β1 integrins [54] ↓ Agrin [110] ↓ AA1 [73] ↑ miRNA-21 → ERK-mediated upregulation of MMP-9 [196] ↑ miRNA-15a [197]
Intracerebral hemorrhage	↑ miRNA-130a → ↑ MMP-2/9 expression [159] ↓ TJ proteins expression by MMPs induced by NF-κB [161]
Multiple sclerosis	↓ α4β1 integrin [57, 58, 229] ↑ MMPs and NO [175] ↓ AA2 and claudin-1 by miRNA-155 [28]
Japanese encephalitis	↓ ZO-1 and claudin-5 by VEGF, IL-6, and MMP2/9 [7]
Autoimmune encephalomyelitis	↓ Claudin-5 by VEGF [180]

ALS amyotrophic lateral sclerosis, ↓ decrease in protein expression or activity, ↑ increase in protein expression or activity

agents result in BBB dysfunction through modulating the expression of TJ proteins or TJ related proteins, particularly occludin, claudin-5, and/or ZO-1 [23, 24]. For example, claudin-5-deficiency induced size-selective loosening of the BBB [25], and we have observed that hyperglycemia caused endothelial hyperpermeability by altering the expression and distribution of ZO-1 in cultured BMECs [26] and in brain microvessels from diabetic mice (unpublished data). It is noteworthy that the role of claudin-1 in BBB permeability has recently been pursued by several groups. Although the presence or absence of claudin-1 in BMECs is still controversial [27, 28], ectopic expression of claudin-1 in Tie-2 tTA/TRE-claudin-1 double transgenic C57BL/6 mice was able to prevent BBB leakage in chronic experimental autoimmune encephalomyelitis [27], indicating the sealing function of claudins. Loss of claudin-1 at the BBB has been associated with barrier dysfunction in human glioblastoma multiforme [29] and hepatitis C infection [30]. In addition, the cross-talk between adherens junctions and TJs in maintaining barrier integrity is an important and emerging topic (see recent review [31] by Tietz and Engelhardt for details).

Integrins

Integrins are cell adhesion proteins that either link to the extracellular matrix (ECM) or mediate dynamic contacts to other cells via binding to cell adhesion molecules [32]. They widely exist and vary among species. Integrins are heterodimers and consist of α and β subunits that bind to each other non-covalently. There are over 18 α and 8 β subunits, which assemble over 24 distinct integrin proteins. Integrins have various cellular functions and can activate many intracellular signaling pathways [33]. Integrins express at the BBB at a high level and play important roles in the interactions between astrocytes and the endothelium. Integrin receptors, working with dystroglycan, contribute to endothelial cell–matrix adhesion and maintain adjacency between abluminal endothelial cell face to astrocyte end-feet [34, 35]. Specifically, $\alpha\nu\beta 8$ integrin has been demonstrated to play an essential role in formation of the adhesive interaction, which was expressed by astrocytes and induced transforming growth factor- β (TGF- β) to stabilize the endothelium [36]. In a primary culture, $\alpha\nu\beta 5$ and $\alpha\nu\beta 8$ induced astrocyte adhesion to vitronectin and migration [37]. $\alpha\nu\beta 8$ derived from neuroepithelial cells was able to activate TGF- $\beta 1$ and TGFBR1-ALK5-Smad3 signaling in endothelial cells to maintain normal vascular patterning and suppress endothelial sprouting [38, 39]. $\alpha 6\beta 4$ expressed by astrocytes was reported to strengthen the attachment between end-feet and basal lamina in order to maintain apposition to endothelium of microvessels [35]. $\alpha\text{phav}\beta 3$ integrin promoted endothelial cell proliferation

and survival through mediating capillary endothelial cell adhesion to fibronectin, which was introduced in the progress of angiogenesis [40]. It is clear that $\beta 1$ integrins (the $\alpha\beta$ heterodimers with $\beta 1$ subunit) play a key role in forming the matrix adhesion between cerebral endothelial cells as a part of the BBB. In order to stabilize the TJ integrity, $\beta 1$ integrins mediate the attachment of ECM to endothelial cells [41]. Expressed by endothelial cells along with other receptors, $\beta 1$ integrins have an important function regulating the permeability of BBB [35]. In addition, in the process of CNS development, an obvious increased $\beta 1$ integrin expression is a sign of cerebral blood vessel maturation [42].

$\beta 2$ Integrins are a group of integrins that is essential for immune cell trafficking. Under inflammatory conditions, it has been found that $\beta 2$ integrins are involved in neutrophil extravasation across inflamed BBB. There are many isoforms of $\beta 2$ integrins such as lymphocyte function-associated antigen (LFA-1) and macrophage-1 antigen (Mac-1) [43, 44]. Many studies have investigated the roles of $\beta 2$ integrins and their ligands, such as intercellular adhesion molecule 1 (ICAM-1), ICAM-2, and junctional adhesion molecule-A, in immune cell extravasation across the BBB under inflammatory conditions [45]. The adhesive interaction between immune cells and BMECs during inflammation requires $\beta 2$ integrin activation to engage with their endothelial ICAMs. Both LFA-1 and Mac-1 with their endothelial ligands (ICAM-1 and ICAM-2) mediate neutrophil crawling on the BBB during inflammation [43, 45]. It seems that LFA-1 mediates shear-resistant arrest of neutrophils by ICAM-1 on the flamed BBB while neutrophil polarization is subsequently mediated by Mac-1 [45]. It was suggested that G protein-coupled receptors (GPCR) mediated $\beta 2$ integrin activation [46] because GPCR inhibition resulted in reduction of neutrophil arrest and polarization. In the complete absence of either $\beta 2$ integrins or their ligands, crawling of neutrophils on the inflamed BBB was abrogated [45].

Alpha 4 integrins are also involved in mediating the migration of immune cells across the BBB during neuroinflammation [47, 48]. Alpha4 integrin family consists of several isoforms such as $\alpha 4\beta 1$ [very late antigen-4 (VLA-4)] and $\alpha 4\beta 7$ [lymphocyte Peyer's patch adhesion molecules (LPAM-1)]. It has been well known that the interaction between $\alpha 4$ integrins and vascular cell adhesion protein 1 (VCAM-1) contributes to T cell recruitment across BMECs. Lacking $\alpha 4$ integrins or VCAM-1 activity reduced the adhesion of encephalitogenic T cell blast to the BBB [49, 50]. It has been reported that VLA-4 predominantly regulated leukocyte extravasation to the inflamed brain while LPAM-1 had a role in immune cell trafficking to other inflamed systems such as the gut [50, 51]. The integrin $\alpha 4$ expression was up-regulated during MS brain

lesions [52]. Moreover, recombinant VCAM-1 has been shown to bind to $\alpha 4$ integrins and to compromise BBB function in vitro through a mechanism that is mediated by activation of intracellular signaling pathways including MAP kinase. Pharmacological inhibitors (e.g. natalizumab) of the $\alpha 4$ integrin-VCAM-1 interaction protected the BBB during inflammatory conditions via blocking leukocyte adhesion to BMECs [51, 52]. Furthermore, preventing the interaction between $\alpha 4$ integrins and VCAM-1 by anti- $\alpha 4$ integrin monoclonal antibodies reduced inflammatory cell recruitment across the BBB during experimental autoimmune encephalomyelitis (EAE) in vivo [47]. Early treatment of EAE mice with a synthetic VLA-4 antagonist significantly reduced the clinical severity of the EAE condition [53].

