Running Head: Endothelial KLF11 protects blood-brain barrier.

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# Abstract

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Microvascular endothelial cell (EC) injury and the subsequent blood-brain barrier (BBB) breakdown are frequently seen in many neurological disorders, including stroke. We have previously documented that peroxisome proliferator-activated receptor gamma (PPARy)mediated cerebral protection during ischemic insults needs Krüppel-like factor 11 (KLF11) as a critical coactivator. However, the role of endothelial KLF11 in cerebrovascular function and stroke outcome is unclear. This study is aimed at investigating the regulatory role of endothelial KLF11 in BBB preservation and neurovascular protection after ischemic stroke. EC-targeted overexpression of KLF11 significantly mitigated BBB leakage in ischemic brains, evidenced by significantly reduced extravasation of BBB tracers and infiltration of peripheral immune cells, and less brain water content. Endothelial cell-selective KLF11 transgenic (EC-KLF11 Tg) mice also exhibited smaller brain infarct and improved neurological function in response to ischemic insults. Furthermore, EC-targeted transgenic overexpression of KLF11 preserved cerebral tight junction (TJ) levels and attenuated the expression of pro-inflammatory factors in mice after ischemic stroke. Mechanistically, we demonstrated that KLF11 directly binds to the promoter of major endothelial TJ proteins including occludin and ZO-1 to promote their activities. Our data indicate that KLF11 functions at the EC level to preserve BBB structural and functional integrity, and therefore confers brain protection in ischemic stroke. KLF11 may be a novel therapeutic target for the treatment of ischemic stroke and other neurological conditions involving BBB breakdown and neuroinflammation.

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# Introduction

Stroke ranks as the fifth leading cause of death, and accounts for 1 out of every 20 deaths in the United States (7, 58). Ischemic stroke leads to cerebrovascular and neuronal damage, both of which contribute to ischemic injury and dictate stroke outcome (24, 66). Although studies have implied the effectiveness of neuroprotectants in animal stroke models (24, 32), there has been very little progress made in the past three decades concerning the successful translation of neuroprotective strategies to the clinical setting. The lack of clinical translation implies that focusing on neuroprotection only is insufficient. Non-neuronal cells and the local microenvironment of the surviving neurons could also serve as therapeutic targets to alleviate ischemic brain injury (61, 65). Cerebral vascular endothelial cells (ECs) are major components

70 of brain microvasculature and play critical roles in maintaining the integrity of the blood-brain 71 barrier (BBB, a brain-specific microvascular structure) and cerebral homeostasis under 72 physiological conditions (1, 5, 50). Cerebral ECs are tightly connected by adherens junctions 73 (AJs) and tight junctions (TJs), including occludin, claudins, and junctional adhesion molecule 74 (JAM), which are anchored to the actin cytoskeleton by scaffold proteins such as zonula 75 occludens (ZO)-1, AF6, and cingulin. Together they form a diffusion barrier that selectively 76 blocks the passage of most compounds through blood to brain compartments (4, 16, 39). When 77 BBB integrity is compromised, proinflammatory factors may pass through the BBB to attract 78 circulating immune cells to the injured brain, leading to secondary ischemic brain parenchymal 79 injuries (15, 33-35, 46, 55, 60). Therefore, inhibition of BBB disruption through stabilizing TJs 80 and protecting brain endothelium has become promising therapeutic strategies for ischemic stroke (18, 22, 45). 81 82 Krüppel-like factors (KLFs) are a large family of zinc finger transcription factors that are known 83 to trans-activate or trans-repress gene expression in various organisms (63). Recent studies have 84 reported the involvement of KLF members in various developmental and pathological vascular processes (2, 52). A recent RNA sequencing study showed that in human endothelium, KLF11 85 86 has modest mRNA expression (20). KLF11 has been reported to play functional roles in the 87 regulation of cell growth and differentiation, cholesterol metabolism, and cell death (21, 25, 42, 88 52). Moreover, population genetics studies have demonstrated that mutations in the KLF11 gene 89 are strongly associated with type 2 diabetes (21, 42). Previous studies indicated that KLF11 90 inhibits endothelial activation in response to inflammatory stimuli (19). In addition, vascular 91 smooth muscle cell-selective deletion of KLF11 aggravates arterial thrombosis in mice (36). 92 However, the role of KLF11 in cerebrovascular biology remains largely undetermined. 93 A previous study by our group reported that KLF11 functions as a peroxisome proliferator-94 activated receptor-y coregulator to attenuate middle cerebral artery occlusion (MCAO)-induced 95 ischemic brain injury (62). In conventional KLF11 knockout mice, we observed larger brain 96 infarct, increased sensorimotor loss, aggravated BBB leakage, higher brain edema, and lower 97 cerebral blood flow (CBF) following cerebral ischemic insult (53). These studies suggest KLF11 98 plays a protective role in the pathogenesis of ischemic stroke. However, the molecular events and 99 regulatory roles of endothelial KLF11 itself in stroke-induced cerebral EC injury and BBB 100 disruption remain virtually unclear. Thus, the present study sought to test the hypothesis that

101 KLF11 stabilizes the cerebrovascular endothelial structure while simultaneously reducing stroke-102 induced inflammatory events through preserving the integrity of TJs in cerebral ECs.

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# Results

105 Generation and characterization of EC-selective KLF11 transgenic mice. ECs are major 106 components of the BBB and are very sensitive to oxidative stress and ischemic insults (23). 107 Previously, KLF11 expression was found to be significantly decreased in cultured mouse 108 primary brain microvascular endothelial cells (BMECs) after 4 h or 16 h oxygen-glucose 109 deprivation (OGD) treatment (Fig. S1A) as well as in isolated cerebral microvessels from 110 C57BL/6J mice that were subjected to 1 h MCAO followed by 24 h reperfusion (Fig. S1B). We 111 also reported that genetic deletion of KLF11 in mice significantly augmented MCAO-induced 112 BBB disruption through exaggerating cerebrovascular permeability and edema (53). However, 113 whether KLF11 maintains the integrity of the BBB at the very early stages (1-3 h) after ischemic 114 insults, and whether endothelial KLF11 plays distinct roles in the pathogenesis of ischemic brain 115 injury remain unknown. To explore the specific role of KLF11 in endothelial cells after ischemic 116 injury, we generated transgenic mice with vascular endothelial cell-selective overexpression of 117 KLF11 on a C57BL/6J background. The EC-specific KLF11 transgenic construct contains the 118 Tie 2 promoter and Tie 2 enhancer (47, 64) to drive expression of the full-length KLF11 cDNA 119 sequence (Fig. 1A). The Tie 2 promoter allows selective targeting of KLF11 to vascular ECs. 120 The transgenic founder mice with KLF11 overexpression (EC-KLF11 Tg) were identified by 121 genomic PCR genotyping (Fig. 1B). To confirm the overexpression of KLF11 in the brain 122 microvasculature, we isolated cerebral microvessels from EC-KLF11 Tg and WT mouse brains. 123 We found a significant upregulation of cerebrovascular KLF11 protein level in EC-KLF11 Tg 124 mice than that of WT controls (Fig. 1C left). Moreover, KLF11 was not overexpressed in the 125 cortical protein extracts from EC-KLF11 Tg mouse brain in comparison with WT controls (Fig. 126 1C right). Of note, in sham-operated mice, no significant differences were observed in CD31-127 labeled microvasculature of EC-KLF11 Tg mice compared to WT controls, as indicated by 128 branch points, capillary numbers, and branch length (Fig. 1 D-E). We also noticed that regional 129 CBF did not differ by transgenic overexpression of KLF11 in endothelium 15 minutes before 130 ischemia, 15 minutes after ischemia, or 15 minutes after reperfusion (Fig. 3 D-E), suggesting that 131 transgenic manipulation of KLF11 in endothelium does not cause possible cerebrovascular

structural (collateral) changes in the brains of EC-KLF11 Tg mice. Thus, EC-KLF11 Tg mice can be used as a powerful tool to investigate the functional role of KLF11 in vascular endothelial biology and BBB pathologies *in vivo*.

