



LETTER

CD146 is essential for PDGFRβ-induced pericyte recruitment

Dear Editor,

The homeostasis of the central nervous system (CNS) is regulated through the blood-brain barrier (BBB), a protective barrier between the peripheral blood and the brain (Zhao et al., 2015). As vascular mural cells of brain microvessels, pericytes are positioned within the neurovascular unit among endothelial cells (ECs), astrocytes, and neurons. During the embryogenesis, pericytes are recruited to and interacts with the nascent CNS vascular ECs for BBB formation and maturation. The recruitment of pericytes to CNS vessels is hence critical for BBB development. A large amount of evidence has suggested that platelet-derived growth factor B (PDGF-B)/PDGFRβ signaling functions as the major regulatory pathway in governing pericyte recruitment (Bell et al., 2010; Daneman et al., 2010; Sweeney et al., 2016; Chen et al., 2017). Homozygous deletion of either Pdgfb or Pdgfrb in mouse embryos results in a complete lack of brain pericytes and CNS microhemorrhages, causing embryonic lethality (Daneman et al., 2010). In other studies, partial disruption of PDGF-B/PDGFR\$\beta\$ signaling leads to BBB breakdown and various CNS pathophysiologies associated with pericyte deficiency (Bell et al., 2010; Sweeney et al., 2016). These studies highlight a fundamental role of PDGF-B/PDGFRβ signaling in pericyte recruitment and CNS homeostasis. Limited studies have shown that in fibroblasts or mesenchymal stem cells, the full activation of PDGFRB requires its association with CD44, TGFβ type I receptor, neuropilin-1, or transglutaminase (Zemskov et al., 2009; Ball et al., 2010; Porsch et al., 2014), indicating that membrane proteins are essential for the functional signaling. However, the regulatory mechanisms governing PDGF-B/PDGFRB activation and how the activated PDGFR\$ induces its intracellular downstream signaling in brain pericytes remain to be elucidated.

CD146, which was originally identified as a novel endothelial biomarker for angiogenesis, has also been found constitutively expressed in pericytes and has subsequently been used for pericyte isolation and characterization (Crisan et al., 2008; Sweeney et al., 2016). Our recent study showed that pericyte-specific deletion of CD146 leads to BBB breakdown and impairment in pericyte recruitment due to the deficiencies of PDGFRB activation in mice (Chen et al., 2017). However, how pericytic CD146 regulates PDGFR\$ activation to transduce its downstream signals properly and efficiently during BBB development is still incompletely unrevealed. In endothelial cells (ECs), CD146 has been reported to activate the Fyn/Fak/paxillin pathway (Anfosso et al., 2001). It could also act as a co-receptor for VEGFR2 to facilitate VEGF-induced VEGFR-2 phosphorylation, AKT/ p38 MAPKs/NF-kB activation during angiogenesis, and this relies on its association with the cell cytoskeleton (Jiang et al., 2012). Besides, we have recently identified CD146 as a novel receptor for netrin-1 to promote p38 MAPKs/ERK activation and physiological angiogenesis (Tu et al., 2015). Above studies prompt us to elucidate the mechanisms and roles of CD146 in regulating PDGF-B/PDGFRß signaling in pericytes during BBB development.

In the present work, we employed various tools, including siRNA to silence the expression of Cd146, monoclonal antibody AA98 that specifically inhibits the function of CD146, dimerization-deficient mutant of CD146 or CD146 mutant lacking cytoskeleton binding site, to investigate the precise mechanism of CD146 in controlling PDGF-B/ PDGFRß signaling during BBB development in in vitro brain pericytic model and in vivo zebrafish model.

To investigate whether CD146 regulates PDGF-B/ PDGFRβ signaling, we incubated the pericyte progenitor cell line 10T1/2 with PDGF-B. In parallel, Cd146 siRNA which specifically knocks down the expression of CD146 was applied as a control. Our results showed that PDGF-B stimulation induced PDGFR\$\beta\$ phosphorylation and activated downstream AKT/JNK/ERK/p38 signaling. This activation was impaired by CD146 knockdown. Furthermore, the inactivation caused by knockdown of CD146 was rescued again by restoring CD146 expression (Figs. 1A and S1A). These results suggest that CD146 plays an important role in the PDGF-B/PDGFRß pathway.