Decreased expression of integrins has been found in neurological disorders and is considered to contribute to abnormal BBB permeability in such conditions. For example, it was shown that $\alpha 6\beta 4$ was lost rapidly from astrocyte end-feet after focal ischemia [54]. Endothelial expression of $\beta 1$ integrin, accompanied with the expression of subunits $\alpha 1$, $\alpha 3$, and $\alpha 6$, was significantly decreased within ischemic core 2 h after middle cerebral artery occlusion [55, 56]. In addition, Rice et al. and von Andrian et al. demonstrated that the counter-receptor on the vascular endothelium interacting with the $\alpha 4\beta 1$ integrin on leukocytes had a significant effect on MS [57, 58]. Taken together, the above-mentioned findings suggest that normalizing integrin expression would be a potential approach to secure BBB integrity in these brain disorders.

Annexins

Annexins are Ca^{2+} and phospholipid-binding proteins that regulate cellular Ca^{2+} concentrations and membrane organization and traffic. Among many annexins, annexin A1 (AA1) is the most investigated. AA1 is a protein of 37 kDa and has been shown to function in intercellular trafficking due to its relation to membrane components and cytoskeletal proteins [59]. It was reported that AA1 possessed anti-inflammatory effects by regulating macrophage phagocytosis and neutrophil migration [60]. It might also play a significant role in differentiation [61], plasma membrane repair [62], proliferation [63], and apoptosis [64]. In the brain, AA1 is expressed in pericytes [65] and BMECs [66], where it may regulate prostaglandin (PG) E2 production [67]. AA1 is also strongly expressed in the ependymal cells of rats [68] and humans [69]. Glial expressions of AA1 are mainly seen in microglia [65], but not in oligodendrocytes [70, 71]. In CNS, the protective role of AA1 has been observed in neurodegeneration [72] and ischemia [73].

There is compelling evidence supporting AA1 as a critical physiological regulator of BBB integrity. First,

endogenous AA1 regulates cytoskeleton by binding to actin. It can directly bind to β -actin, especially the cytosolic form rather than the fibrillar one. With predominant binding to soluble actin, AA1 may have a potential function in stabilizing actin oligomers before forming cytoskeleton [74]. It also works as a linkage between cytoskeleton and plasma membrane and helps to facilitate TJ functions. Second, AA1 signaling inhibits GTPase RhoA [75] that is able to lead to a downstream reaction, destabilizing actin cytoskeleton and enhancing paracellular permeability [76]. Third, AA1 knockout mice have a significant decrease in occludin and vascular endothelial (VE)-cadherin and an increase in BBB permeability although levels of claudin-5 and ZO-1 remain the same [75]. Furthermore, neutrophil extravasation in an animal model of stroke can be blocked by AA1 active fragments [73], corroborating AA1's protective effect on BBB. In addition, AA2 has been shown to regulate endothelial cell morphology and junctional integrity via its association with VE-cadherin because silencing AA2 expression increased phosphorylation of VE-cadherin and BBB permeability in vitro [28, 77].

Vascular endothelial cell-specific phosphotyrosine phosphatase

Vascular endothelial cell-specific phosphotyrosine phosphatase (VE-PTP) is a member of the receptor-type protein tyrosine phosphatase family that is exclusively expressed in endothelial cells. VE-PTP plays a critical role in embryonic development and angiogenesis. Mice lacking VE-PTP exhibited severe vascular malformation and died during early embryonic period [78, 79]. Furthermore, VE-PTP, working with its adhesive VE-cadherin partner, regulates the endothelial barrier function by an increased VE-PTP/VE-cadherin interaction [80]. Dissociation of VE-PTP from VE-cadherin is mediated by many inflammatory factors such as VEGF and then results in destabilization of endothelial cell integrity. Indeed, down-regulation of VE-PTP expression increased trans-endothelial permeability due to reduction of VE-cadherin adhesiveness [81, 82]. Moreover, cytokines (e.g. IL-1) that inhibit PTP activity increased BBB permeability [83]. Collectively, these findings suggest that modulating PTP activity could be an approach to regulate BBB permeability properties.

Platelet endothelial cell adhesion molecule-1

Platelet endothelial cell adhesion molecule-1 (PECAM-1) is a transmembrane glycoprotein that is expressed in the intercellular junctions of the endothelial cells. PECAM-1 mediates angiogenesis in BBB development as well as migration and interaction of leukocytes to the endothelial

cells. These effects indicate that PECAM-1 is a crucial molecule in neuroinflammation [84, 85]. Mice lacking PECAM-1 had deficiency in their inflammatory responses and exhibited an increased ability of inflammatory cells to cross the BBB due to prolonged changes in the BBB permeability [86, 87]. In addition, PECAM-1 seems to have an indirect role in modulating leukocyte extravasation behavior via increased $\beta 1$ and $\beta 2$ integrin affinity. This increase in integrin affinity mediates adhesion of leukocytes to vascular endothelial cells and enhances their trans-endothelial migration during neuroinflammation [87–89].

Proteins of the basement membranes of the BBB

The BBB is composed of both cellular components and basement membranes (BMs). Two BMs exist between blood and brain, endothelial BM and parenchymal BM. Endothelial BM lining the capillaries consists of collagen IV, fibronectin, laminins, etc. [90]. Each of these membrane components contributes to BBB integrity [91]. Some cells such as pericytes are embedded within the endothelial membrane. Parenchymal BM is primarily formed by the end-foot processes of astrocytes and neurons, ECM, and laminins. The BMs line the perivascular space and act as a barrier for leukocyte migration into the CNS during inflammatory perivascular cuff conditions [92].