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EC-selective KLF11 transgenic overexpression ameliorates post-ischemic cerebrovascular permeability. Next, we performed Evans Blue and TMR-Dextran extravasation assays to observe and quantify changes in BBB permeability. EC-KLF11 Tg and WT mice were subjected to 1 h MCAO followed by 1 h, 3 h, or 24 h reperfusion. The results showed that EC-targeted transgenic overexpression of KFL11 markedly reduced, but not completely blocked the leakage of both Evans blue dye and TMR-dextran into ischemic brain after 24 h reperfusion (Fig. 2 A-D). To assess BBB permeability at earlier stages (1-3 h) after ischemic insult, we also analyzed the extravasation of a small fluorescent tracer, Alexa 555 cadaverine (0.95 kDa), which was injected through the tail vein, and the relatively larger endogenous plasma IgG (150 kDa) into the ischemic brain parenchyma (48, 49). The extravasation of the small molecule cadaverine in both the ipsilateral cortex and striatum was detected as early as 1 h reperfusion, whereas endogenous IgG was detected in the same area at 3 h reperfusion (Fig. 2E). Compared with WT controls, EC-KLF11 Tg mice consistently showed significantly reduced extravasation of cadaverine (at 1h, 3h, and 24h reperfusion; Fig. 2F) and IgG (at 3h and 24h reperfusion; Fig. 2G). These results suggest that early-onset BBB impairment occurred following ischemic insult and endothelial-specific KLF11 transgenic overexpression protects against MCAO-induced early-onset and progressive (late-onset) BBB impairments. Moreover, anti-MAP2 immunostaining indicated that the brain regions which present signs of BBB breakdown at earlier stages after MCAO progressed into infarct zones at 24 h after MCAO (Fig. 2E). Quantification analysis of anti-microtubuleassociated protein 2 (MAP2) immunostaining showed that EC-KLF11 Tg mice had significantly reduced brain infarction compared with WT controls (Fig. 2H). We further measured and quantified the water content in ipsilateral and contralateral hemispheres of EC-KLF11 Tg and WT mice 72 h after MCAO. The results demonstrated that brain water content in the ipsilateral hemisphere of EC-KLF11 Tg mice was significantly less than that in the WT group (Fig. 2I), whereas water content in the contralateral hemisphere from both groups showed no significant differences (Fig. 2I). It is known that blood neutrophils can migrate through an impaired BBB and carry proinflammatory mediators into the injured brain, thereby causing more serious BBB

damage and secondary expansion of ischemic brain parenchymal injury (49). We used immunohistochemistry methods to detect neutrophils with a labeled Ly-6B antibody (Fig. 2J). Quantification analysis of Ly-6B immunostaining confirmed that in comparison with WT controls, MCAO-induced increases in brain infiltration of Ly-6B<sup>+</sup> neutrophils were significantly reduced in EC-KLF11 Tg mouse brains (Fig. 2K). Taken together, these results suggest that endothelial-selective KLF11 transgenic overexpression not only protects against ischemia-induced BBB damage but also provides further neurovascular protection against ischemic stroke.

EC-targeted KLF11 overexpression reduces ischemic brain injury and improves shortterm neurological function. The protective role of EC-KLF11 overexpression against ischemic insult was further confirmed by 2% 2,3,5-triphenyltetrazolium chloride (TTC) staining in another cohort of mice, in which neurological deficits were also scored. Compared with WT controls, EC-KLF11 Tg mice presented much smaller cerebral infarcts (Fig. 3 A-B) and significantly reduced neurological scores (Fig. 3C), indicating better neurological outcome after stroke. Of note, regional CBF showed similar changes between EC-KLF11 Tg and WT mice 15 minutes before ischemia, 15 minutes after ischemia, or 15 minutes after reperfusion (Fig. 3 D-E). In humans, cerebral ischemia always leads to sensorimotor dysfunction. We then investigated whether EC-targeted KLF11 overexpression affects post-stroke functional recovery by a battery of three different behavioral tests: adhesive tape removal, foot fault, and rotarod tests before and up to 7 days following MCAO (Fig. 3 F-I). Compared with WT control mice, EC-KLF11 Tg mice showed better recovery of sensorimotor function: longer staying latency in the rotarod test (Fig. 3F), lower rate of fault steps in the foot fault test (Fig. 3G), and shorter touching or removing time in the adhesive tape removal test (Fig. 3 H-I). These cumulative findings suggest that endothelial-specific overexpression of KLF11 facilitates spontaneous sensorimotor recovery after ischemic brain insult.

**EC-selective KLF11 overexpression preserves the integrity of junctional proteins after ischemic brain injury.** TJ proteins are essential for maintaining the integrity of the BBB (28). Therefore, we further examined the underlying mechanism of endothelial KLF11-mediated BBB protection *in vivo*. EC-KLF11 Tg and WT mice were subjected to 1 h MCAO followed by one or three days of reperfusion. Mouse brain tissue was harvested at the indicated time points after

MCAO and subjected to total RNA isolation and qPCR analysis (Fig. 4 A-B) or western blotting analysis (Fig. 4 C-E). Endothelial-targeted KLF11 transgenic overexpression significantly preserved the expression of TJ proteins, ZO-1 and occludin in ischemic brain tissue at both the mRNA (Fig. 4 A-B) and protein (Fig.4 C-E) levels, consistent with the protective role of EC-KLF11 overexpression in MCAO-induced BBB damage. Reduction in ZO-1 and occludin expression in vascular endothelial cells (labeled by CD31) following ischemic brain insults were further confirmed by immunofluorescent staining (Fig. 4F). As expected, EC-KLF11 overexpression markedly retained their *in situ* expressions (Fig. 4F). These results indicate that endothelial-targeted KLF11 overexpression robustly protected the structural integrity of the BBB.

# **EC-targeted KLF11 transgenic overexpression mitigates inflammatory activity in ischemic brain regions.** Neuroinflammation following ischemic stroke contributes to BBB damage (57). To test whether EC-targeted KLF11 overexpression alleviates MCAO-induced inflammatory activities in the ischemic brain, we measured a panel of inflammatory mediators in the cortex of the ipsilateral brain hemisphere at one or three days after MCAO. The mRNA expression levels of six pro-inflammatory mediators: *TNF-α*, *IL-1β*, *MCP-1*, *IL-6*, *ICAM-1*, and *P-selectin* markedly elevated following MCAO, and EC-targeted KLF11 overexpression attenuated this elevation (Fig. 5 A-F). We then measured the expression of the abovementioned inflammatory factors using Quantikine ELISA kits. Consistent with our previous observations, the six inflammatory mediators all reduced in EC-KLF11 Tg mice when compared with WT controls following MCAO (Fig. 5 G-L). Taken together, these results demonstrate that KLF11 transgenic overexpression in endothelium mitigates post-stroke inflammatory responses.

KLF11 alleviates OGD-induced BBB leakage in an *in vitro* BBB model. To further investigate endothelial KLF11-mediated BBB protection against ischemic insults, we employed a two-cell based *in vitro* BBB model. The co-culture system consists of mouse astrocytes that are seeded on the bottom of a 12-well plate and adenovirus-transduced mouse BMECs that are seeded on the polyester membrane of transwell inserts (Fig. 6A). The co-culture system was then subjected to ischemic-like insult of OGD for 16 h, followed by 2 h reoxygenation. Transepithelial/transendothelial electrical resistance (TEER) was measured to monitor the functional integrity of the cellular barrier (51). When KLF11 was overexpressed in mBMECs by

adenovirus-mediated infection before being subjected to OGD, the barrier functional disruption induced by OGD was attenuated in the Ad. KLF11group when compared with Ad. LacZ control group (Fig. 6B). On the contrary, OGD-induced barrier functional disruption was worsened in the adenoviral KLF11 knockdown group (Ad. shKLF11) compared with Ad. shLacZ control group (Fig. 6C). Next, luminal-to-abluminal barrier permeability (paracellular permeability) of the small fluorescent tracer (3 kDa Dextran) was measured at indicated time points after OGD. When KLF11 was upregulated by adenovirus in mBMECs prior to OGD, leakage of the 3 kDa tracer was significantly reduced, but not completely blocked, at 2-8 h post-OGD reoxygenation (Fig. 6D). In contrast, when KLF11 was downregulated in mBMECs by adenovirus-mediated infection (Ad. shKLF11) prior to OGD, leakage of the 3 kDa tracer was significantly increased at 2-24 h post-OGD reoxygenation compared to the Ad. shLacZ group (Fig. 6E). Based on the above observations, we conclude that KLF11 overexpression attenuates OGD-induced endothelial barrier functional disruption *in vitro*, while KLF11 knockdown reverses this effect.

KLF11 inhibits OGD-induced disruption of junctional proteins in cultured brain microvascular endothelial cells. To dissect functional roles and molecular mechanisms of KLF11 in stabilizing BBB integrity in response to ischemic stimuli, mouse primary BMECs were subjected to ischemic-like insult of OGD for 16 h, followed by 24 or 48 h reoxygenation. We found adenovirus-mediated overexpression of KLF11 blocked OGD-induced upregulation of pro-inflammatory cytokines IL-6 and MCP-1 at both the mRNA (Fig. S2 A-B) and protein levels (Fig. S2 C-D). Moreover, OGD stimuli significantly reduced the expression of TJ proteins ZO-1 and occludin in mouse BMECs (Fig. 7 A-D). As expected, adenovirus-mediated KLF11 overexpression successfully preserved the expression of both TJ proteins at the mRNA (Fig. 7 A-B) and protein (Fig. 7 C-D) levels, consistent with previous observations in EC-KLF11 Tg mice (Fig. 4). Immunocytochemical studies also confirmed decreased ZO-1 and occludin expression after 4h OGD treatment (Fig. 7E). This effect was markedly attenuated in mouse BMECs overexpressing KLF11 (Fig. 7E), indicating KLF11 overexpression preserves the integrity of tight junction proteins after OGD.

