Research Paper

Exosome-mediated delivery of an anti-angiogenic peptide inhibits pathological retinal angiogenesis

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Abstract

Background: Pathological angiogenesis is the hallmark of many vision-threatening diseases. Anti-VEGF is a primary treatment with substantial beneficial effects. However, such agents require frequent intravitreal injections. Our previous work established a method for effectively modifying exosomes (EXOs) for loading therapeutic peptides. Here, we used this system to load the anti-angiogenic peptide KV11, aiming to establish an EXO-based therapy strategy to suppress neovascularization in the retina.

Methods: Using an anchoring peptide, CP05, we linked KV11 to endothelial cell (EC) derived EXOs, yielding EXOKV11. We tested the delivery efficiency of EXOKV11 via two commonly used ocular injection methods: retro-orbital injection and intravitreal injection. Deploying an oxygen-induced retinopathy (OIR) model and a VEGF injection model, we tested the effects of EXOKV11 on neovascular formation, EC proliferation, and vascular permeability. In vitro experiments were used to test the mechanism and to analyze the effects of EXOKV11 on EC proliferation, migration, and sprouting.

Results: By using the EXO loading system, KV11 was more efficiently delivered to the blood vessels of the mouse retina via retro-orbital injection. In both OIR model and VEGF injection model, EXOKV11 was more effective than KV11 alone in inhibiting neovascularization and vessel leakage. The therapeutic effect of retro-orbital injection of EXOKV11 was comparable to the intravitreal injection of VEGF-trap. Mechanistically, KV11 alone inhibited VEGF-downstream signaling, while EXOKV11 showed a stronger effect.

Conclusions: We used EXOs as a carrier for intraocular delivery of KV11. We showed that KV11 itself has an anti-angiogenic effect through retro-orbital injection, but that this effect was greatly enhanced when delivered with EXOs. Thus, this system has the potential to treat proliferative retinopathy via retro-orbital injection which is a less invasive manner compared with intravitreal injection.

Key words: Exosomes (EXOs), drug delivery, proliferative retinopathy, pathological angiogenesis, VEGF signaling.

Introduction

Angiogenesis is vital for embryonic development [1, 2], and abnormal blood vessel growth contributes to many diseases, such as tumors and neovascular eye diseases [3, 4]. Pathological retinal neovascularization is a key pathological change in diseases such as retinopathy of prematurity (ROP)
and proliferative diabetic retinopathy (PDR), which are the major causes of vision loss in preterm infants and working-age adults [5-7]. Abnormal blood vessel growth is also characterized by compromised blood-retinal barrier (BRB) function, which further leads to hemorrhage, retinal detachment, and eventually blindness [8].

An elevated intraocular level of vascular endothelial growth factor (VEGF) is considered as the major cause that drives abnormal vessel growth and BRB breakdown. This has led to the use of intravitreal injection of VEGF inhibitors (such as the Fab-Fragment ranibizumab and VEGF decoy proteins aflibercept), which have been proven to have substantial beneficial effects [9, 10]. However, whether or not anti-VEGF is detrimental for the retina is controversially discussed. Although some studies have reported that anti-VEGF injection is safe [11, 12], others have had a concern that a major potential risk of this treatment is that during long-term therapy [13], intravitreal injections need to be repeated, which is invasive and may cause blinding sequelae such as endophthalmitis and retinal detachment [14, 15]. Furthermore, due to the neurosupportive properties of VEGF, some studies indicated that long-term of anti-VEGF treatment can also be associated with neuronal toxicity and developmental side effects [16-18]. Thus, it remains desirable to develop safer alternative therapeutic methods.

Exosomes (EXOs) are nanoparticles with sizes of 50 to 150 nm that are secreted by many different cell types [19]. EXOs are generated by the inward budding of endosomal membrane during maturation of multivesicular bodies (MVBs) and secreted into the extracellular environment upon fusion with the cell surface [20]. Exosomes may carry a large variety of cargoes, including cytosolic and cell-surface proteins, lipids, nucleic acids and metabolites [21]. Due to their natural, non-toxic, and biodegradable characteristics, EXOs are considered as ideal candidates for drug delivery to treat many diseases such as cancer, neurodegenerative disease, cardiovascular disease or muscular dystrophy [22, 23]. EXOs carrying propeptide enhanced muscle regeneration and growth in muscular dystrophy [24]. EXOs loaded with different cargoes, such as chemotherapeutics, multiple tumor-associated antigen or potent adjuvant, can enhance tumor recognition and suppress tumor growth [25, 26]. Moreover, their ability to cross various biological barriers, including the BRB, highlights the potential of using EXOs as drug delivery carriers that can penetrate formidable biological barriers [27, 28]. In the retina, EXO-associated AAV vectors have shown robust delivery efficiency into the murine retina compared with conventional AAV [29]. Thus, we aimed to test whether EXOs could be loaded with therapeutics and introduced into the eyes for treating retinopathies. HUVECs (human umbilical vein endothelial cells), which are derived from the umbilical cord endothelium, have been used extensively for studying the function of endothelial cells (ECs). We chose HUVECs to produce endogenous EXOs, with the intention of increasing the potency of anti-angiogenesis peptide delivery into the retina vasculature.