Laminins

Laminins belong to a family of high-molecular weight (~ 400 kDa) glycoproteins of the ECM. There are four major isoforms of laminins at the BBB, including laminins 1, 2, 8, and 10. Laminins 8 and 10 are secreted in the endothelial BM while laminins 1 and 2 are mainly secreted by astrocytes in the parenchymal BM [93–95]. It has been known that pericytes regulate the secretion of these laminins [96]. Results from studies performed on mice lacking the major $\gamma 1$ subunit of laminins support the notion that laminins play a crucial role in BM assembly. This targeted mutation resulted in early embryonic lethality due to defects in BM formation [97]. Yao et al. recently demonstrated that in conditional knockout mice and an acute adenovirus-mediated knockdown model, lack of astrocytic laminins induced BBB breakdown, possibly by increasing pericyte differentiation from the BBB-stabilizing resting stage to the BBB-disrupting contractile stage and decreasing TJ protein expression [98]. Moreover, it has been shown that decreased levels of laminins contribute to BBB damage caused by ischemia. Hamann et al. reported that ischemia significantly reduced the content of diverse basal lamina antigens including laminins [99]. Mostly recently, Cai et al. revealed that degradation of collagen IV

and laminins was accompanied by increased BBB leakage after recombinant tissue plasminogen activator treatments, indicating that the BM works as an important barrier for preventing BBB disruption and hemorrhagic transformation after thrombolysis in stroke [100].

Collagen IV and fibronectin

All three major cells at BBB, endothelial cells, astrocytes, and pericytes, secrete collagen IV and fibronectin. Collagen IV chains form a covalently stabilized polygonal framework. It is responsible for the mechanical resistance of the basal lamina and mediates BM stabilization and integrity through retaining laminins [101]. Fibronectin exists in two forms, insoluble glycoprotein dimer and soluble disulphide linked dimer. The insoluble fibronectin serves as a linker in the ECM and is an important component of the ECM. Fibronectin binds other ECM components such as collagen, fibrin, and heparan sulfate proteoglycans. Collagen IV- or fibronectin-mutated mice exhibited embryonic death due to defects in vascular development and loss of BM stability, respectively [101, 102]. Similar to laminins, collagen IV and fibronectin are susceptible to ischemic damage and contribute to BBB hyperpermeability in ischemic stroke [99]. Collagen IV has been reported to decrease significantly at 3 and 24 h after focal ischemic brain injury [103, 104]. The loss of basal lamina components including laminins, collagen IV and fibronectin during ischemic brain injury seems to be a result of protein degradation caused by plasmin, MMPs and cathepsins (see review for detail [105]).

Agrin

Agrin is an extracellular heparan sulfate proteoglycan and accumulates extensively in the BM of the brain microvasculature [106]. It may play a role in BBB development due to its presence around brain microvessels during embryonic development [107]. Many lines of evidence have revealed that agrin is involved in maintaining BBB integrity and function. It was shown that loss of agrin was correlated with the loss of TJ protein expression in glioblastoma [108], Alzheimer's disease (AD) [109], and cerebral ischemia [110]. Recently, the role of agrin in AD has attracted more attention due to its contribution to BBB dysfunction. Donahue et al. reported that a large fraction of the agrin in AD brains is insoluble in 1 % SDS while all the agrin in a normal brain is soluble [111]. The authors proposed that the insolubility was due to agrin's tight association with β -amyloid and that altered agrin expression in the microvasculature and the brain parenchyma contributed to the pathogenesis of AD. Rauch et al. stated that in mice lacking endothelial cell expression of agrin,

the level of aquaporin 4, a BBB-associated component, was reduced, and overexpression of agrin decreased A β deposition, indicating that agrin is important for maintaining BBB composition and influences A β homeostasis in mouse models of AD [112]. Most recently, Steiner et al. reported that agrin contributed to barrier characteristics of brain endothelial cells by stabilizing junctional localization of the adherent junction proteins VE-cadherin, β -catenin, and ZO-1 [113]. Their results demonstrated that agrin significantly enhanced the barrier characteristics of bEnd5 monolayers, accompanied by enhanced localization of VE-cadherin, β -catenin, and ZO-1, but not of claudin-5 and occludin. They further argued that agrin might stabilize these proteins at cell-to-cell junctions rather than inducing their enhanced-expression because protein levels of VE-cadherin, β -catenin, and ZO-1 did not change. Their argument was strengthened by results from an agrin deficiency mouse model that expressed mini-agrin, in which the vascular BMs had decreased junctional staining for VE-cadherin.

Signaling mediators that regulate BBB permeability

Cytokines

Cytokines are a group of polypeptides involved in inflammatory responses. These polypeptides include many families, such as interleukins (ILs) and tumor necrosis factors (TNFs), which are predominantly present in the brain during inflammation [114]. The normal activity of neuroinflammation is mainly to restore the homeostasis in the brain [115]. However, upon prolonged CNS inflammation, inflammatory responses may influence the BBB integrity and further result in a wide variety of CNS pathologies. In fact, many studies have recently demonstrated that neuroinflammation is a risk factor mediating BBB dysfunction in various brain disorders [116]. For example, early inflammation of the BBB was observed together with changes in permeability in experimental models of neurodegeneration [117]. Emerging evidence suggests that the chronic release of pro-inflammatory mediators from neurovascular cells is a major player in the disruption of BBB integrity in MS [118]. Although the mechanisms behind inflammation-induced BBB disruption and hyperpermeability are not fully understood, there is strong evidence that integrity of the BBB components can be threatened in response to pro-inflammatory mediators.

It has been revealed that the BBB disruption during neuroinflammation is largely associated with the release of numerous cytokines, such as IL-1, IL-6, TNF α , and eicosanoids [119–121]. Inflammation initiates cytokine

production mainly through activation of leukocytes, astrocytes, and microglial cells in the brain [122, 123]. Increase in alteration of TJ function and BBB permeability is strongly linked with elevation of the cytokines (IL-1, IL-6, and TNF α) in the brain. For example, Quagliarello et al. reported that an increase in the level of IL-1 induced meningitis and BBB breakdown in the rat [124]. Wang et al. demonstrated that IL-1 β increased BBB permeability via suppression of astrocytic sonic hedgehog production, leading to down-regulating TJ proteins such as claudin and occludin [125]. In addition, IL-6 and TNF α were able to elevate paracellular permeability in BMECs through a down-regulation of TJ proteins and an increase in reactive oxygen species (ROS) generation [126]. Furthermore, it was found that TNF α mediated BBB permeability via induction of cyclooxygenase-2 (COX2) and PG release in BMECs [127]. Results reported by Nishioku et al. showed that lipopolysaccharide (LPS)-activated microglia could release TNF α and result in BBB dysfunction [128]. HPI201 and minocycline, which prevented the up-regulation of TNF- α , IL-1 β , and IL-6, improved TJ protein expression and BBB integrity [129, 130]. In addition, TNF α up-regulated expression of chemokines and cell adhesion molecules (e.g. ICAM-1) in BMECs; and these molecules played a critical role in adhesion and migration of leukocytes across BMECs [131]. Most recently, Rosenberg et al. proposed that inflammatory cytokines that increase BBB permeability could be used as biomarkers for vascular cognitive impairment [132]. Taken together, it is clear that certain cytokines have the ability to increase BBB permeability through down-regulating TJ proteins as well as inducing other pro-inflammatory mediators such as PGs.

