

Structural Alterations to the Endothelial Tight Junction Complex During Stroke

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1 Introduction

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

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

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

2 Normal BBB Structure and Function.

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

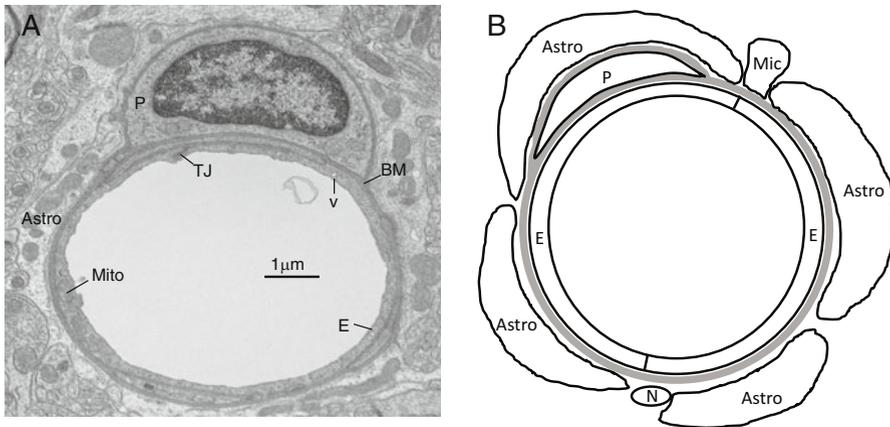


Fig. 1 (a) An electron micrograph showing the structure of a normal mouse cerebral capillary. (b) A schematic showing the relationship between the endothelium and other elements of the neurovascular unit. Brain capillary endothelial cells (E) are linked by tight junctions (TJ), show few vesicles (v)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

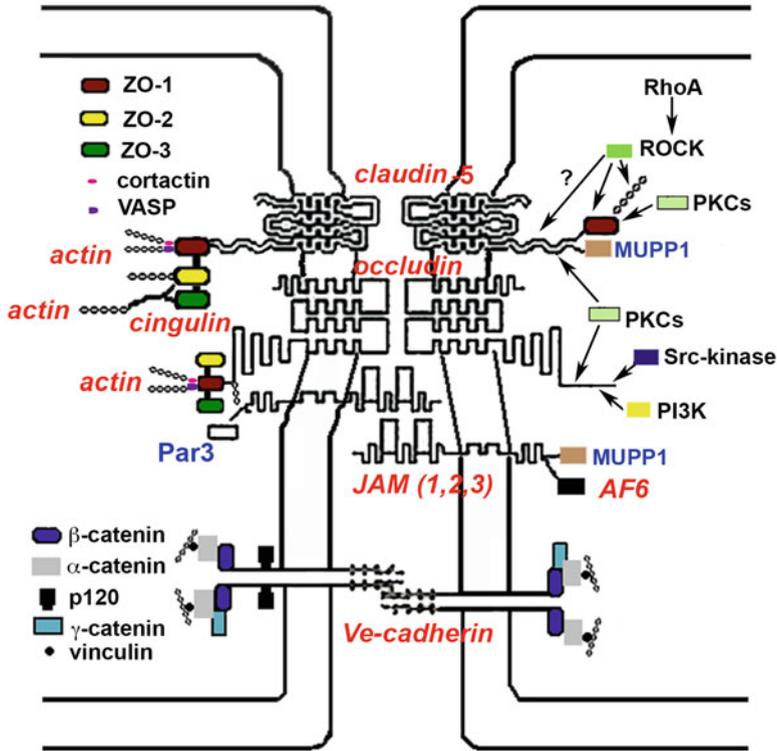


Fig. 2 Schematic representation of the basic components of the brain endothelial junctional complex. The *left* of the panel shows potential structural components of the TJ complex. Claudin-5, occludin, and the JAMs are transmembrane proteins that link adjacent cells. Claudin-5 is the major occlusive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

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

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

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

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

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

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

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

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

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

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

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

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

3 Alterations in BBB Function After Stroke

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

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

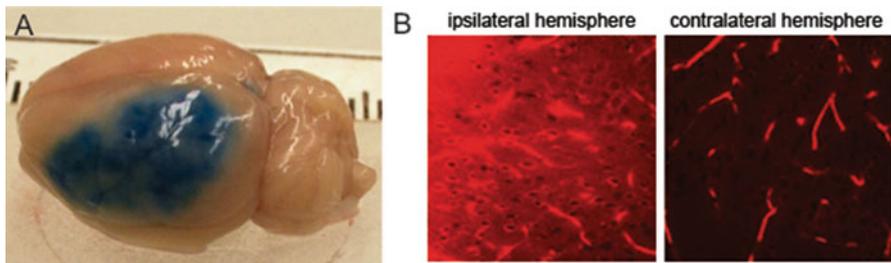


Fig. 3 (a) An example of BBB disruption after rat focal cerebral ischemia at the macroscopic level. A hyperglycemic rat underwent 2 h of middle cerebral artery (MCA) occlusion followed by 2 h of reperfusion. The rat was injected intravenously with Evans blue, which binds to albumin in the bloodstream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