CD63 is a tetraspanin that is enriched on the surface of EXOs. Using phage display, we previously identified a peptide, CP05, which bound specifically to CD63 and enabled direct painting and effective modification of EXOs [30]. KV11, which consists of 11 amino acid residues from the human apolipoprotein(a) kringle 5-like domain (KV), showed a potential anti-angiogenic effect in vitro and in vivo [31, 32]. In this study, we hypothesized that by using CP05, KV11 could be loaded on EXOs and efficiently delivered to ECs, thereby improving its therapeutic efficacy. We demonstrate that this system has the potential to treat proliferative retinopathy in a highly effective but less invasive way.

Results

EXOs mediate efficient cellular uptake of KV11 in vitro

To test the feasibility of using the CP05-EXO system as a delivery tool in ECs, we first isolated HUVEC-derived EXOs by centrifugation and examined their properties. Transmission electron microscopy image showed that EXOs had a typical sauce-cup shape (Figure S1A). Nanosight showed a peak diameter of 112.0 nm, which is within the normal size range [23] (Figure S1B). By labeling EXOs with DiR, we observed that EXOs can be effectively taken up by HUVECs (Figure S1C). Western blotting showed that isolated EXOs possessed a high level of EXO-associated markers compared with whole cell lysate of HUVECs or supernatant, including Alix and CD81. CD63 is particularly abundant in the isolated EXOs, which should allow further modification using the CP05 peptide system (Figure S1D).

Next, we aimed to construct the EXO-linked KV11 peptide (EXOKV11) using CP05 as an anchor peptide (Figure 1A). For this, KV11-CP05 was incubated with EXOs for 6 h, followed by diaphragm to remove unbound peptides. For experiments intended to visualize EXOKV11, fluorescein isothiocyanate (FITC)-labeled KV11-CP05 was used. To evaluate whether CP05 can efficiently load KV11 onto EXOs, we performed flow cytometry for
EXOKV11. The results showed that 83.1% of KV11 peptides were anchored on EXOs, indicating an efficient EXO loading system using CP05 (Figure 1B).

Figure 1. Preparation of EXOs loaded with anti-angiogenesis peptide and efficiency of HUVEC delivery. (A) Schematic illustration of preparation of EXOKV11 by painting an anti-angiogenic peptide (KV11) on EXOs via the exosomal anchor peptide CP05. (B) Flow cytometry for measuring the binding efficiency of KV11-CP05 on EXOs. For checking cellular uptake, FITC-labeled KV11-CP05 was used. (C) EXOs were labeled with DiR and incubated with FITC-labeled KV11 (EXO+KV11) or FITC-labeled KV11-CP05 (EXOKV11) for 6 h at 4 °C. Each mixture was then added to HUVECs in culture. Representative confocal images from at least three different experiments show the cellular uptake of KV11 in HUVECs at 24 and 48 h. (D) EXO+KV11 and EXOKV11 prepared as in (C) were added to HUVECs in culture. After 24 or 48 h, cells were trypsinized for flow cytometry to analyze the cellular uptake efficiency of EXO+KV11 and EXOKV11. The red line represents the no-treatment control (Ctrl). The percentage of FITC-positive events out of total events is indicated in each histogram. Scale bars, 20 µm in (C).
We then tested the delivery efficiency of EXOKV11 compared with a mixture of KV11 and EXOs in the absence of CP05 (EXO+KV11). For this, DiR-labeled EXOs were used to visualize EXO uptake; FITC-KV11-CP05 or FITC-KV11 were then incubated individually with DiR-labeled EXOs (Figure 1C). After incubation, unbound peptides were removed by dialfiltration, and the products were then incubated with HUVECs or human retinal microvascular endothelial cells (HRMECs) for 24 or 48 h to test the delivery efficiency. Unsurprisingly, DiR-labeled EXOs could be taken up by both HUVECs and HRMECs (Figure 1C, Figure S2A). However, while strong FITC signals were observed in EXOKV11 (FITC-KV11-CP05 anchored EXOs), only very weak fluorescence signals were observed in EXO+KV11 (FITC-KV11 incubated with EXOs in the absence of CP05) group (Figure 1C, Figure S2A). Consistent with this, flow cytometry analysis also revealed that at 24 h, 100% of HUVECs and HRMECs were FITC-positive in EXOKV11-treated group, while in the EXO+KV11 group the proportion of FITC+ cells was only 0.395% and 29.1%, respectively (Figure 1D, Figure S2B). Additionally, there were still 98.0% FITC+ HUVECs and 100% FITC+ HRMECs at 48 h, indicating that this system was stable (Figure 1D, Figure S2B). The above results strongly suggested EXOKV11 as an efficient delivery system in vitro.