KLF11 regulates the expression of tight junction proteins ZO-1 and Occludin at the transcription level. As a member of the KLF family of transcription factors, KLF11 is known

to bind to GC-rich target sequences on promoters (38). We analyzed mouse ZO-1 and occludin promoters and identified three potential KLF11 binding sequences on each respective promoter (Fig. 8A), suggesting KLF11 may transcriptionally regulate ZO-1 and occludin. To confirm that KLF11 regulates ZO-1 and occludin through direct interaction with predicted binding sequences located in the promoter regions of these two genes, we cloned the promoter region of mouse ZO-1 and occludin, respectively into a dual-luciferase reporter vector (pGL4.10 [luc2]). We also clone the deleted promoters of mouse ZO-1 and occludin (without the three potential KLF11 binding sites) into the pGL4.10[luc2] vector (Fig. 8B, 8E). We then co-transfected mouse bEnd.3 cells with these luciferase reporter constructs and a Renilla luciferase control reporter vector (pRL-TK). We found adenovirus-mediated KLF11 overexpression significantly increased luciferase activity of the reporter constructs containing wild type ZO-1 (pGL4 mZO-1) and occludin (pGL4 mOccludin) promoter sequences (Fig. 8C, 8F), whereas adenovirus-mediated KLF11 knockdown significantly decreased the luciferase activity of these two promoter constructs (Fig. 8D, 8G). In addition, neither adenovirus-mediated KLF11 overexpression nor KLF11 knockdown affects the luciferase activities of the two deleted promoter constructs (pGL4 Δ mZO-1, pGL4 Δ mOccludin). Taken together, these results confirm that KLF11 transactivates ZO-1 and occludin by directly binding to their promoter region.

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# Discussion

This study investigated the protective function of endothelial KLF11 against BBB damage in a mouse ischemic stroke model. The results indicate that KLF11 transgenic overexpression in endothelium attenuates post-ischemic cerebrovascular permeability, brain water accumulation, and brain infarction, presumably through blocking the disruption of the junctional proteins, ZO-1 and occludin. By preserving the integrity of the BBB after ischemic insult, EC-targeted KLF11 overexpression effectively reduces the infiltration of neutrophils into brain parenchyma and the production of proinflammatory factors in the injured brain, thereby eliciting neurovascular protection against neurological deficits. Our results also demonstrate that KLF11 transactivates ZO-1 and occludin through direct binding to their promoters. Taken together, our findings suggest EC-targeted KLF11 transgenic overexpression is effective in mitigating BBB disruption and improving overall stroke outcome.

286 The functional involvement of KLF family members has been reported in various developmental 287 and pathological vascular processes (2, 52). KLF2 has been demonstrated to regulate endothelial 288 proliferation, migration, and angiogenesis (8, 9, 13). KLF2 has also been reported to regulate 289 vasoreactivity and vascular tone (13, 14, 43). Another KLF family member, KLF4, has been 290 reported as an endothelial regulator in response to pro-inflammatory stimuli (27) and shear stress 291 (40). A study using EC-targeted KLF4 transgenic and conditional knockout mouse models 292 demonstrated vascular anti-inflammatory and anti-atherothrombotic functions of KLF4 (68). 293 KLF4 was also reported to activate VE-cadherin at the transcriptional level, and thereby maintain normal endothelial barrier function (11). In addition, KLF6 has been demonstrated to play 294 295 functional roles in regulating vascular development, remodeling, and responses to injury (10, 12). 296 KLF14 was reported to mitigate atherosclerosis through modulating hepatic ApoA-I production 297 in mice (26). Moreover, KLF14 was demonstrated to reduce endothelial inflammation by 298 inhibiting the NF-κB signaling cascade (30). However, KLFs are less studied in the context of 299 cerebrovascular biology and diseases such as cerebral ischemia. 300 We have previously performed genome-wide screening for PPARy coregulators, and identified 301 KLF11 as a novel PPARy coregulator (62). We further confirmed a physical interaction between 302 PPARy and KLF11, and the regulatory effects of KLF11 on PPARy-mediated cerebrovascular 303 protection in primary BMEC cultures and mouse brain after in vitro (OGD) and in vivo (MCAO) 304 ischemic insults. Later (53), we provided direct evidence of KLF11 itself in the regulation of 305 ischemic brain injury. We found that conventional KLF11 KO mice had larger brain infarcts, 306 along with worsened neurobehavioral performance, increased edema, and greater BBB 307 disruption compared with WT mice (53). Our studies suggest KLF11 as an endogenous 308 protective mediator of ischemic stroke. However, the regulatory role of vascular KLF11 and the 309 underlying mechanisms of KLF11 itself in mediating protection against BBB leakage, especially 310 at the early onset (1-3 h) following ischemic insult, remain unknown. 311 In this manuscript, we found that KLF11 expression was significantly downregulated in cultured 312 BMECs and mouse cerebral microvessels after in vitro (OGD) and in vivo (MCAO) ischemic 313 insults (Fig. S1). Brain microvascular ECs are the major components of the BBB, and along with 314 pericytes, astrocytes, perivascular microglia, and the basal lamina, they form the neurovascular 315 unit (29, 31). Given the importance of brain microvascular ECs in the formation of the BBB as 316 well as in the pathogenesis of stroke, we therefore generated transgenic mice with vascular EC-

317 targeted transgenic overexpression of KLF11 driven by the Tie-2 promoter. Our results have 318 shown that KLF11 overexpression in ECs not only reduces MCAO-induced BBB permeability 319 and disruption (Fig. 2) but also leads to better neurological outcomes (Fig. 3). Moreover, in our 320 two-cell based in vitro BBB model, we determined that KLF11 overexpression effectively 321 preserved endothelial barrier integrity following ischemic-like conditions (Fig. 6). These findings 322 suggest that endothelial KLF11 contributes to its overall protective effects against ischemic 323 insults, therefore opening the possibility of potential therapeutic applications for KLF11. 324 There are numerous well studied mechanisms that contribute to ischemic-induced BBB 325 disruption, such as increased oxidative stress, vascular inflammation, activation of matrix 326 metalloproteinases (MMPs), loss of BBB cellular components, and abnormal pathologies of 327 endothelial tight junctions (33, 37, 44, 46, 49, 55, 60). A recent study demonstrated that MCAO-328 induced early structural disruption in brain microvascular ECs, including actin stress fiber 329 formation and redistribution of junctional proteins, resulted in impaired BBB integrity (49). This 330 early BBB disruption further facilitates MMP-mediated (especially gelatinase B/MMP-9) 331 degradation of EC tight junctions/basal lamina and the consequent infiltration of circulating 332 immune cells, leading to further BBB breakdown and secondary expansion of ischemic injury (41, 49). In our current study, we found that EC-targeted KLF11 transgenic overexpression 333 334 ameliorates MCAO-induced early (1-3 h) and late (24 h) onset of BBB impairment (Fig. 2E), 335 through preserving the integrity of tight junction proteins, ZO-1 and occludin (Fig. 5). It remains 336 to be determined whether and how endothelial KLF11 affects MMP-mediated late-stage BBB 337 disruption. 338 Brain ECs express TJ proteins at relatively high levels, thereby maintaining low permeability 339 between blood components and the cerebrospinal fluid (6). The three primary TJ proteins 340 expressed in brain ECs, occludin, claudins, and JAM, are tethered to the actin cytoskeleton by 341 scaffold proteins such as ZO-1, AF6, and cingulin (28). As a member of the zinc finger family of 342 transcription factors, KLF11 binds to GC-rich regions on gene promoters (38). We analyzed the 343 promoter regions of all TJ proteins, and discovered three KFL11 binding motifs on mouse ZO-1 344 and occludin promoters. Further, we confirmed the direct binding of KLF11 to these promoters 345 and transactivation activities on ZO-1 and occludin by dual-luciferase assays (Fig. 8). Although 346 in the current study we did not take into consideration that TJs are known substrates of MMPs 347 (37), whether EC-KFL11 mediated transactivation and upregulation of ZO-1 and occludin are