To investigate the impact of CD146 dimerization on the downstream signal transduction of PDGFR\$\beta\$ in pericytes, we transfected 10T1/2 cells with Cd146/C454A or Cd146/ C501A to block dimerization of mice CD146. Our results showed that the PDGF-B-induced PDGFRβ phosphorylation and the downstream signal transduction were totally inhibited by either Cd146/C454A or Cd146/C501A (Figs. 1B and S1B), suggesting that CD146 dimerization is required for

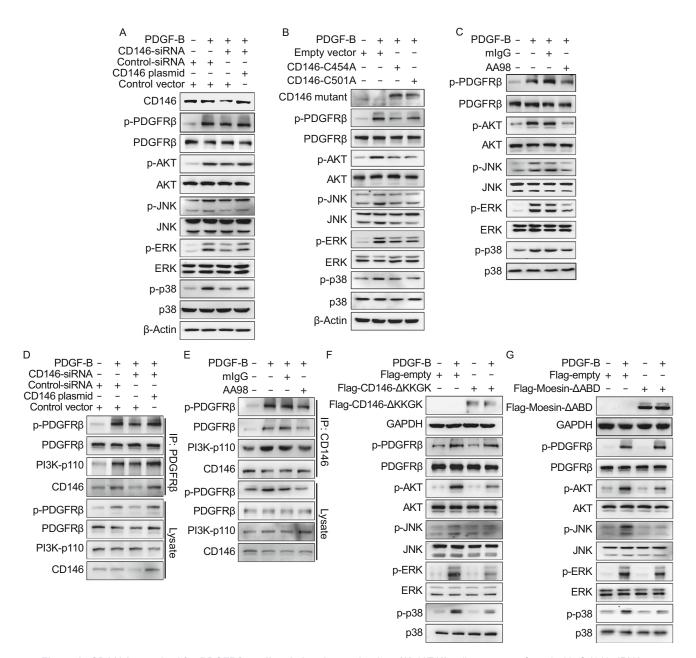


Figure 1. CD146 is required for PDGFRβ-mediated signal transduction. (A) 10T1/2 cells were transfected with *Cd146*-siRNA or co-transfected with *Cd146*-siRNA and a *Cd146*-expressing plasmid, and then stimulated with PDGF-B (20 ng/mL). CD146 expression and the activation of PDGFRβ, AKT, JNK, ERK, and p38 were detected by Western blotting. (B) *Cd146/C454A* or *Cd146/C501A* was transfected into 10T1/2 cells and the activation of PDGFRβ, AKT, JNK, ERK, and p38 induced by PDGF-B (20 ng/mL) was detected by Western blotting. (C) 10T1/2 cells were stimulated with PDGF-B (20 ng/mL) in the presence of mlgG or AA98 (50 μg/mL). The activation of PDGFRβ, AKT, JNK, ERK, and p38 was detected by Western blotting. (D) Co-immunoprecipitation (Co-IP) of endogenous CD146, PDGFRβ, p-PDGFRβ, and p110 induced by PDGF-B (20 ng/mL) was measured after the transfection of 10T1/2 cells with *Cd146*-siRNA or co-transfection with *Cd146*-siRNA and a *Cd146*-expressing plasmid. (E) 10T1/2 cells were stimulated with PDGF-B (20 ng/mL) in the presence of mlgG or AA98 (50 μg/mL). The interactions among endogenous CD146, PDGFRβ, p-PDGFRβ, and p110 were detected by Co-IP. (F) *Cd146*-Δ*KKGK* was transfected transiently into 10T1/2 cells and the activation of PDGFRβ induced by PDGF-B in addition to AKT, JNK, ERK, and p38 signaling was analyzed by western blotting. (G) *Moesin*-Δ*ABD* was transfected transiently into 10T1/2 cells and the activation of PDGFRβ after induction with PDGF-B in addition to AKT, JNK, ERK, and p38 signaling was examined. Data represent three independent experiments.

PDGF-B/PDGFR β signaling. To confirm the role of this dimerization on the signal activation of PDGFR β , we further applied anti-CD146 AA98, which has been reported to abrogate CD146 dimerization (Zheng et al., 2009). The results turned out that PDGF-B-activated PDGFR β phosphorylation and the AKT/JNK/ERK/p38 signaling were impaired by AA98 treatment, while the mlgG-treated cells were unaffected (Figs. 1C and S1C), indicating that CD146 dimerization plays an essential role in PDGFR β phosphorylation and AKT/JNK/ERK/p38 signal activation.