EXOs efficiently deliver an anti-angiogenic peptide into the retina vasculature

We then asked whether the above system could be used as a tool to deliver cargo into the retina vasculature in vivo. Intravitreal injection and retro-orbital injection are two commonly used ocular drug delivery methods, we tested the delivery efficiency of these two methods. Between them, intravitreal injection is known to be associated with the risk of retinal detachment, retinal hemorrhage and endophthalmitis [14, 15], while retro-orbital injection is a less invasive method with lower risks associated with the intravitreal route of administration [33-35].

A previous study has also proved that retro-orbital injection method to be effective for perfusing retina vessels [36], thus, we first used retro-orbital injection to determine the delivery efficiency. We injected EXOKV11 and KV11 into CD-1 mice, an albino mouse strain that allows us to acquire images with an IVIS imaging system (Figure 2A). Six hours after injection, abundant FITC signals were detected in the eyes of mice injected with EXOKV11, while a much weaker signal was observed in KV11-injected group, suggesting that EXOKV11 was delivered into the eye more efficiently than KV11 (Figure 2B). Furthermore, at 12 h, the FITC signal was still detectable (Figure 2B), indicating high stability of EXOKV11. We then checked the distribution of EXOKV11 in the retina. For this, 6 h after retro-orbital injection, the eyes were dissected, snap-frozen and cryosectioned. Blood vessels were labeled with Isoclectin B4 (IsoB4) (Figure 2C). Confocal images showed that the FITC signal was strongly enriched in blood vessels in EXOKV11-injected eyes, but much more weakly in KV11-injected ones (Figure 2C,D). Similarly, FITC signals were still detectable in the retina vasculature 12 h after injection, further highlighting the high stability of this system.

Intravitreal injection is another common method for delivering drugs into the eye. We next explored the delivery efficiency of EXOKV11 via intravitreal administration. To this end, we dissected eyes at 6 or 12 h after intravitreal injection of KV11 or EXOKV11. Confocal images showed that regardless of injection with either KV11 or EXOKV11, there was no clear co-localization of FITC signals and blood vessels (Figure S3).

Taken together, the above-mentioned results suggested that retro-orbital injection of a CP05-linked anti-angiogenic peptide and EXOs can be used as an efficient method for targeting retinal blood vessels, highlighting the possibility of treating pathological angiogenesis in the retina.

EXOKV11 suppresses neovascularization of the retina in the OIR model

The above results encouraged us to test the therapeutic effects of EXOKV11 in the oxygen-induced retinopathy (OIR) mouse model, which resembles human ROP and certain aspects of human PDR [37-39]. Mouse pups were exposed to hyperoxia conditions (75% oxygen) from postnatal day 7 (P7) to P12 as previously described [38] (Figure 3A). Hyperoxia causes oxygen toxicity and reduces the level of VEGF, which leads to vaso-obliteration. At P12, when the pups were returned to ambient air, the relatively hypoxic environment caused upregulation of VEGF, which then led to pathological neovascularization, consisting of overgrowth of immature vasculature and the consequent formation of extraretinal vascular tufts (Figure 3A) [40, 41]. To test whether EXOKV11 has anti-angiogenic effects in the OIR model, EXO, KV11 or EXOKV11 was retro-orbitally injected into the pups at P12. Saline vehicle (Ctrl) injected in the same way served as the control. Analysis of the avascular areas of the retina revealed that EXOKV11 did not alter vessel regression (Figure 3B,C). However, quantifying vessel tuft formation indicated that retinal neovascularization was significantly reduced in the KV11 and EXOKV11 groups (Figure 3B,D) [42]. In addition, when we compared KV11 and EXOKV11, the latter showed a
stronger inhibitory effect on neovascularization in the OIR model (Figure 3B,D).