348 further regulated by MMPs need further investigation. Moreover, increased transcytosis in CNS 349 endothelial cells may also lead to BBB disruption (3). Ben-Zvi et al. reported that genetic 350 deletion of major facilitator superfamily domain containing 2a that is specifically expressed in 351 CNS ECs results in a leaky BBB due to increased transcytotic vesicles within ECs, without 352 affecting tight junctions (6). In our studies, EC-KLF11 Tg mice preserved relatively intact TJs 353 but still developed BBB breakdown to some extent after MCAO, suggesting the possible 354 involvement of transcellular pathway. Whether KLF11 plays a role in transcytosis needs further investigation. 355 Loosening of EC junctions due to ischemic insult facilitates the infiltration of circulating 356 357 immune cells and inflammatory factors, thereby producing serious clinical consequences other 358 than ischemic injuries, such as vasogenic edema and hemorrhagic transformation (49). We 359 previously observed that KLF11 KO mice show aggravated brain edema after ischemic insult 360 when compared with WT controls (53), whereas in the current study, EC-targeted KLF11 361 transgenic overexpression alleviates edema in the ischemic brain (Fig. 2I). We also observed less 362 infiltration of neutrophils into the ischemic brain of EC-KLF11 Tg mice (Fig. 2K). Moreover, 363 the expression of six inflammatory factors (TNF-α, IL-1β, MCP-1, IL-6, ICAM-1, p-selectin) 364 that are frequently upregulated after ischemic insult were found to be significantly reduced in the 365 ischemic brain of EC-KLF11 Tg mice compared with WT controls (Fig. 5). This is consistent 366 with our previous report that genetic deletion of KLF11 in mice further aggravated MCAO-367 induced upregulation of IL-6 (53), a major stroke-related pro-inflammatory cytokine. Our 368 observations are also consistent with one previous publication that demonstrated 369 lipopolysaccharide (LPS)-induced endothelial inflammation can be inhibited by KLF11 (19). 370 They reported that compared with WT mice, LPS-induced leukocyte rolling and adhesion on 371 postcapillary venular endothelium were further augmented in KLF11 KO mice. In addition, the 372 expression levels of VCAM-1 and E-selectin were significantly increased in the aortas of KLF11 373 KO mice (19). Therefore, KLF11-triggered anti-inflammation mechanisms may contribute to its 374 overall protective functions against BBB damage and ischemic neurovascular injuries. Taken 375 together, our results demonstrate that endothelial KLF11-mediated improvements in BBB 376 tightness or stabilization limited the infiltration of blood neutrophils and reduced the release of 377 inflammatory mediators into the ischemic brain, ultimately reducing secondary brain injuries

378 caused by neuroinflammation. Also, the regulatory effects of vascular KLF11 on other blood

379 immune cells need further attention.

380 In conclusion, our study demonstrates that EC-targeted KLF11 transgenic overexpression 381 mitigates both early and late BBB disruption through preserving the integrity of TJ proteins and 382 hindering the progression of neuroinflammation, leading to an overall functional improvement 383 after ischemic stroke. Our results may have potential clinical applications because cerebral 384 endothelial barrier disruption is important contributor to several pathological processes in the 385 brain, especially those associated with inflammatory brain disorders. Therefore, the findings of 386

this study may be applied to other disease processes including but not limiting to ischemic stroke.

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# Materials and methods

- 389 All procedures using laboratory animals were approved by University of Pittsburgh Institutional
- 390 Animal Care and Use Committee, and performed in accordance with the National Institutes of
- 391 Health Guide for the Care and Use of Laboratory Animals. All mice were randomly assigned to
- 392 various experimental groups using a lottery box. All surgical preparation, stroke outcome
- 393 assessments, neurobehavioral tests and data analysis were performed in blinded manner.
- 394 EC-KLF11 Tg and littermate wild-type control (WT) mice (male, 8-10 weeks-old, body weight
- 395 23-25g) were housed in a temperature- and humidity-controlled animal facility with a 12 h
- 396 light/dark cycle, and with unlimited access to food and water. Animals that did not show a more
- 397 than 75% CBF reduction or a less than 60% reperfusion over baseline levels or died after
- 398 ischemic induction (~10% of stroke animals) were excluded from further experimentation. In
- this study, we used 138 MCAO-operated mice (WT, n=69; EC-KLF11 Tg, n=69). Sham-399
- 400 operated mice were used as controls (WT, n=28; EC-KLF11 Tg, n=28).
- 401 Generation of EC-selective KLF11 transgenic (EC-KLF11 Tg) mice. Transgenic mice with
- 402 endothelial cell-selective KLF11 overexpression were generated as described (64). A 1539-bp
- 403 DNA fragment containing a KLF11 coding sequence was PCR amplified from human genomic
- 404 DNA using the primers 5'-GATATCATGCACACGCCGGACTTCGCAGGCC-3' (Forward)
- 405 and 5'-GATATCTCAGGCAGAGGCTGGCATGCTCACC-3' (Reverse). The DNA fragment
- 406 was then cloned into the EcoRV site of the pBluescript II SK (+) vector to generate a pBlue-
- 407 KLF11 plasmid. To generate the Tie-2 promoter-driven pBlue-KLF11 construct, we then
- 408 inserted the Tie-2 promoter (2,089 bp) (a generous gift from Dr. Sato, (47)) into the HindIII site

409 and the polyA plus full Tie-2 enhancer (10,367 bp) (a generous gift from Dr. Sato, (47)) into the 410 Xba I/Not I cloning site on pBlue-KLF11 plasmid, respectively. The construct was injected into 411 fertilized C57 mouse oocytes and implanted into pseudo-pregnant female mice. Transgenic 412 were founder \_mice\_ identified by **PCR** genotyping using primers 413 CTGTGCTCAGACAGAAATGAGAC-3' (forward) and 5'-ATCATCTGGCAAAGGACAGG-414 3' (reverse). The PCR amplification conditions were 94 °C x 5 min, 40 cycles of 94 °C x 30 sec, 415 55 °C x 30 sec, 72 °C x 80 sec, followed by 72 °C x 5 min. This produced a 1.2 kb DNA 416 fragment that contains both plasmid and Tie-2 promoter sequences. EC-KLF11 Tg mice are viable, fertile, normal in size, and do not display any gross physical or behavioral abnormalities. 417 418 Mouse model of transient focal cerebral ischemia. Focal cerebral ischemia was induced in 419 adult male mice (8-10-weeks-old, 25-30 g) by intraluminal middle cerebral artery occlusion 420 (MCAO) as described previously (53, 59, 61, 67). Briefly, mice were anesthetized with 1.5-3% 421 isoflurane (Henry Schein Animal Health). A 2-cm length of a 7-0 rounded tip nylon suture was 422 gently advanced from the internal carotid artery up to the origin of the middle cerebral artery 423 (MCA) until regional cerebral blood flow (rCBF) was reduced to less than 25% of baseline. 424 After 60 minutes of MCAO, blood flow was restored by removing the suture, and mice were allowed to recover for 1-7 days. In sham-operated mice, the same surgical procedure was 425 426 performed except for suture insertion. Changes in rCBF, arterial blood gases, mean arterial 427 pressure, and heart rate were monitored in animals 15 min before, during, and 15 min after 428 MCAO. Animals that did not show a CBF reduction of at least 75% over baseline levels were 429 excluded from further experimentation. Approximately 90% survival rate was observed in EC-430 KLF11 Tg or WT mice at 1-7 d after MCAO. Animals that died after ischemic induction were 431 also excluded. The rectal temperature was controlled at  $37.0 \pm 0.5$ °C during surgery. 432 Measurement of infarct volume, neurological deficits, and sensorimotor function. 2,3,5-433 triphenyltetrazolium (TTC) staining was performed to measure brain infarct after MCAO in 434 some mice. These mice were sacrificed, and the brains were harvested 24 h after MCAO. The 435 forebrain was sliced into seven coronal sections, each 1 mm thick. Sections were stained with 436 2% TTC in 0.9% NaCl for 20 min, followed by fixation with 4% paraformaldehyde in PBS. 437 Infarct volume was determined with ImageJ (National Institute of Health) as the volume of the 438 contralateral hemisphere minus the non-infarcted volume of the ipsilateral hemisphere.