Since it has been demonstrated that PDGF-B engagement triggers PDGFRβ phosphorylation and subsequent PI3K recruitment for downstream signal activation, we next investigated whether CD146 was required for PDGFRβ/PI3K complex formation using AA98 and siRNA against *Cd146*. Co-immunoprecipitation revealed that CD146 binds to phosphorylated PDGFRβ (p-PDGFRβ) and PI3K in 10T1/2 cells. This association was enhanced by PDGF-B stimulation, whereas knockdown of *Cd146* (Figs. 1D and S1D) or AA98 pre-treatment (Figs. 1E and S1E) significantly impaired the complex formation. These suggested that CD146 regulates PDGF-B signaling by recruiting p-PDGFRβ and PI3K.

We previously demonstrated that downstream signal transduction and cell migration induced by CD146 requires physical interaction between CD146 and the cytoskeleton, which is mediated by actin-linking ezrin-radixin-moesin (ERM) proteins. The juxtamembrane KKGK motif of CD146 is required for its firm association with ERM proteins (Luo et al., 2012). Therefore, we asked whether this interaction is also essential for PDGFRB signaling. To this end, a Cd146 construct lacking the KKGK motif (coding Cd146-∆KKGK) or a Moesin construct lacking the actin-binding domain (coding Moesin-△ABD) was transfected into 10T1/2 cells. We observed that either Cd146-∆KKGK or Moesin-∆ABD impaired PDGF-B-induced PDGFRB downstream signaling without affecting PDGFR\$ phosphorylation (Figs. 1F, 1G, S1F, and S1G). In addition, transfection with *Cd146-∆KKGK* or Moesin-AABD caused a loss of PDGF-B-induced migratory and proliferation ability of 10T1/2 cells (Fig. S2). These suggest that the interaction between CD146 and the cytoskeleton, bridged by ERM proteins, is critical for PDGF-B-induced signal transduction, as well as cell migration and proliferation.

Since it has been reported that the PDGF-B/PDGFRβ signaling is important for pericyte recruitment to the vascular wall, we next used zebrafish (*Danio rerio*) as an *in vivo* model to investigate the impact of CD146 on the pericyte recruitment and the BBB development. For this, *cd146* and/ or *pdgfrb* were first knocked down using specific antisense oligonucleotide morpholinos (MOs). Next, fluorescent *in situ* hybridization using *pdgfrb* probe as a pericyte-specific marker was combined to investigate the recruited pericytes. As a result, we found that *cd146*-MO or *pdgfrb*-MO, alone or in combination, resulted in a significant decrease of pericyte recruitment at 72 h post-fertilization (hpf) (Figs. 2A and S3A),

demonstrating that CD146 and PDGFRβ synergistically regulate pericyte recruitment.

To further investigate the role of CD146 in BBB integrity, we micro-injected the tracing dye of rhodamine B-dextran (70 kDa) into the vasculature of *cd146*-MO zebrafish larvae at 3 dpf, when the function of BBB was initially formed. Our results showed that the dye leaked from the cerebrovasculature into the brain tissues in *cd146*-MO embryos. Knockdown of *pdgfrb* or co-injection of *cd146*-MO and *pdgfrb*-MO resulted in a similar phenotype of leakage. However, rescue of CD146 expression by co-injection of *cd146*-MO with exogenous *cd146* mRNA restored CNS vessel permeability, where the fluorescent dye was retained within the cerebrovasculature 30 min post-injection, similar to that observed in control larvae (Fig. 2B and 2C). This *in vivo* evidence indicates that CD146 is required for the functional formation of the BBB, and functions by regulating PDGFRβ pathway.

While since PDGFR\$\beta\$ is only expressed in pericytes, pdgfrb knockdown could not affect cerebral vasculogenesis nor EC number in zebrafish embryos before 56 hpf. In contrast, knockdown of cd146 or simultaneous treatment with cd146 and pdgfrb MOs resulted in severe defects in the cerebrovasculogenesis (Fig. S3). This is consistent with the early researches showing that CD146 is also expressed in ECs and is required for vascular sprouting and branching in zebrafish (Tu et al., 2015). Since CD146 is expressed in both ECs and pericytes, it is not easy to ascertain a causal relationship between pericytic cd146 knockdown and BBB integrity using cd146-MO zebrafish, as CD146 in ECs is also suppressed. The generation of pericyte-specific cd146 knockout zebrafish may be further needed to answer this question. However, combining the in vitro evidence that siRNA against Cd146 or anti-CD146 AA98 treatment in pericytes blocked PDGFR\u00e3-mediated signal transduction, we confirmed that CD146 is required for pericyte recruitment by regulating the PDGFR_{\beta} pathway.