Eicosanoids

Eicosanoids are derived from arachidonic acid and include PGs and leukotrienes (LTs). They are released by COX1 and COX2 in response to inflammation [133]. In addition to the cytokines, these inflammatory mediators are predominant in CNS inflammation. Many studies have shown that PGE2 causes BBB damage and dysfunction. For instance, it was reported in 1998 that PGE2 contributed to BBB disruption during experimental rat meningitis [134]. A recent study showed that increased PG production facilitated the permeability of an in vitro model of BBB with human BMECs [135]. Moreover, studies have demonstrated that PGs enhanced BBB permeability in LPS- and TNF α -administered mice. Furthermore, Candelario-Jalil et al. demonstrated that COX inhibitors limited BBB disruption during neuroinflammation [136]. Besides PGs, LTs are also known to contribute to BBB hyperpermeability. They are inflammatory mediators derived from the arachidonic acid/lipoxygenase pathway. Injecting LTs

(e.g. LTB₄, LTC₄, and LTE₄) directly into the brain parenchyma increased BBB permeability. This effect could be limited by pretreatment with lipoxygenase inhibitors [137]. Although the specific mechanisms underlying arachidonic acid-induced BBB disruption have not been fully elucidated, some studies demonstrated that arachidonic acid could increase BBB permeability via ROS, which can be generated from COX and lipoxygenase signaling pathways [138, 139]. In addition, up-regulation of MMPs during neuroinflammation could be a link between PG production and BBB dysfunction because increased MMPs can destruct the BMs [136].

MMPs

MMPs belong to a family of at least 25 zinc-dependent endopeptidases and are extracellular enzymes that play a pivotal role in CNS physiology and physiopathology. Numerous studies have demonstrated that dysregulation of MMPs is a major cause of increased BBB leakage, cerebral edema, hemorrhage, leukocyte infiltration, progressive inflammatory reactions, and cerebral injury in many neurological conditions, such as Parkinson's disease [140, 141], Huntington's disease [142], migraine [143–145], and stroke [20]. MMP-2 and -9 are two prominent proteins that cause BBB disruption in many conditions. For example, A β 1-42 significantly up-regulated the level of MMP-2 and -9 and induced alterations in TJ scaffold and BBB leakage through a mechanism that involved receptor for advanced glycation end-products [146]. MMP-2 and -9 were reported to be responsible for BBB damage in HIV encephalopathy [147]. Early appearance of MMP-9 triggered BBB dysfunction after focal cerebral ischemia in mice [148]. Increased MMP-9 expression was observed in pericontusional brain; and MMP-9 inhibitors reduced brain swelling and final lesion volume in focal injury models [149]. MMP-2 seemed not to play a role in the vasogenic edema [149]. In addition, it was reported that MMP-1 was highly expressed in brain metastatic cells and was capable of degrading claudin and occludin but not ZO-1 [150]. The expression of MMP-1 seemed to be induced by COX2-mediated PGs. MMP-3 was also able to induce BBB abnormality in Parkinson's disease [141]. Mechanistically, MMP-induced endothelial barrier disruption was accompanied by MMP-mediated proteolytic degradation of claudin-5 in Japanese encephalitis-associated BBB breakdown [7]. Suppressing MMP-9-mediated degradation of the claudin-5 and occludin reduced BBB permeability in ischemic stroke [151].

Recent research has focused on pericytes as an important source of MMPs and inflammatory cytokines. Takata et al. reported that MMP-9 derived from pericytes induced migration of the cells from the endothelium, leading to loss

of pericytes and ensuing BBB damage [152]. Machida et al. recently demonstrated that pericytes played a pivotal role, as a highly thrombin-sensitive and MMP-9-producing cell, at the BBB in brain damage [153]. Halliday's research confirmed that pericytes maintained the integrity of the BBB and degenerated in AD. It was also revealed that apolipoprotein E4 (APOE4), a major genetic risk factor for late-onset AD, led to accelerated pericyte loss and enhanced activation of the low-density lipoprotein receptor-related protein-1-dependent cyclophilin A-MMP-9 pathway that caused BBB opening via degradation of endothelial TJ proteins and BMs in AD models [154, 155].

Another focus of research in the field is to identify potential pharmacological approaches to inhibit MMPs, at least in animal models. Takeuchi's data demonstrated that hydrogen was able to attenuate BBB disruption via suppressing MMP-9 activity in the hippocampus and improved neurological function outcome in a hypertensive stroke model [156]. Hydrogen sulfide inhalation was found to be able to decrease early BBB permeability and brain edema induced by cardiac arrest and resuscitation, possibly by decreasing MMP-9 and VEGF activity [157]. Allahtavakoli et al. revealed that ascorbic acid might be a useful candidate to reduce the side effects of delayed application of recombinant tissue plasminogen activator in stroke therapy by suppressing overexpression of MMP-9 [158]. Yang et al. reported that minocycline reduced reperfusion injury by inhibiting MMPs and microglial activity and enhancing TJ protein expression in ischemic brains [130]. Neurotensin receptor agonist HPI201 and TGF- β 1 significantly decreased MMP-2 and -9 levels in stroke and traumatic brain injury (TBI) models of adult rodents [100, 129]. Shi et al. argued that ethyl pyruvate conferred long-term neuroprotection against TBI, possibly through breaking the vicious cycle among MMP-9-mediated BBB disruption, neuroinflammation, and long-lasting brain damage [153]. Wang et al. observed that miRNA-130a inhibitors reduced brain edema, BBB permeability, and increased neurological deficit scores, possibly due to decreased MMP-2/9 expression in an acute intracerebral hemorrhage (ICH) model [159]. Wu et al. found that angiotensin 1–7 could effectively restore claudin-5 and ZO-1 expression in rats by down-regulating hypoxia-induced MMP-9 [160]. In addition, BBB disruption and brain edema formation play important roles in the secondary neuronal death and neurological dysfunction induced by ICH. Poloxamer 188, a multiblock copolymer surfactant, was found to protect against ICH, and the protective effect was associated with preventing BBB disruption through NF- κ B-MMP-mediated TJ protein degradation [161]. Furthermore, tissue destruction by MMPs is regulated by their endogenous tissue inhibitors (TIMPs). TIMPs prevent excessive MMP-related degradation of ECM components. Promoting

secretion of TIMPs has been explored as an effect way to inhibit MMP activities. Reuter et al. showed that simvastatin decreased MMP expression in human BMECs and experimental stroke mainly by means of increasing expression and secretion of TIMP-1 and TIMP-2 [162]. Resveratrol was also reported to attenuate the cerebral ischemia by maintaining the integrity of BBB via regulation of MMP-9 and TIMP-1 [163] (Fig. 1).