439 Neurobehavioral deficits were determined by the adhesive removal test, foot fault test, and 440 rotarod test 1-3 days before MCAO and also at 3, 5 and 7 days of reperfusion after MCAO (53, 441 54, 59, 67). Following cerebral ischemia, mice were also tested for neurological deficits and 442 scored on a 5-point scale: 0, no observable neurological deficits (normal); 1, failure to extend 443 right forepaw (mild); 2, circling to the contralateral side (moderate); 3, falling to the right 444 (severe); 4, mice could not walk spontaneously; 5, depressed level of consciousness (very 445 severe). Immunohistochemistry and image analysis. At different time points following MCAO, EC-446 KLF11 Tg and WT mice were anesthetized with carbon dioxide and transcardially perfused with 447 448 0.9% NaCl followed by 4% paraformaldehyde in PBS. Brains were harvested and cryoprotected 449 in 30% sucrose in PBS, and frozen serial coronal brain sections (30 µm thick) were prepared on 450 a cryostat (CM1900, Leica). Brain sections were blocked with 5% normal goat serum in PBS for 451 1 h, followed by overnight incubation (4 °C) with the following primary antibodies: mouse anti-452 MAP2 (1:200; EMD Millipore, Billerica, MA), rabbit anti-NeuN (1:500; EMD Millipore), rat 453 anti-CD31 (1:200; BD Biosciences, San Jose, CA), mouse anti-ZO-1 (1:100; Invitrogen, 454 Carlsbad, CA), rat anti-Ly-6B (1:100; Abcam, Cambridge, UK), rabbit anti-Occludin (1:100; Thermo Fisher Scientific, Waltham, MA) (Supplementary Table S1). Secondary antibodies: 455 456 Cy3-conjugated goat anti-rat IgG, Alexa Fluor 488-conjugated goat anti-rabbit IgG, and goat 457 anti-mouse IgG (all at 1:400; The Jackson ImmunoResearch Lab). Images were collected by 458 confocal microscopy (FV1000-II; Olympus; Center for Biological Imaging, University of 459 Pittsburgh Medical School) and processed in Adobe Photoshop for compositions (17). Five 460 different sections from each animal were photographed, and six ROIs were randomly selected 461 from the infarct core and inner infarct border, respectively, on each section. Infarct volume was 462 measured on seven equally spaced MAP2-stained sections encompassing the MCA territory 463 using ImageJ. Capillary densities were examined by counting the number of capillaries stained 464 with the anti-CD31 antibody as previously described (56). 465 Quantification of BBB permeability/leakage after MCAO. For analysis of cerebrovascular 466 permeability by Evans Blue extravasation, mice were injected with 100 µl of 4% Evans Blue 467 (EB) (Sigma-Aldrich, St. Louis, MO) through tail vein 23 h after MCAO. After 1 h, mice were 468 perfused with 0.9% NaCl, mouse brains were then removed and separated into ipsilateral and

contralateral hemispheres. Each hemisphere was homogenized in N, N-dimethylformamide

- 470 (Sigma-Aldrich) and centrifuged for 45 min at 25,000 rcf. The supernatants were collected, and
- EB levels in each hemisphere were calculated using the formula:  $(A_{620 \text{ nm}} (A_{500 \text{ nm}} + A_{740 \text{ nm}}) /$
- 2) / mg wet weight. Background EB levels in the nonischemic hemisphere were subtracted from
- the ischemic hemisphere ipsilateral to the MCAO.
- 474 For analysis of cerebrovascular permeability by intravenous injection and detection of
- fluorescent-labeled Dextran and fluorescent tracer, mice were subjected to tail vein injection of
- either Tetramethylrhodamine-Dextran (TMR-Dextran, 70 kDa; 0.1 mg/g body weight,
- 477 Invitrogen) or Alexa Fluor 555-conjugated cadaverine (0.95 kDa; Invitrogen) 30 min before
- sacrifice. Mice were anesthetized and transcardially perfused with 0.9% NaCl followed by 4%
- paraformaldehyde in PBS. Mouse brains were collected and cryoprotected in 30% sucrose in
- 480 PBS, and frozen serial coronal brain sections (30 µm thick) were prepared on a cryostat. Brain
- sections were visualized directly under a fluorescent microscope. In parallel, brain hemispheres
- were homogenized in 1% Triton X-100 (Sigma-Aldrich) in PBS and fluorescent intensity was
- quantified by a SpectraMax i3x plate reader (Molecular Devices, San Jose, CA) using 555-nm
- 484 excitation and 580-nm emission.
- To measure the extravasation of endogenous IgG, sections were blocked in avidin and biotin
- 486 solution (two drops of avidin/biotin solution into 10 mL of PBS; Vector Laboratories,
- Burlingame, CA) for 15 min each, followed by blocking in 5% NDS for 1 h. Sections were then
- incubated with biotinylated anti-mouse IgG (1:500; Vector Laboratories) at 4 °C overnight.
- 489 Sections were then incubated with Alexa 488 Streptavidin (1:1,000; The Jackson
- 490 ImmunoResearch Labs). Whole-section images were acquired using an inverted Nikon
- 491 fluorescence microscope. Six equally spaced sections encompassing the MCA territory were
- 492 quantified for cross-sectional area of fluorescence. These areas were summed and multiplied by
- the distance between sections (1 mm) to yield a volume of leakage (in mm<sup>3</sup>).
- 494 Adenovirus-mediated gain- or loss- of- KLF11 expression in BMECs. To generate adenoviral
- vectors overexpressing KLF11 (19), the coding sequences of human KLF11 was PCR-amplified
- 496 with primers 5'-ATGCACACGCCGGACTTCGCAGG-3' (Forward) and 5'-TCAGGCA
- 497 GAGGCTGGCATGCTCA-3' (Reverse), and subcloned into the pCR8/GW/TOPO entry vector
- 498 (Invitrogen). To generate adenoviral vectors for overexpressing LacZ, the coding sequences of
- 499 Escherichia coli LacZ were PCR amplified with primers 5'-
- 500 ATGTCGTTTACTTTGACCAACA-3' (Forward) and 5'-

501	TTATTTTTGACACCAGACCAACT-3' (Reverse), and subcloned into the pCR8/GW/TOPO
502	entry vector (Invitrogen). After sequencing, the LR recombination reactions were carried out
503	between the entry clone pCR8/GW/TOPO/KLF11 and the destination vector pAd/CMV/V5-
504	DEST according to the manufacturer's protocol (Invitrogen). For knockdown experiments, an
505	siRNA oligo, which targets a region 100% conserved between human and mouse, was purchased
506	from Invitrogen. To prepare adenovirus-containing shRNA for KLF11 or LacZ, synthesized
507	oligos were annealed and inserted into the BLOCK-iT U6 entry vector. The U6 promoter and
508	shRNA were cloned into the adenoviral plasmid pAd/BLOCK-iT-DEST according to the
509	manufacturer's instructions. The sequences for shRNA were as follows:
510	shKLF11, 5'-CACCGGGTAGACTTTTCCCGAAGGCGAACCTTCGGGAAAAGTCTACC-3',
511	5'-AAAAGGTAGACTTTTCCCGAAGGTTCG CCTTCGGGAAAAGTCTACCC-3'.
512	shLacZ, 5'-CACCGCTACACAAATCAGCGATTTCGAAAAATCGCTGATTTGTGTAG-3',
513	5'-AAAACTACACAAATCAGCGATTTTTCGAAATCGCTGATTTGTGTAGC-3'.
514	To package the adenoviruses, adenoviral vectors were linearized with PacI and transfected into
515	HEK293 AD cells using Lipofectamine 2000. The recombinant adenoviruses were purified by
516	CsCl2 density gradient ultracentrifugation. Adenovirus genomic DNA was purified with
517	NucleoSpin Virus Kit (Macherey Nagel), the adenovirus titration was determined using the
518	Adeno-X <sup>TM</sup> qPCR Titration Kit (Clontech). The generated adenovirus was used to infect BMECs
519	for 48-72 h.
520	Two cell based in vitro BBB model. Mouse astrocytes were purchased from Sciencell research
521	laboratories (M1800-57). Astrocytes (2-8 passages) were grown in astrocyte medium (AM-a
522	1831; Sciencell) and were seeded in a regular 12-well plate at 37 °C in a humidified incubator
523	until reaching confluence. Transwell chamber inserts with PET membranes (0.4 µm pore;
524	Corning) were coated with collagen type IV (0.3 mg/ml) and fibronectin (0.5 mg/ml). Mouse
525	BMECs were seeded onto the membrane at a density of 1 x 10 <sup>5</sup> cells per insert. The inserts were
526	then placed into the 12-well plates containing confluent astrocytes (Figure 7A) and co-cultures
527	were maintained in endothelial growth medium (M1168; Cell Biologics) at 37 °C in a humidified
528	incubator overnight. The inserts seeded with mBMECs were then separated from the co-culture
529	system and mBMECs were infected with either Ad. LacZ/Ad. KLF11 or Ad. shLacZ/Ad.
530	shKLF11 in endothelial cell infection medium without phenol red and antibiotics (M1168PF;
531	Cell Biologics) for 6 hours. The inserts were placed back to the 12-well plate containing

532 confluent astrocytes and co-cultures were maintained in endothelial growth medium to reach confluence.

Changes in transendothelial electrical resistance (TEER) at designated time points or conditions were used to detect permeability changes in the BBB. The reading of total resistance ( $R_{Total}$ ) was measured with an Epithelial Volt/Ohm Meter (WPI, FL) at room temperature. The value of each sample ( $R_{TEER}$ ) was corrected by the reading of an empty coated-insert cell ( $R_{Blank}$ ) and calculated with the polyester membrane area ( $S_{Membrane}$ ) using the following formula and TEER values were reported in units of  $\Omega$ -cm<sup>2</sup>:

 $R_{TEER} = (R_{Total} - R_{Blank}) \times S_{Membrane}$ 

To assess paracellular permeability, Dextran Alexa Fluor 488 (3,000 MW) was added to the luminal chamber at a concentration of 1µg per 1ml medium. At 0.5h, 1h, 2h, 4h, 8h and 24h after OGD treatment, 50 µl medium was collected from the abluminal chamber. Fluorescence intensity was measured with a SpectraMax i3x Multi-Mode Detection Platform. The accumulated fluorescence intensity of abluminal Dextran Alexa Fluor 488 was measured at each time point and corrected by the respective blank (same treatment without adding Dextran in the luminal). Relative fluorescence unit (RFU) was recorded.