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

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

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

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

4 Alterations to TJ Structure After Stroke

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

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

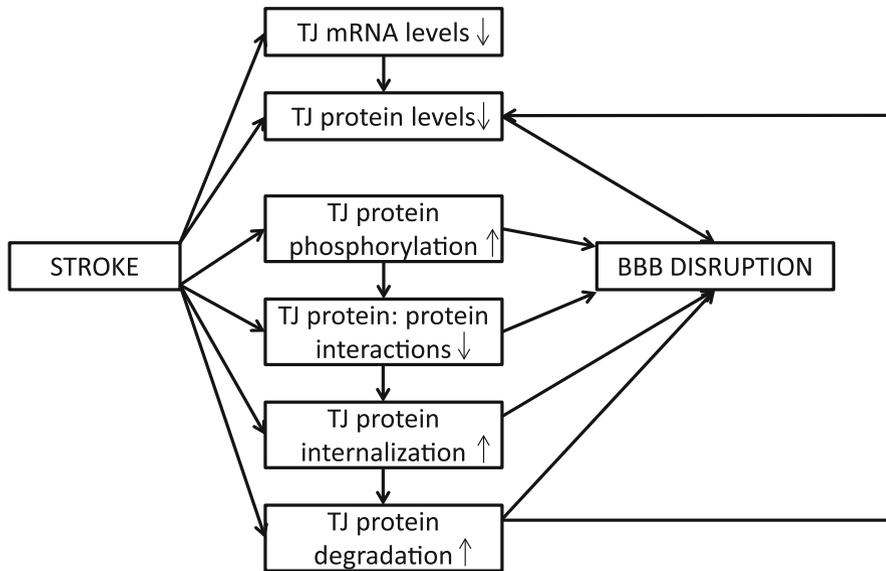


Fig. 4 Schematic showing changes occurring in TJ proteins after stroke that participate in BBB hyperpermeability. Although a potential link between stroke-induced TJ protein phosphorylation, loss of TJ protein interactions, TJ protein internalization and degradation is shown, stroke may affect those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

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

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

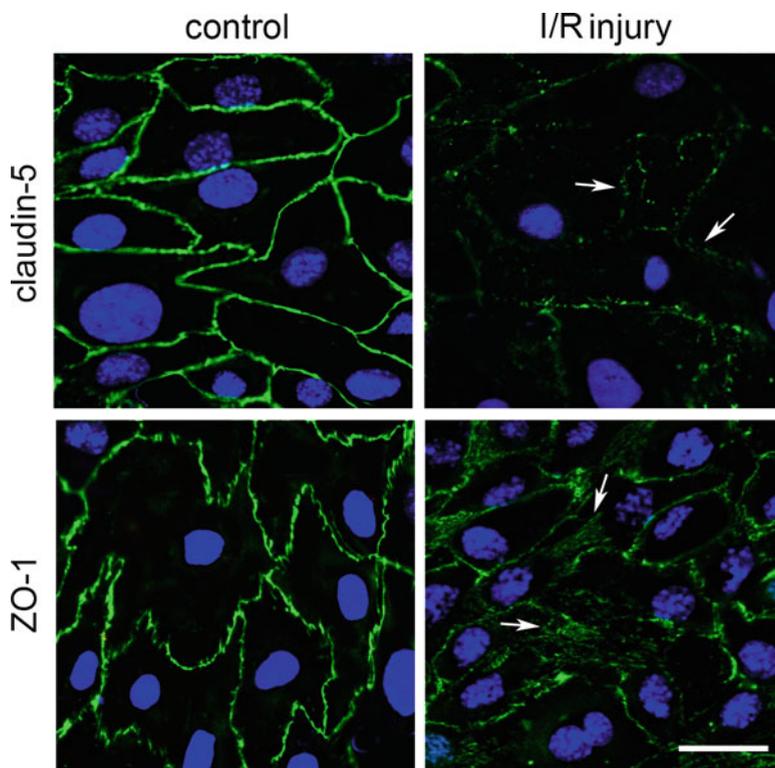


Fig. 5 Mouse brain endothelial cells were exposed to oxygen–glucose deprivation (OGD/ischemia) for 5 h followed by normal oxygen and glucose condition (reperfusion) for 1 h as a model of ischemia/reperfusion (I/R) injury. Control represents cells exposed to normal oxygen and glucose. Immunofluorescent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

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

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

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

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

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

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

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

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

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

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

5 Signaling Mechanisms Underlying Alterations in TJ Structure in Stroke

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

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

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

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

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

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

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

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

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

6 Non-TJ Pathways Involved in BBB Hyperpermeability After Stroke

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

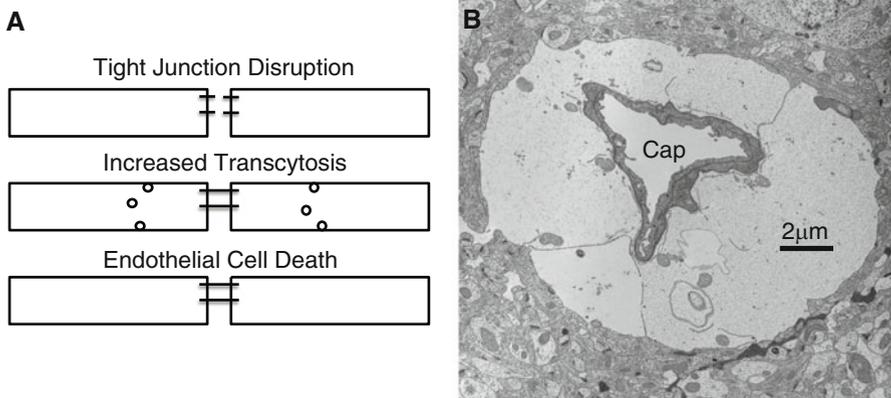


Fig. 6 (a) Routes proposed to participate in BBB disruption after stroke. Note that these routes are not mutually exclusive. (b) Electron micrograph of a cerebral capillary (cap) from the ischemic penumbra in a mouse that underwent 3 h of middle cerebral artery occlusion. Note the marked perivascular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

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

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

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

7 Therapeutic Interventions

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

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

8 Conclusions

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

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