Free radicals

Human body normally produces free radicals at low concentrations as a defense mechanism. Overproduction of free radicals is implicated in the pathogenesis of many neurological diseases [140, 164, 165]. Reactive oxygen and nitrogen species are important in the early and delayed BBB disruption as the neuroinflammatory responses progress (see Rosenberg's reviews for details [166, 167]). For example, studies have shown that LPS-induced BBB disruption is mediated by ROS and reactive nitrogen species (RNS) production [168, 169]. LPS-induced microglial activation can increase NADPH oxidase, one of the major enzymes that generate ROS. Many reports have clearly demonstrated that ROS play a central role in BBB dysfunction during ischemia–reperfusion [166, 167]. For instance, Kamada et al. reported that hyperglycemia increased ROS levels and MMP-9 activity, exacerbating BBB dysfunction after ischemia and reperfusion [23]. Other studies showed that ROS induced BM degradation, enhanced tyrosine phosphorylation of TJ proteins by activating MMP-1/2/9 and decreasing TIMP-1 and -2, and led to increased permeability and monocyte infiltration [24]. In addition, HIV-1 envelope protein gp120 up-regulated MMP-2/9 expression, probably by gp120-mediated ROS [147].

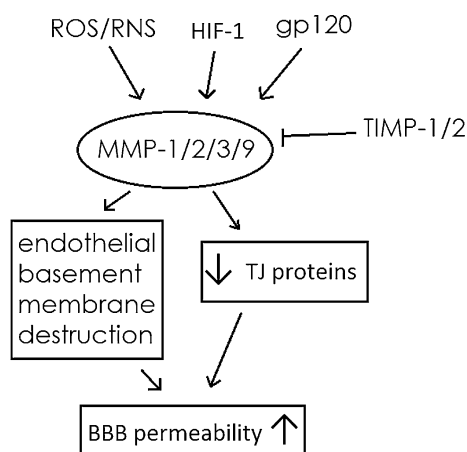


Fig. 1 A brief summary of MMPs as a factor controlling BBB permeability

Although the exact mechanism by which ROS induce BBB breakdown has not fully clarified, their harmful effects on BBB integrity could be mediated by many mechanisms such as MMP activation and TJ protein down-regulation [170–172]. Moreover, ROS can change the vascular tone and therefore influence cerebral blood flow. Their vascular effects also include increasing platelet aggregability and endothelial cell permeability, altering reactivity to vasodilators, and leading to the formation of focal lesions in endothelial cell membranes [14].

The brain has an essential defense system in response to free radical generation, and this antioxidant system contains small molecules such as glutathione and enzymes such as superoxide dismutase and GSH peroxidase that decrease levels of free radicals. It is believed that free radical scavengers could reduce oxidative stress and subsequently rescue BBB integrity [173]. Clinically, glucocorticoids have been used to treat neuroinflammatory conditions such as MS. Glucocorticoids seem to restore BBB integrity via a broad spectrum by down-regulating multiple inflammatory genes, such as cytokines [174], enzymes (e.g. inducible NO synthase and MMPs) [175], adhesion molecules [176]. Antioxidant gene delivery of Cu/Zn superoxide dismutase or GSH peroxidase blunted gp120-induced MMP production though down-regulating TIMP-1 and -2 [147]. Moreover, administration of a peroxynitrite decomposition catalyst, 5,10,15,20-tetrakis (4-sulfonatophenyl) porphyrinato iron (III), protected against hemoglobin-induced neurovascular injuries, which possibly in part by suppressing MMP-9 activation [177].

VEGF

VEGF is a specific mitogen; it binds to the vascular BM and causes proliferation and migration of endothelial cells [178]. Besides being the most prominent member of the angiogenic growth factor family, VEGF has been known since the 1980s to increase vascular permeability. VEGF has been characterized as an inducer of vascular leakage in response to hypoxia [17]. It has been reported that VEGF alters the expression and distribution of TJ proteins, leading to BBB hyperpermeability in hypoxia and autoimmune encephalomyelitis [179, 180]. VEGF exerted a major role in BBB disruption leading to subsequent edema during ischemic brain injury [181, 182]. It is noteworthy that VEGF seemed to have a dual role in both enhancing cerebral microvascular perfusion and increasing the BBB leakage in the ischemic brain. Zhang et al. reported that rhVEGF165 significantly enhanced cerebral microvascular plasma perfusion and improved functional neurological recovery when it was administered to ischemic rats at 48 h after a stroke while administration at 1 h after stroke exacerbated BBB leakage [182]. Pericytes and astrocytes

seem to be the major sources of VEGF in pathological conditions such as stroke [11, 183].

The mechanism that contributes to VEGF-mediated BBB hyperpermeability is not fully understood. It was reported that VEGF bound to the receptor tyrosine kinase, VEGF receptor 2, to induce BBB permeability [184]. In vitro studies suggest that the receptors for VEGF are located on the abluminal side of microvessels [185, 186]. Although it is known that VEGF increases BBB permeability by down-regulating the expression of TJ proteins [180], the specific target proteins of VEGF may vary, depending on the pathological condition and other factors. For example, our study has shown that claudin-5 seemed not to play a role in hyperglycemia-induced BBB hyperpermeability [26]. In contrast, Argaw's data suggest that down-regulation of claudin-5 by VEGF was central to disruption of the BBB in autoimmune encephalomyelitis [180]. Most recently, Japanese encephalitis virus-infected astrocytes were found to release VEGF that activated ubiquitin proteasome, degraded ZO-1, and disrupted endothelial barrier integrity in cultured BMECs [7]. In addition, several groups have reported that effect of exogenous VEGF on endothelial cells is partially attributed to NO generation from these cells through a NO synthase/cGMP-dependent pathway [182, 187]. VEGF may also induce microvascular hyperpermeability through activation of vesicular–vacuolar organelles [188] (Fig. 2).