Cerebral microvessel isolation. Cerebral microvessels from mouse brain were isolated for determining the expression of KLF11, as previously described with modification (61). Briefly, EC-KLF11 Tg and littermate WT control mice were anesthetized with carbon dioxide and transcardially perfused with ice cold 0.9% NaCl. The brain was immediately removed from the skull and immersed in ice cold PBS. Brainstem, meninges and pia vessels were quickly removed, and the brain was cut into 1mm block and transfer to a 15 mL Dounce Tissue Grinder Tube (Kimble Chase, TN) with 5 mL buffer TE (a mixture of 0.25% Trypsin-EDTA and DMEM (Invitrogen, CA) at 1:1). The brain was homogenized by 5 strokes with a small clearance pestle. The homogenate was mixed with another 5 mL of buffer TE and incubate at 37 °C for 1 h with occasional agitation. After triturating 10-20 times with a 5 mL pipette, the creamy texture was centrifuged at 500 x g for 10 min at room temperature. The pellet was collected and dissolved in 8 mL HBSS (Invitrogen, CA) by mixing twice with a 5 mL pipette, and the homogenate was suspended in 11 mL HBSS dissolved 25% BSA by gently triturate 5 times with a 10 mL pipette. The homogenate was centrifuged at 3,000 x g for 15 min at 4 °C to separate the lipid and the capillary fraction. The pellet was collected and rinsed with 20 mL PBS once followed by another

- centrifuge at 1,800 x g for 10 min. The pellet was rinsed with 1 mL PBS, transferred to a 1.5 ml centrifuge tube and spun down at 16,000 x g for 1 min. The final microvessel pellet was stored at -80 °C freezer until use.
- Brain water content. Brain water content was measured by the dry-wet method as described previously (53). Following MCAO, mice were sacrificed by exposure to CO<sub>2</sub>. The weights of the ipsilateral and contralateral hemispheres were recorded separately as wet weights. The dry weights of the ipsilateral and contralateral hemispheres were obtained after being heated at 100 °C in an oven for 24 h. Brain content was calculated by the following formula: brain content = (wet weight dry weight)/wet weight x 100%.

- Cell cultures and oxygen-glucose deprivation (OGD). Mouse primary brain microvascular endothelial cells (mBMECs) were purchased from Cell Biologics (C57-6023). Mouse brain microvascular endothelial cells, bEnd.3 were purchased from American Type Culture Collection (CRL-2299, ATCC, Manassas, VA). Mouse BMECs and bEnd.3 cells (2-8 passages) were grown to 85-95% confluency before using for studies. To mimic ischemia-like conditions *in vitro*, mouse BMEC cultures were exposed to OGD for up to 16 h. Briefly, culture medium was replaced with deoxygenated glucose-free Dulbecco's Modified Eagle Media (DMEM; Gibco, Grand Island, NY), and cultures were flushed with 95% N<sub>2</sub> and 5% CO<sub>2</sub> for 5 min in a Billups-Rothenberg modular incubator chamber (Del Mar, San Diego, CA). The modular incubator chamber was then sealed and placed in a water-jacketed incubator (Forma, thermo Fisher Scientific, Waltham, MA) at 37 °C for the indicated period of time in each experiment before returning to humidified 95% air and 5% CO<sub>2</sub> and glucose-containing medium (62, 67).
- Immunocytochemistry. Mouse BMEC cells were seeded in 4-chamber polystyrene vessel tissue culture treated glass slides (Corning, NY) and infected with Ad. LacZ or Ad. KLF11. Seventy-two hours after infection, BMEC cultures were subjected to OGD treatment and reperfusion for indicated time points. The cells were then fixed in 4% paraformaldehyde followed by blocking with 5% normal goat serum in PBST for 1 h at room temperature. The cells were then incubated with the following primary antibodies overnight at 4 °C: mouse anti-ZO-1 (1:100; Invitrogen, Carlsbad, CA), rabbit anti-Occludin (1:100; Thermo Fisher Scientific, Waltham, MA). After rinse in PBS, cells were incubated with secondary antibodies: Alexa Fluor 488-conjugated goat anti-rabbit IgG, and goat anti-mouse IgG (all at 1:400; The Jackson ImmunoResearch Laboratories). After PBS rinses, cells were counterstained with DAPI for nuclear labeling and

- mounted with antifade Vecta-Shield solution (Vector Laboratories). Images were collected on an
- 595 Olympus FV1000-II confocal microscope and processed in Adobe Photoshop for compositions
- 596 (17). Immunofluorescence intensities of ZO-1 and occludin were quantified by ImageJ. Three
- ROIs were randomly selected from each chamber, and 2-3 chambers were quantified for each
- experimental condition in each independent culture.
- 599 **Molecular cloning.** The pGL4.10[*luc2*] vector was purchased from Promega. To make
- pGL4\_mZO-1 and pGL4\_mOccludin promoter reporter constructs, a 1,013 bp fragment of the
- promoter region of mouse ZO-1 or a 1,014 bp fragment of mouse occludin promoter region
- 602 containing the putative KLF11 binding sequence was PCR-amplified from mouse genomic DNA
- 603 using primers 5-CGAGACGCTAGCCTTGACTTTGAAACCTTAATTGATG and 5-
- 604 CTGGACCTCGAGGCAAAACCTGCCGGACCGGGCCACT, or 5-
- 605 CGAGACGCTAGCAGATAGTTAACTAACAAGAACTAAAATCTC and 5-
- 606 CTGGACCTCGAGCCTACCCCGGGCATGCGCACCAATT. Both promoter fragments were
- then subcloned into the pGL4.10[luc2] vector, respectively. Mutant promoter constructs where
- all three putative KLF11 binding sequences were deleted were generated by OuikChange XL
- 609 Site-directed Mutagenesis kit (Stratagene, Santa Clara, CA) (64). All constructs were validated
- 610 by DNA sequencing (Genewiz, South Plainfield, NJ). NCBI RefSeq ID: ZO-1 (mouse):
- 611 NP 001157046.1, occludin (mouse): NP 001347465.1. Supplementary Table S2.
- Dual-luciferase reporter assays. Mouse bEnd.3 cells (ATCC, Manassas, VA) were seeded at
- 0.5 x 10<sup>5</sup> cells/well in 24-well plates. After overnight incubation, cells were infected with
- different adenoviruses aimed at achieving KLF11 overexpression or knockdown. The cells were
- also co-transfected with pGL4 mZO-1/ΔmZO-1 or pGL4 mOccludin/ΔmOccludin luciferase
- reporter vector and a *Renilla* luciferase control reporter vector (pRL-TK; Promega), along with
- 617 Lipofectamine 2000 (Invitrogen) for 5 h. Luciferase activity was measured 48 h after transfection
- by Dual-Luciferase assay kits (Promega) using a SpectraMax i3x Multi-Mode Detection
- Platform. Individual luciferase activity was normalized to the corresponding *Renilla*-luciferase
- 620 activity (62, 64).
- Quantitative real time PCR. Total RNA was isolated from BMEC cultures or cerebral cortex
- by using RNeasy Mini Kit (Qiagen, Valencia, CA) or Trizol (Invitrogen, Carlsbad, CA).
- Quantitative real-time reverse-transcriptase polymerase chain reaction (RT-PCR) was carried out
- 624 with a Bio-Rad CFX Connect thermocycler, iScript cDNA synthesis kit and iTaq Universal