As we observed reduced neovascularization in the KV11- and EXO_{KV11}-treated OIR model, we next assessed whether EC proliferation was affected. For this, we injected 5-ethynyl-2′-deoxyuridine (EdU) to label proliferating cells (Figure 3E). The endothelial nucleus was stained with ERG1. KV11 only mildly reduced the number of proliferating ECs, as determined by the number of EdU\(^+\) ERG\(^+\) nuclei, while EXO_{KV11} showed a stronger reduction effect (Figure 3F). This effect was not due to the EXOs themselves as EXOs alone did not change EC proliferation significantly (Figure 3F). qPCR also showed that the expression of key molecular that drive neovascularization such as HIF1\(\alpha\) and VEGF were significantly reduced upon EXO_{KV11}-treatment, while EXOs alone and KV11 had no clear effect (Figure 3 G,H).

![Figure 2. Ocular delivery efficiency of EXO_{KV11}.](http://www.thno.org)

(A) FITC-labeled KV11 or EXO_{KV11} was injected into CD-1 mice retro-orbitally. Fluorescence of the eyes was imaged at 6 and 12 h after the indicated injection. Ctrl represents the uninjected group. (B) Quantification of fluorescence intensity of eyes in (A) (n = 11 - 12 mice/group). (C) Ocular distribution of KV11 and EXO_{KV11} in cryosections of retinas from CD-1 mice treated as in (A). For checking in vivo distribution, FITC-labeled peptides were used. Arrowheads indicate co-localization of ECs (red, IsoB4) and peptide (green, FITC). (D) Quantification of fluorescence intensity of FITC-peptide in IsoB4\(^+\) area in cryosections of retinas in (C) (n = 3 mice/group; at least 3 images per mouse were analyzed and the values were averaged. The pictures for analysis were taken from the whole retinas, including peripheral part and retinas next to the optic nerve). The data represent as mean ± SEM. \(*p < 0.05, \ ***p < 0.01, \ **p < 0.001,\) Mann-Whitney test in (B), student's t test (two-tailed) in (D). Scale bars, 20 µm.
Figure 3. EXOKV11 inhibits retinal neovascularization and suppresses EC proliferation in the OIR model. (A) Schematic depiction of the mouse OIR model. Pups were placed in 75% oxygen from P7 to P12, and then returned to normal oxygen conditions. Saline vehicle (Ctrl), EXO, KV11, or EXOKV11 were injected retro-orbitally at P12. Pups were sacrificed and eyes were dissected at P17. (B) Representative confocal images of retina vasculature stained with IsoB4 in OIR retinas of saline vehicle (Ctrl)-, EXO-, KV11-, and EXOKV11-treated pups at P17. Upper panels indicate the avascular area and lower panels show the neovascular tufts. (C and D) Avascular area and neovascularization quantification of (B) (n = 6 - 9 eyes per condition). (E) Representative confocal images of retina sagittal sections costained with IsoB4, EdU (labels proliferating cells), and ERG (labels EC nuclei) in pups treated as in (A). (F) Quantitative analysis of EdU* ERG+ cell number in (E) (n = 5 - 7 per condition). (G and H) qPCR analysis of the expression of Hif1α and Vegfa in OIR retinas of saline vehicle (Ctrl)-, EXO-, KV11-, and EXOKV11-treated pups at P17 (n = 5 per condition). The data represent as mean ± SEM, *p < 0.05, **p < 0.01, ***p < 0.001; one-way ANOVA followed by Tukey's multiple comparisons test in (C), (D), (F), (G). Scale bars, 1 mm in (B), 50 µm in (E).
**EXOKV11 suppresses vascular leakage in the retina of OIR model mice**

The abnormal vascular growth in OIR model mice also results in a breakdown of the BRB; thus, severe and extensive retinal hemorrhages were found in the retinas of OIR pups. This was dramatically decreased in EXOKV11-treated pups (Figure 4A). We further performed immunohistochemical analysis for TER119 (an erythrocyte marker) and IsoB4. EXOKV11 inhibited erythrocyte extravasation, as determined by quantifying TER119+ IsoB4- area in the whole-mount retina, while EXO or KV11 alone showed no significant effect (Figure 4B,C). In addition, FITC-dextran was injected to determine vessel leakage. Extravasated FITC-dextran was detected in OIR pups, but the degree of leakage was markedly decreased in EXOKV11-treated pups (Figure 4D,E). Again, the protective effect of EXO or KV11 was not significant (Figure 4A-E).