Hypoxia inducible factor 1

The transcriptional factor hypoxia inducible factor 1 (HIF-1) is implicated in many cerebral vascular pathological disorders. HIF-1 is a heterodimeric complex consisting of a hypoxia inducible subunit HIF-1 α and a constitutively expressed subunit HIF-1 β . Under hypoxic conditions, HIF-1 α is stabilized and translocated to the nucleus where it dimerizes with HIF-1 β . The activated HIF-1 complex subsequently binds to hypoxic response elements in the regulatory regions of targeted genes. It induces transcription of more than a hundred genes with various functions [62]. VEGF is one of the most well-known HIF-1 target genes in vascular biology. Thus, one mechanism of HIF-1-mediated BBB disruption is VEGF up-regulation. Moreover, HIF-1 may cause BBB damage via the induction of MMP-2 and -9 [180]. It was reported that increased MMP activity during ischemia resulted in fragmentation/relocalization of occludin and claudin-5, leading to BBB opening [189]. Inhibition of HIF-1 ameliorated hypoxia-induced BBB disruption and the subsequent brain injury in animal ischemic models [190, 191]. Suppression of HIF-1 α and its downstream VEGF and MMPs reduced hemorrhagic transformation induced by hyperglycemia in ischemic rat brains, which indicates a recovery of BBB function [192].

Yeh et al. showed that inhibition of HIF-1 by YC-1 decreased VEGF expression and suppressed the increase in BBB permeability [191]. We were the first to report that high glucose induced a time-dependent up-regulation of HIF-1 α in mouse and human primary BMECs, in which it induced disarrangement of TJ proteins [26]. In addition, HIF-1 was responsible for MMP-9-mediated BBB disruption that contributed white matter injury in hypertensive rats with carotid artery occlusion and a Japanese permissive diet [193].

MicroRNAs

MicroRNAs (miRNAs) are a family of non-protein-coding small RNA molecules that negatively regulate protein expression. Many miRNAs have recently been found to regulate cellular signaling pathways and cell survival. Various miRNAs have been discovered in different cellular components of the BBB (Table 2). Several groups have reported that certain miRNAs control endothelial cell functions and BBB permeability. For example, Reijkerker et al. demonstrated that a large number of miRNAs were down-regulated in BMECs with impaired BBB function and that strengthening BBB function was generally associated with increased miRNA expression [194]. Specifically, they reported that brain endothelial cells overexpressing miRNA-125a-5p formed thicker and more continuous junctional complexes of VE-cadherin and ZO-1 and increased BMEC barrier function. Conversely, specific knockdown of the miRNA reduced the levels of VE-cadherin and ZO-1 along the BBB. In contrast, others have demonstrated negative effects of miRNAs on BMECs and BBB function. Lopez-Ramirez et al. observed that miRNA-155 negatively affected BBB function during neuroinflammation [28]. MiRNA-155 was up-regulated by inflammatory cytokines and expressed at the neurovascular unit of MS patients and of mice with experimental autoimmune encephalomyelitis. The miRNA up-regulation mimicked cytokine-induced alterations in junctional organization and BBB permeability. Furthermore, they demonstrated that miRNA-155 modulated brain endothelial barrier function by targeting cell–cell complex molecules, such as AA2, claudin-1, and molecules that are critical in cell-to-ECM interactions including dedicator of cytokinesis 1 and syntenin-1. Kalani et al. reported that miRNA-29b caused BBB hyperpermeability by regulating DNA (cytosine-5-)-methyltransferase 3 beta and MMP-9 that can disrupt the membrane and junction proteins leading to leaky vasculature [195]. Furthermore, Deng et al. showed that miRNA-21, but not -224, involved in ERK-mediated upregulation of MMP-9 in rat hippocampus and BBB dysfunction following cerebral ischemia [196]. Yin's results suggested that miRNA-15a was involved in

Fig. 2 A brief summary of VEGF as a factor controlling BBB permeability

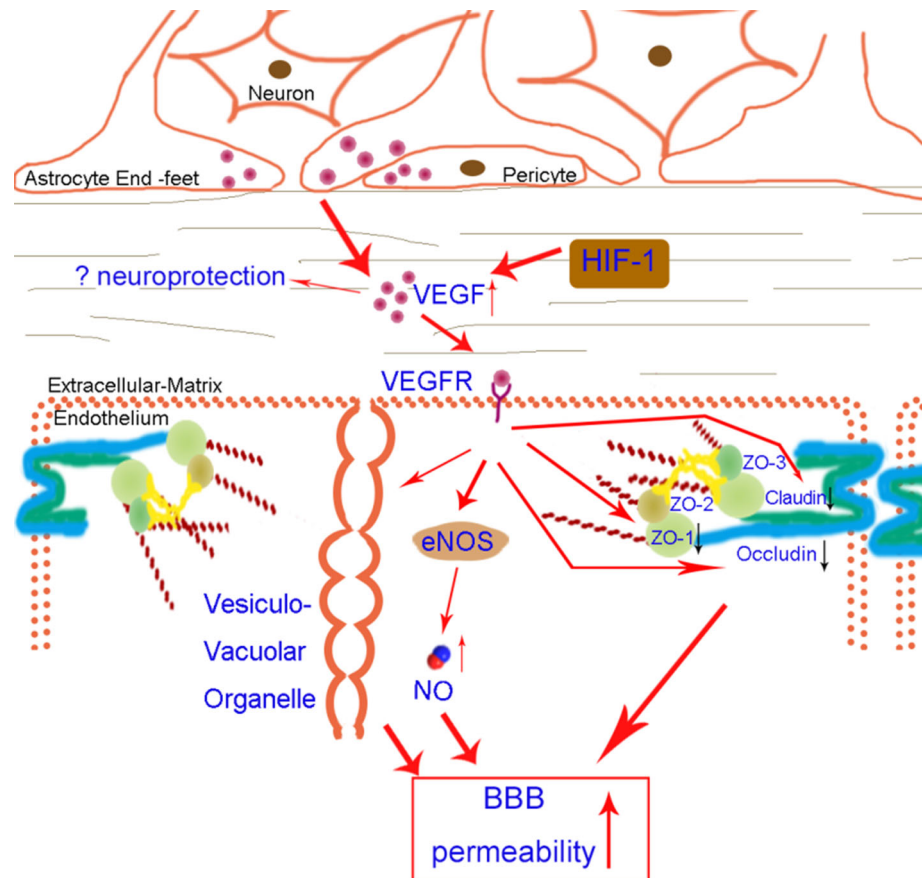


Table 2 MiRNA expression in different cells at the BBB

miRNAs	Cellular components of the BBB
miRNA-15a	Astrocytes, microglia [230]
miRNA-21	BMECs, astrocytes, microglia [231]
miRNA-26b	BMECs [198]
miRNA-28	BMECs [198]
miRNA-29	Astrocytes [232]
miRNA-98	BMECs [199]
miRNA-125a-5p	BMECs [194]
miRNA-155	BMECs [28], astrocytes [233]
miRNA-181c	Microglia [200]