- 625 SYBR green supermix (Bio-Rad, Hercules, CA). Specific primers used for the reactions are ZO-
- 1 Forward, 5'-gccgctaagagcacagcaa-3'; ZO-1 Reverse, 5'-tccccactctgaaaatgagga-3'. Occludin
- 627 Forward, 5'-tgaaagtccacctccttacaga-3'; Occludin Reverse, 5'-ccgcataaaaagagtacgctgg-3'. TNF-α
- 628 Forward, 5'-etecteaccacacegteage-3'; TNF-α Reverse, 5'-aacacccattecetteacagagea-3'. IL-1β
- 629 Forward, 5'-aggagaaccaagcaacgacaaaatac-3'; IL-1β Reverse, 5'-tggggaactctgcagactcaaact-3'.
- MCP-1 Forward, 5'-gcaccagcaccagcaccagcacactctcact-3'; MCP-1 Reverse, 5'-cattccttcttggggtcagcacag-
- 3'. IL-6 Forward, 5'-agttgccttcttgggactga-3'; IL-6 Reverse, 5'-tccacgatttcccagagaac-3'. Icam-1
- Forward, 5'-tteacactgaatgccagctc-3'; Icam-2 Reverse, 5'-gtctgctgagacccctcttg-3'. P-selectin
- 633 Forward, 5'-gtccacggagagtttggtgt-3'; P-selectin Reverse, 5'-aagtggtgttcggaccaaag-3'.
- 634 Cyclophilin Forward, 5'-actcctcatttagatgggcatca-3'; Cyclophilin Reverse, 5'-
- 635 gagtatccgtacctccgcaaa-3' (Supplementary Table S3). The relative mRNA expression was
- normalized to cyclophilin RNA levels. PCR experiments were repeated 3 times, each using
- separate sets of cultures (53, 59, 67).
- Western blot analysis. Samples from the 0.9% NaCl perfused mouse cerebral cortex or cultured
- 639 mBMECs were homogenized in T-PER Tissue Protein Extraction Reagent (Pierce, IL)
- containing Mini Protease Inhibitor Cocktail (cOmplete<sup>TM</sup>, Sigma) by Dounce Tissue Grinder or
- sonication for 30 seconds with 30% pulse on ice respectively. The tissue or cell lysates were
- centrifuged at 10,000 x g for 5 min to pellet tissue or cell debris. Then supernatants were
- collected to measure protein concentrations by using the Bio-Rad Protein Assay (Bradford, Bio-
- Rad). As described previously (53, 59, 67), equal amounts of protein were loaded into 4-15%
- precast gels (Bio-Rad) and transferred to polyvinylidene difluoride (PVDF, Bio-Rad)
- membranes. Membranes were blocked for 1 h with TBS/0.1%-Tween buffer plus 5% (w/v) non-
- 647 fat dried milk and incubated overnight at 4 °C with primary antibodies diluted in the same
- blocking buffer. Membranes were then incubated with secondary antibodies diluted in blocking
- buffer for 1 h and developed using a Pierce® ECL Western blotting detection kit (Thermo
- 650 Scientific) and Amersham High-Performance Chemiluminescence Films (GE Healthcare).
- Various primary antibodies were used and are listed in Supplementary Table S1, including
- mouse anti-KLF11 (1:500; NovusBio), rabbit anti-occludin (1:1,000; Invitrogen), rabbit anti-ZO-
- 653 1 (1:250; Abcam), and mouse anti-β-actin (1:2000; Sigma-Aldrich).
- 654 Enzyme-linked immunosorbent assay (ELISA). Cerebral cortices of mice were collected and
- sonicated by an ultrasound homogenizer. After centrifugation, supernatants were collected and

- concentrations of a series of pro-inflammatory chemokines, cytokines, and adhesive molecules in
- 657 the supernatants were determined by Quantikine ELISA kits (R&D Systems, Minneapolis, MN)
- according to the manufacturer's instructions (67). All assays were performed in triplicates.
- 659 **Statistical analysis.** Quantitative data are expressed as mean ±SEM or ±SD. Differences among
- three or more groups were statistically analyzed by one or two-way ANOVA followed by
- Bonferroni's post-hoc test. Comparisons between two experimental groups were conducted by
- the two-tailed Student's t-test. A *p-value* less than 0.05 was considered significant. Statistical
- analyses and graphic representations were obtained with GraphPad Prism 7.0 software.

# Data availability statement

- The authors confirm that the data supporting the findings of this study are available within the
- article and from the corresponding author upon reasonable request.

- Abbreviation: Endothelial cell = EC; Krüppel-like factor = KLF; Blood-brain barrier = BBB;
- Peroxisome proliferator-activated receptor gamma = PPARγ; Tight junction = TJ; Brain
- 671 microvascular endothelial cells = BMECs; Oxygen-Glucose Deprivation = OGD;
- 672 Transepithelial/transendothelial electrical resistance (TEER); Middle cerebral artery occlusion =
- 673 MCAO; 2,3,5-triphenyltetrazolium chloride = TTC; Cerebral blood flow = CBF; Microtubule-
- associated protein 2 = MAP2.**Reference**
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- 883 Author contribution statement
- 884 X.Z., X.T., Y.E.C., and K.-J.Y. designed the research; X.Z., X.T., F.M., Y.F., T.Z., and J.Z.
- performed research; X.Z., X.T., F.M., and K.-J.Y. analyzed data; and X.Z., M.H.H, and K.-J.Y.
- wrote the paper.

- 888 Competing interests
- 889 None.
- 890 Figure Legends
- Figure 1. Generation of endothelial cell (EC)-selective KLF11 transgenic mice. (A) Schematic
- diagram of EC-selective KLF11 transgenic structure. The transgene cassette is composed of a
- 893 ~2.1-kb Tie-2 promoter, a ~10.4-kb Tie-2 enhancer, and a 1539-bp DNA fragment containing the
- 894 human KLF11 coding sequence. (B) Genomic PCR for genotyping EC-selective KLF11
- transgenic mice (EC-KLF11 Tg). A 960 bp band is expected from EC-KLF11 transgenics (Lanes
- 896 1, 3, and 16). (C) Representative western blotting images indicated enhanced protein levels of
- KLF11 in the cerebral microvessels (cerebral vessels, left) but not in brain parenchyma (cerebral
- cortex, right) of EC-KLF11 Tg mice than WT controls. n=5 mice per group. (D) Representative
- 899 immunofluorescent images show CD31-positive capillaries in the cortex and striatum of both
- 900 WT and EC-KLF11 Tg mice. n=5 mice per group. Scale bar: 50 µm. (E) Quantification of

- 901 CD31-labeled branch points, capillary numbers, and branch length are shown. In comparison 902 with WT controls, EC-targeted transgenic overexpression of KLF11 does not affect cerebral 903 vascular density and structure in mice. Data are expressed as mean  $\pm$ SEM. 904 **Figure 2.** Targeted overexpression of KLF11 in endothelium ameliorates both early- and late-905 onset BBB impairments, edema, and neutrophil infiltration. EC-KLF11 Tg mice and WT 906 controls were subjected to 1 h MCA occlusion followed by 1, 3, or 24 h reperfusion. (A) 907 Extravasation of Evans blue dye and (B) 70 kDa TMR-Dextran 24 h after MCAO in whole 908 brains or coronal sections was shown. Scale bar, 50 µm. (C-D) Quantification of Evans blue in 909 panel A and TMR-Dextran in panel B. n=6 mice per group. (E) Representative images 910 demonstrate the extravasation of Alexa Fluor 555 (0.95 kDa, red) or endogenous plasma IgG 911 (~150 kDa, green) into brain parenchyma 1, 3, or 24 h after MCAO. At 24 h after MCAO, the 912 area with loss of microtubule-associated protein 2 (MAP2) immunofluorescence illustrates the 913 infarct zone on adjacent sections from the same brains. (Scale bar: 1 mm). (F-G) Quantitative 914 analysis of the volume of the brain with leakage of cadaverine and IgG at indicated times of 915 reperfusion after MCAO. n=5 mice per group. (H) Brain infarct volume at 24 h after MCAO 916 was quantitatively measured on MAP2-stained coronal sections. n=5 mice per group. (I) Brain 917 water content was measured by the wet/dry weight protocol as described in the "Material and Methods". n=10 per group, \*\* p < 0.01 vs. WT contralateral hemisphere, # p < 0.05 vs. WT 918 919 ipsilateral hemisphere. (J) Representative images taken from the ipsilateral periinfarct cortex 920 after MCAO or the corresponding region after sham operation; markers used: Ly-6B (neutrophil), 921 and NeuN (neuron) (Scale bar: 50 µm). Rectangle: the region enlarged in high-power images 922 (third column). (Scale bar: 10 µm.) (K) Ly-6B+ cells were counted in the areas described in (J) 923 and data are expressed as the number of cells per mm2. After MCAO, the number of infiltrated 924 neutrophils is significantly less in EC-KLF11 Tg mouse brains compared with WT controls. n=5 mice per group. Data are expressed as mean  $\pm$  SEM. \* p < 0.05 vs. WT + sham group, # p < 0.05 925 926 vs. WT + MCAO group. 927 Figure 3. Transgenic overexpression of KLF11 in endothelium improves short-term histological 928 and functional outcomes after focal cerebral ischemia. (A) EC-KLF11 Tg mice and WT controls
- and functional outcomes after focal cerebral ischemia. (A) EC-KLF11 Tg mice and WT controls were subjected to 1 h MCAO and 72 h reperfusion. Two percent TTC-stained coronal sections were shown at different brain levels from the frontal to the posterior pole. (B) Quantitative analysis was performed on infarct volume and (C) neurological deficits in these mice after