Tight junctions, including Claudin-5 and ZO-1 are essential for regulating vascular barrier function and determine vascular permeability [43]. In the OIR retina, the Claudin-5 and ZO-1 junction distribution was discontinuous. Strikingly, EXOKV11 treatment clearly rescued the junction defects, indicating that EXOKV11 treatment protecting vascular leakage by preventing destruction of tight junctions (Figure 4F).

To determine the inflammatory status, we analyzed macrophage infiltration in the whole-mount OIR retina. The number of F4/80+ macrophages in both the KV11 and EXOKV11 treated group decreased, but the latter had a stronger effect (Figure S4A,B). Consistently, the expression of interleukin-6 (IL-6) and vascular cell adhesion molecular-1 (VCAM-1) was markedly reduced by EXOKV11 treatment (Figure S4C,D). Thus, these results suggested that EXOKV11 limits inflammation in the OIR retina.

**EXOKV11 suppresses VEGF-induced vascular leakage in vivo**

In the mouse OIR model, increased VEGF secretion caused by hypoxia is the major cause of neovascularization and vascular leakage. Thus, we investigated whether EXOKV11 could also suppress vascular endothelium activation by VEGF in an in vivo model. We first explored whether EXOKV11 has prevention effects in VEGF-induced vascular model. For this, EXOKV11, KV11, EXO or vehicle was retro-orbitally injected into adult C57BL/6j mice. Twenty-four hours later, 100 ng VEGF was intravitreally injected (Figure 5A). Vascular leakage was visualized by FITC-dextran perfusion. VEGF-induced vascular leakage was strongly suppressed by EXOKV11, while KV11 was less effective (Figure 5B,C). When we intravitreally injected a VEGF-trap, which is currently being used for treating neovascular retinal disease, we found that it also significantly inhibited VEGF induced vascular leakage in our mouse model, and the effect of retro-orbital injection of EXOKV11 was comparable with the effect of intravitreal injection of the VEGF-trap (EXOKV11 50.71% ± 7.18% reduction vs. VEGF-trap 53.09% ± 8.62% reduction) (Figure 5B,C and Figure S5A,B). Consistently, VEGF injection affected the organization of Claudin-5, and this was protected by EXOKV11 treatment (Figure 5D). We also performed immunofluorescence staining for F4/80 and the endothelial marker CD105. We found that although F4/80+ macrophages were abundantly enriched in the retinas of VEGF-treated mice, these macrophages were scarce in EXOKV11-treated retinas, suggesting that EXOKV11 also limits inflammation in the VEGF-treated retina (Figure 5E,F). Again, the effect of retro-orbital injection of EXOKV11 was comparable with intravitreal injection of the VEGF-trap (EXOKV11 60.04% ± 8.36% reduction vs. VEGF-trap 62.87% ± 10.0% reduction), while retro-orbital injection of KV11 was less effective (Figure 5E,F and Figure S5C,D). By using this pre-treatment model, our results indicated that EXOKV11 improved vascular integrity and suppressed inflammation in the VEGF-treated mouse retina.

The previous experiments revealed EXOKV11 prevent vascular leakage after VEGF injection. Next, we investigated whether it has also therapeutic effect by using a post-treatment model. For this, we injected 100 ng VEGF intravitreally to induce vascular leakage. 24 h later, EXOKV11, KV11, EXO or vehicle was retro-orbitally administrated (Figure 6A). Vascular leakage was visualized by FITC-dextran perfusion 48 h after the treatment. Analysis of dextran extravasation showed that in the post-treatment model, EXOKV11 significantly inhibited vascular leakage (Figure 6B,C). Similarly, quantification of F4/80+ macrophage demonstrated that EXOKV11 dramatically reduced macrophage infiltration (Figure 6D,E). Again, KV11 also had an inhibitory effect, but was less effective (Figure 6B-E). Collectively, we showed that EXOKV11 treatment is not only useful when given before VEGF injection, but as a post-treatment strategy can also inhibit vascular leakage and limit inflammation. Additionally, an auricular Miles assay showed that EXOKV11 reduced VEGF-induced vascular leakage, suggesting a conserved protection mechanism in the mature vasculature (Figure 5G).
Assessment of toxicity in EXOKV11-treated mice