ischemia-induced cerebral vascular endothelial injury [197]. Another group showed that miRNA-26b and -28 seemed to contribute to BBB dysfunction in lupus mouse brain [198]. Slava et al. demonstrated that miRNA-98 overexpression rescued the BBB during neuroinflammation through reduction of leukocyte adhesion and cytokine production. Their study also showed that inhibition of glycogen synthase kinase 3 β increased miRNA-98 expression in inflamed brain [199]. Moreover, miRNA-181c has been reported to be induced in many malignancies

such as breast cancer [50]. Despite the exact mechanism by which miRNA-181c mediates BBB breakdown is not yet fully understood, Tominaga et al. found an increased level of miRNA-181c in the blood circulation of breast cancer patients with brain metastasis, indicating that the miRNA plays a crucial role in brain metastasis [52]. In addition, a recent study showed that miRNA-181c triggered the Toll-like receptor 4 pathway, resulting in microglial activation and neuroinflammation [200]. These observations suggest the miRNAs are a new set of controllers of BBB permeability under stress and pathological conditions.

Anesthetic agents

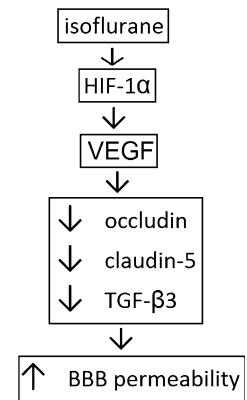
Besides endogenous mediators that control/regulate BBB function and permeability, many pharmacological agents have great impact on BBB permeability through regulating the signaling mediators and the structural components. One widely studied pharmacological agent is isoflurane, a popular volatile anesthetic routinely used in human patients. As early as in 1992, Chi et al. reported that isoflurane decreased the transport of small hydrophilic molecules across the BBB by measuring blood–brain transfer coefficient and capillary permeability-surface area

[201]. Later on, Tetrault et al. demonstrated that high doses of isoflurane (3 %) opened the BBB in the cortex and thalamus while milder doses (1 %) only opened the BBB in the thalamus in cats by monitoring the extravasation of Evans blue [202]. Their results also showed that the opening of the BBB was associated with an evaluated increase in cerebral volume of 2–2.8 %, indicating brain edema [202].

The mechanism of isoflurane-mediated hyperpermeability of BBB seems to involve in its direct or indirect effects on the TJ proteins. Most recently, Cao et al. studied the effects of isoflurane on BBB in aged rats. Their results showed that 1.5 % isoflurane resulted in reversible time-dependent morphological damage of the BBB ultrastructure and significant decreases in expression of the TJ protein occludin in 24-month old rats. They concluded that occludin down-regulation was one of the mediators of isoflurane-induced BBB disruption, and might contribute to hippocampus-dependent cognitive impairment after isoflurane exposure in aged rats [203]. In a model of TBI, ZO-1 expression was more significantly disrupted in sevoflurane than isoflurane-exposed animals with controlled cortical impact, while claudin-5 was less affected in the pericontusional area [155]. Moreover, Zhao et al. reported in 2014 that isoflurane treatment induced a time- and concentration-dependent decrease in occludin mRNA and protein levels in human BMECs, which was partially abrogated by silencing the alpha subunit of HIF-1 [204]. Isoflurane-mediated HIF-1 α expression up-regulated the level of VEGF, which decreased the expression of occludin and TGF- β 3 that accelerated the endocytosis of occludin [204]. If HIF-1 activation is indeed responsible for isoflurane-induced BBB abnormality, it is still not clear how isoflurane up-regulates the transcriptional factor (Fig. 3).

Furthermore, several recent reports suggest that under pathological conditions, isoflurane may exaggerate brain injury by further opening the BBB. Dittmar et al. investigated the effect of a 2-h isoflurane exposure on apoptosis of the cerebral endothelium following 24 h of hypoxia in an *in vitro* BBB model. They found that isoflurane treatments resulted in severe cellular morphological changes and a significant dose-dependent increase in DNA fragmentation in trans-differentiated human umbilical vein endothelial cells, possibly due to increases in pro-apoptotic Bax levels and decreases in the level of anti-apoptotic Bcl-2 [205]. In a rat stroke model of embolic occlusion of middle cerebral artery, Bezerra et al. observed that BBB permeability measured by blood-to-brain forward rate constant was elevated in the ipsilateral hemisphere in animals under either halothane or isoflurane anesthesia. However, isoflurane-anesthetized rats demonstrated less BBB damage than halothane-anesthetized group [155]. In addition, BBB disruption causes cerebral edema formation and is a major

Fig. 3 Proposed mechanism responsible for isoflurane-mediated BBB hyperpermeability



cause for high mortality after TBI. Results from Thal's study demonstrated that 24 h after controlled cortical impact, brain water content of mice subjected to either sevoflurane or isoflurane increased significantly while in healthy mice, anesthesia did not influence brain water content [206].

In summary, isoflurane is able to affect BBB permeability at normal conditions, to induce brain edema, and to exaggerate BBB injuries under pathological conditions such as stroke and TBI. This effect may be attributed to its impact on TJ protein expression, possibly through regulating HIF-1 activity. In addition, other anesthetic agents have also been shown to affect BBB permeability. Similar to isoflurane, sevoflurane was able to induce structural changes in BMECs and increase BBB permeability [155, 206, 207]. An *in vitro* study showed that thiopental induced BBB opening at high concentrations (~100 μ g/ml) while fentanyl and methohexital did not affect BBB permeability [208]. Future research is required to investigate the exact molecular mechanisms of anesthetic-mediated BBB opening and to mitigate the adverse effect on patients.

Factors that regulate transcellular transport of iron through BBB

The above discussion focuses on paracellular permeability of the BBB. Transcellular transport across BBB might be affected by the same or different signaling mediators as those discussed above. In this section, recent progress on transcellular transport of iron ion is briefly reviewed. Brain is one of the most abundant iron-containing organs. When there is iron deficiency or overload in the brain, the nervous system suffers from disturbed functions. Hence, it is essential to maintain brain iron homeostasis. The key to iron balance in the brain lays in the regulation of brain iron uptake. The regulation of iron transport across the BBB is of major importance to brain iron uptake [209]. Available data suggest that uptake of plasma transferrin binding iron (Tf-Fe) occurs at the luminal membrane of BMECs, mainly

through the classic transferrin receptors (TfR)-mediated endocytosis [210]. In BMECs, due to the acidification of the microenvironment within endosomes, iron ions dissociate from Tf and are reduced from Fe^{3+} to Fe^{2+} . Ferrous ion can translocate across the endosomal membrane via a proton driven transporter, divalent metal transport 1 [211]. Most of the Tf and TfR will return to the luminal membrane. Iron ion is proposed to be transported across the abluminal membrane of the BBB into the interstitial fluid of the brain [212]. This process likely involves iron exporter ferroportin1 (FPN1) on the abluminal membrane, but the exact mechanism remains to be explored. Available studies showed that both hephaestin and FPN1 have been identified in BMECs. In addition, iron efflux from BMECs by FPN1 may require the action of an exocytosomal ferroxidase, possibly ceruloplasmin produced by glial cells [213].