Our study demonstrated that CD146, a reliable marker of pericytes, participates in PDGF-B/PDGFRβ-induced pericyte recruitment during BBB development. Based on the in vitro studies, we conclude that CD146 regulates PDGF-B/ PDGFRB signaling in a multifaceted manner, involving CD146-PDGFRß binding, CD146 dimerization, PI3K recruitment, and CD146-cytoskeleton binding. Combined with the zebrafish in vivo studies showing that either knockdown of cd146 or pdqfrb impaired pericyte recruitment and BBB integrity, we propose a model for CD146 to sequentially regulate PDGF-B/PDGFRB signaling and the recruitment of pericytes during BBB formation: within the cell membrane, the dimerization of CD146 associates with PDGFRB to enable efficient PDGF-B-induced receptor autophosphorylation, thereby enhancing the recruitment of PI3Kp110 to the p-PDGFRβ; subsequently, the CD146-ERM-cytoskeleton complex functions as scaffold for diverging signaling cascades downstream of PDGFRB, thus promoting pericyte recruitment and BBB integrity.

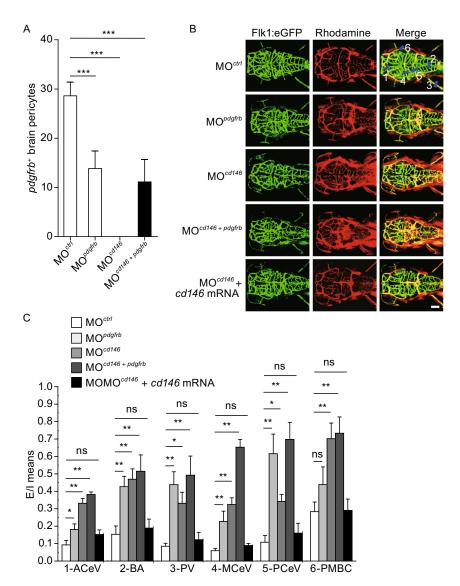


Figure 2. Knockdown of cd146 impairs BBB integrity and pericyte recruitment via PDGF-B signaling in zebrafish. (A) Quantification of $pdgfrb^+$ pericytes attached to cerebrovascular walls in zebrafish injected with cd146 morpholino, pdgfrb morpholino, or co-injected with cd146 and pdgfrb morpholinos. (B) Cerebral microangiography to analyze the BBB permeability of the 72 hpf zebrafish larvae injected with cd146 morpholino, pdgfrb morpholino, or co-injected with cd146 and pdgfrb morpholinos or with cd146 morpholino and cd146 mRNA. Scale bar represents 100 μ m. (C) Quantification of the Rhodamine signal for the intensity value of the vasculature lumen (I) relative to that out of the cerebral vasculature (E) in different central nervous system vessels of zebrafish injected with morpholinos as indicated. For each analyzed vessel, five positions along the vessel were chosen for calculating the E/I value; $^*P < 0.05$, $^{**P} < 0.01$, $^{***P} < 0.001$. Data represent five independent experiments. ACeV, anterior (rostral) cerebral vein; BA, basilar artery; PV, pectoral vein; MCeV, middle cerebral vein; PCeV, posterior (caudal) cerebral vein; PMBC, primordial midbrain channel.

Previous studies have proposed several mechanisms through which PDGF-B induces PDGFR β activation in pericytes, including PDGF ligand diversity and availability as well as interplay with membrane proteins or co-receptors (Andrae et al., 2008). In this study, we observed an association between CD146 and PDGFR β , which enhances PDGFR β activation in pericytes. This is consistent with numerous studies demonstrating that the interaction

between PDGFR β and membrane proteins such as neuropilin-1, CD44, TGF β type I receptor, and transglutaminase, is essential for the functional signaling of PDGFR β . Therefore, the association with membrane proteins might be a common mechanism exploited by pericytes for PDGFR β activation. In conclusion, our study identifies the cell adhesion receptor CD146 as a novel regulator in PDGF-B/PDGFR β signaling and sheds light on the mechanisms

governing pericyte recruitment in both the development of BBB and neurological disorders associated with BBB breakdown

FOOTNOTES

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Jianan Chen and Yongting Luo designed and performed experiments, analyzed data, and wrote the manuscript; Hongxin Huang, Shuilong Wu, and Jing Feng performed experiments and analyzed the data; Jingjing Zhang and Xiyun Yan initiated the study, designed experiments, reviewed the data, and wrote the manuscript.

Jianan Chen, Yongting Luo, Hongxin Huang, Shuilong Wu, Jing Feng, Jingjing Zhang, and Xiyun Yan declare that they have no conflict of interest. All institutional and national guidelines for the care and use of laboratory animals were followed.

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