932 ischemic stroke. n= 8 mice per group. (D) Regional CBF was measured by using a laser speckle 933 imager. EC-KLF11 Tg and WT mice were subjected to 1 h MCAO followed by 72 h reperfusion. 934 Representative CBF images were shown at 15 min before MCAO (baseline), 15 min after the 935 onset of MCAO (ischemia), and 15 min after the onset of reperfusion (reperfusion). (E) Two 936 identical elliptical ROIs were selected as indicated on the same brain region of the ipsilateral and 937 contralateral hemispheres. The relative CBF was first determined as the ratio of ischemic to 938 nonischemic cerebral blood flow, and then as the percentage value normalized to the presurgical 939 baseline for each animal. n=6 each group. (F-I) Sensorimotor deficits were assessed before and up to 7 days after MCAO by a battery of behavioral tests, including Rotarod test (F), Foot fault 940 941 (G), and Adhesive tape removal test (H-I). Compared with WT controls, mice with endothelial-942 specific overexpression of KLF11 showed dramatically improved sensorimotor function 943 (increased staying latency in the rotarod test, lower rate of fault steps in the foot fault test, and 944 shorter touching or removing time in the adhesive tape removal test. Data are represented as mean  $\pm$  SD. n=12 per group. \* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001 EC-KLF11 Tg MCAO vs. 945 946 WT MCAO by one-way ANOVA (individual time point) or two-way ANOVA (bracket). 947 Figure 4. EC-targeted KLF11 overexpression preserves the integrity of tight junctions after focal 948 cerebral ischemia. EC-KLF11 Tg mice and WT controls were subjected to 1 h MCA occlusion 949 and 1d or 3d reperfusion. (A-B) Total RNA was extracted from the ipsilateral cortex of EC-950 KLF11 Tg and WT mice 1d and 3d after MCAO. ZO-1 and occludin mRNA expression levels 951 were determined by qPCR and normalized to cyclophilin (n=6 per group). (C) Total protein was 952 extracted and subjected to gel electrophoresis. The protein levels of ZO-1 and occludin were 953 determined with β-actin as the loading control. (D-E) Quantification of ZO-1 and occludin 954 protein. Experiments were repeated three times and representative blots are displayed. n=6 per 955 group. (F) Representative images and quantification of ZO-1 (green) and occludin (green) on 956 CD31+ microvessels (red) in the cortex of the ischemic hemisphere at 1d after MCAO. (Scale 957 bar: 25 µm.) Endothelial KLF11 overexpression suppressed MCAO-induced disruption of 958 junctional proteins ZO-1 and occludin. n=6 mice per group. \* p < 0.05 vs. WT + sham group, # p 959 < 0.05 vs. WT + MCAO group. 960 **Figure 5.** Effects of EC-targeted overexpression of KLF11 on inflammatory mediators in mouse 961 cortex after focal cerebral ischemia. EC-KLF11 Tg and WT mice were subjected to sham 962 operation or 1h of MCAO followed by 1d or 3d of reperfusion. (A-F) mRNA expression levels

963 of different inflammatory markers were measured in the ipsilateral cortex by qPCR and 964 normalized to cyclophilin. (G-L) Protein levels of different inflammatory markers were analyzed 965 by ELISA. EC-targeted overexpression of KLF11 inhibited MCAO-induced upregulation of 966 TNF $\alpha$ , IL-1 $\beta$ , MCP-1, IL-6, ICAM-1, and P-selectin expression. n=6 mice per group. \* p < 0.05 967 vs. WT + sham group, # p < 0.05 vs. WT + MCAO group. 968 Figure 6. Adenoviral gain- or loss-of-KLF11 expression in mouse BMECs affects endothelial 969 barrier function after OGD in vitro. (A) Illustration of the transwell co-culture system. Mouse 970 astrocytes were seeded at the bottom of a 12-well plate with transwell insert. Cultured mBMECs were seeded on top of a collagen- and fibronectin-coated membrane of the insert and grown to 971 972 form a confluent monolayer. The mBMECs were infected for 48 h with either an empty 973 adenovirus (Ad. LacZ), adenoviral vector carrying mouse KLF11 (Ad. KLF11), adenoviral 974 vector shLacZ (Ad. shLacZ), or adenoviral vector carrying short hairpin sequences targeting 975 mouse KLF11 (Ad. shKLF11), and then subjected to OGD for 16 h before being placed back to 976 the plate with astrocytes. (B-C) Transepithelial electrical resistance (TEER) in the co-culture 977 model was measured at 2 h after OGD and in non-OGD conditions, n=3 per group, \* p < 0.05 vs. 978 Ad. LacZ/Ad. shLacZ + non-OGD group, # p < 0.05 vs. Ad LacZ/Ad. shLacZ + 16h OGD-2h 979 Reoxygenation. (D-E) Paracellular permeability was quantified at 0-24 h after OGD by 980 measuring the fluorescence intensity of abluminal Dextran Alexa Fluor 488 (3,000 MW). 981 Adenovirus-mediated KLF11 overexpression improved endothelial barrier function after OGD 982 treatment, whereas KLF11 knockdown worsened endothelial barrier function. Data are expressed 983 as mean  $\pm$  SEM of three independent experiments with triplicate wells (n=3 per group). \* P < 984 0.05 vs. Ad. LacZ/Ad. shLacZ + 16h OGD group. 985 **Figure 7.** Adenovirus-mediated KLF11 overexpression in mouse BMEC cultures preserves the integrity of tight junction proteins after OGD. Cultured mBMECs were infected with Ad. LacZ 986 987 or Ad. KLF11 for 48 h prior to 16 h or 4 h OGD followed by 24 h or 48 h reperfusion. (A-B) The 988 mRNA expression levels of ZO-1 and occludin were determined by qPCR and normalized to cyclophilin (n=3 per group). (C-D) Expression levels of tight junction proteins, ZO-1 and 989 990 occludin were evaluated by western blotting. β-Actin was used as an internal loading control. 991 Data represent three independent experiments and representative blots are displayed. n=3 per group. \* P < 0.05 vs. Ad. LacZ + non-OGD group, # P < 0.05 vs. Ad. LacZ + OGD group. (E) 992

Representative images and quantification of ZO-1 (green) and occludin (green) in cultured

995 KLF11 overexpression suppressed OGD-induced disruption of junctional proteins, ZO-1, and 996 occludin. n=3 per group. \* P < 0.05 vs. Ad. LacZ + non-OGD group, # P < 0.05 vs. Ad. LacZ + OGD group. H16R0 = 16h OGD + 0h reperfusion, H16R24 = 16h OGD + 24h reperfusion, 997 998 H16R48 = 16h OGD + 48h reperfusion, H4R0 = 4h OGD + 0h reperfusion, H4R24 = 4h OGD + 999 24h reperfusion. 1000 Figure 8. KLFII transcriptionally activates ZO-1 and occludin. (A) Potential KLF11 binding 1001 site in the promoter region of mouse ZO-1 and occludin. TSS: transcription start site. The green 1002 triangles mark the binding sites of KLF11. The numbers indicate the relative distance to TSS, 1003 which is labeled as +1. (B) 1kb of the mouse ZO-1 promoter or a mutated ZO-1 promoter 1004 (deleted all three KLF11 binding sequences,  $\Delta$  mZO-1) was cloned into the pGL4.10[luc2] vector. The pGL4 promoter constructs were then transfected into bEnd.3 cultures. bEnd.3 cells 1005 1006 were also co-transduced with an empty adenovirus (Ad. LacZ) or adenoviral vectors carrying 1007 mouse KLF11 (Ad. KLF11) for 48-72 h prior to performing luciferase reporter activity assays. (C) KLF11 overexpression in bEnd.3 cells significantly increased luciferase activity of 1008 pGL4 mZO-1, which contains all three KLF11 binding sequences (n=4 per group). (D) On the 1009 contrary, the luciferase activity was significantly reduced in the pGL4 mZO-1 group co-1010 transfected with Ad. shKLF11 (n=4 per group). (E) The occludin promoter or a mutated occludin 1011 1012 promoter (deleted all three KLF11 binding sequences,  $\Delta$  mOccludin) was cloned into the 1013 pGL4.10[luc2] vector. (F) KLF11 overexpression significantly increased luciferase activity of 1014 pGL4 mOccludin, but not pGL4 Δ mOccludin (n=4 per group). (G) On the contrary, luciferase 1015 activity was significantly reduced in the pGL4 mOccludin group co-transfected with Ad. 1016 shKLF11 (n=4 per group). Results shown are representative of three separate experiments with 1017 similar results. Data are expressed as mean  $\pm$  SEM. \* P < 0.05 vs. Ad. LacZ group or Ad. 1018 shLacZ group.

mBMECs after 4 h OGD, followed by 24 h reoxygenation. (Scale bar: 25 µm.) Adenoviral