At the systematic level, brain iron homeostasis may be regulated by a peptide ‘hormone’ named hepcidin. Hepcidin binds to the extracellular loop of FPN1 and causes its internalization and degradation, and thereby reduces cellular iron efflux. Recent data indicate that brain hepcidin stimulates internalization of FPN1 to regulate iron efflux from the BMECs [214]. The observation indicates that hepcidin may play an important regulatory role in brain iron metabolism. The expression of hepcidin can be regulated by high iron levels, inflammation, erythropoiesis, or hypoxia [215]. When iron levels are high, hemojuvelin and TfR2 increase hepcidin expression. In addition, IL-6, a cytokine that regulates paracellular permeability, is able to stimulate hepcidin expression through molecular pathways that could include binding of STAT3 to the hepcidin promoter.

Perspectives

Blood–brain barrier permeability is tightly controlled by many component factors of BMECs, astrocytes, pericytes and microglial cells. The TJ structure is essential to the integrity of BBB. It establishes a linkage between endothelial cells, which is tight enough to form a barrier to separate brain cells from exposure to the systemic circulation. As discussed above, altered TJ protein expression and structural arrangement underlines the mechanism of BBB dysfunction in many brain disorders. Hence, normalizing TJ protein expression and arrangement is an important pharmacological goal to enhance the BBB integrity under pathological conditions. The regulation of the expression of TJ proteins including their interactions, stereochemistry, and their signaling mediators are directions of future studies. Meanwhile, many other proteins play important roles in maintaining BBB permeability and functions. As evident from many recent reports, integrins

binding to ECM together with annexins and agrin play an essential role in BBB integrity and functions. Yet, the functions of integrins, annexin and agrin in BMEC are still not well revealed, especially their interactions with TJs. These are interesting areas to be explored in the future.

Inflammation plays a central role in the pathogenesis of neurodegenerative and neurological disorders. Evidence is being accumulated for the harmful effects of neuroinflammation on BBB integrity. Inflammatory mediators such as cytokines, eicosanoids, or free radicals have been reported to cause BBB damage. Results from recent studies are consistent in supporting that TJ protein alteration is the most predominant mechanism underlying neuroinflammation-induced BBB disruption. Although the mechanism has not fully been understood, activation of microglial cells may act as the first step in BBB disruption in CNS inflammation. Recent studies have established a link between microglial activation and BBB breakdown [216]. Interestingly, microglial cells seem to be a double-edge sword. On one hand, these cells regulate the integrity of the mature BBB. On the other hand, the microglial response may activate different signaling pathways through pro-inflammatory mediators. Inactivation of microglial cells is most likely not the suitable therapeutic approach because their activation can mediate many beneficial physiological functions such as phagocytosis of damaged neurons [217]. Thus, targeting the microglial signaling pathways that mediate inflammation-induced BBB breakdown would be a promising therapeutic strategy. Sonic hedgehog has been known to maintain BBB integrity via TJ protein up-regulation [218]. Overexpression of sonic hedgehog could be an effective experimental approach to restore BBB properties after inflammation. Furthermore, many studies have shown that anti-inflammatory and antioxidative agents could limit BBB disruption after neuroinflammation. Inhibition of the COX signaling pathway with indomethacin protected against TNF- α -induced BBB disruption in vivo. This COX inhibition was associated with a decrease in MMP-9 and -3 activities [136]. Recently, a report showed that ginsenosides, a group of anti-inflammatory and antioxidant bioactive compounds, reduced BBB disruption following cerebral ischemia [53] by suppressing the expression of pro-inflammatory mediators such as NO synthase, IL-1 β , and MMP-9 [53, 200]. Moreover, caffeic acid phenethyl ester, an anti-inflammatory and antioxidative agent [219], limited BBB opening in a TBI rodent model through preserving TJ proteins such as claudin-5 [220].

Emerging evidence suggests that the brain is a target of diabetes, resulting in diabetic encephalopathy [221]. The pathophysiology of diabetic encephalopathy is still largely elusive. Cerebral microvascular disturbances could play an important role in diabetes-induced brain abnormality. In fact, BBB dysfunction (i.e. hyperpermeability) has been

demonstrated in diabetic patients by magnetic resonance imaging [222] and in animal models of diabetes [223]. However, the mechanism responsible for BBB hyperpermeability in diabetes is not known. To understand the mechanism of diabetes-induced BBB dysfunction is crucial for the prevention and treatment of diabetes-induced neurological diseases. Future studies may focus on the following to reveal the mechanism, inflammatory mediators, MMPs, HIF-1, VEGF, ROS, and miRNAs that control the permeability of BBB. In fact, excessive generation of ROS has been recognized as an integral part to the pathophysiology of diabetes, particularly in diabetes-induced vascular diseases [224]. Expression and activation of HIF-1 in hyperglycemia-exposed brain endothelial cells are believed to be closely associated with the brain microvascular damage and to be induced by free radicals. We recently demonstrated that high glucose activated HIF-1 in mouse and human BMECs [26]. It will be useful to determine what specific ROS and inflammatory mediators are generated and responsible for the enhanced activities of HIF-1 and BBB dysfunction in diabetes. In addition, anesthetic agents such as isoflurane may change BBB permeability in diabetic patients. More research is needed to find its molecular mechanism to minimize the adverse effects on patients. Moreover, change in iron transport across BBB in diabetic patients is also an important and interesting topic to be addressed.

In summary, BBB is a very important component in the body regulating the brain homeostasis. Permeability of BBB is regulated by many factors such as junctional proteins, BM proteins, cytokines, VEGF, HIF-1, ROS, miRNAs, etc. Recent research has made great progress in revealing the mechanisms by which they regulate BBB permeability. Changes of BBB permeability are involved in the pathogenesis of many brain disorders. Further studies are needed to reveal a better understanding of BBB permeability regulation, particularly in neuroinflammatory and diabetic conditions, and to develop novel and specific therapeutic approach to mitigate the harmful effect of BBB dysfunction.

Acknowledgments This work was supported in part by the University of Kansas Center for Research and National Natural Science Foundation of China (31520103908 and 31528013). Mohammed M. A. Almutairi was supported by a scholarship from King Saud University (Riyadh, Saudi Arabia).

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