We next examined the toxicity of EXOKV11 to the retina as well as the major organs, including the liver, spleen, lung, and kidney. TUNEL assay showed that TUNEL+ cells in the retina of the EXOKV11 treated mice were not more than those in saline-treated controls in either the OIR model (Figure 7A) or the VEGF-induced vascular leakage model (Figure 7B). Histological analysis also showed no evidence of in vivo toxicity of EXOKV11 in either the model (Figure S6A,B). The concentrations of the hepatic enzymes alanine aminotransferase (ALT) and alanine aspartate aminotransferase (AST) in the serum of EXOKV11-treated mice were also not significantly different from those in saline-treated controls in either model (Figure S6C-F), suggesting that EXOKV11 was not significantly toxic.
**EXOKV11 suppresses VEGF-downstream signaling in ECs**

Although the mechanism of how KV11 inhibits angiogenesis is not clearly understood, it has been shown KV11 suppressed extracellular-signal-regulated kinase (ERK) activation as well as Src activation under VEGF stimulation [31]. As ERK is crucial in regulating EC proliferation [44, 45], while Src is involved in regulating cell migration,
endothelial junctions, and barrier function [46], we then performed western blot to test whether EXO_{KV11} could also suppress such VEGF-downstream signaling in HUVECs. Consistent with the previous report [31], we indeed observed that KV11 reduced ERK1/2 phosphorylation in response to VEGF stimulation. This effect was more robust in HUVECs pretreated with EXO_{KV11} (Figure 8A,B). We also found that EXO_{KV11} significantly reduced Src phosphorylation upon VEGF stimulation, while KV11 showed only a milder reduction (Figure 8C,D). VEGF also induces phosphorylation of Y685 of VE-Cadherin, which is involved in vascular permeability [47]. Pretreatment with KV11 also suppressed VEGF-mediated Y685 phosphorylation in HUVECs, while EXO_{KV11} showed a stronger reduction (Figure 8E,F). Moreover, these effects were not observed in the EXO group, suggesting that the HUVEC-derived EXO itself has no clear effect in blocking VEGF-downstream signaling (Figure 8A-F).

Figure 6. EXO_{KV11} has therapeutic effect in post-treatment model. (A) Schematic depiction of the post-treatment procedure in VEGF-induced vascular leakage model. 100 ng VEGF was intravitreally injected to induce vascular leakage in the retina. After 24 h, EXO_{KV11}, KV11, EXO, or vehicle was retro-orbitally injected in adult wild type C57BL/6j mice. The mice were sacrificed for analysis after another 48 h. (B) FITC-dextran was injected and the retinas were harvested after treatment. Representative images of flat-mounted retina show extravasated FITC-dextran and CD105+ vessels. (C) Quantification of dextran leakage in (B) (n = 4 - 6 mice per condition). (D) Representative images of F4/80+ macrophages (green) and CD105+ vessels in retinas treated as in (A). (E) Quantification of macrophage infiltration in (D) (n = 4 - 6 mice per condition). The data represent as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001, one-way ANOVA followed by Tukey’s multiple comparisons test in (C), (E). Scale bars, 50 µm.
Figure 7. Assessment of eye toxicity in EXOKV11-treated mice. (A and B) Representative TUNEL assay images of eye sections from saline-treated controls (Ctrl), EXO-, KV11-, or EXOKV11-treated mice in the OIR at P17 (A) and pre-treated VEGF-induced vascular leakage model at indicated endpoint (B).

Figure 8. EXOKV11 suppresses VEGF-induced endothelial proliferation and VE-cadherin phosphorylation. (A, C) Starved HUVECs were pretreated with EXO, KV11, or EXOKV11 for 24 h and then stimulated with 25 ng/mL VEGF for 10 min. Representative western blots for pERK (A) and pSrc (C) showing VEGF-downstream signaling. Conbercept (VEGF-trap) was used as a positive control, the pretreatment time was 5 min before VEGF stimulation. 30 μg total protein from each sample was loaded. (E) Starved HUVECs were pretreated with EXO, KV11, or EXOKV11 for 24 h and then stimulated with 25 ng/mL VEGF for 15 min. Representative western blot for phosphorylation of VE-cadherin at site Y685 showing how the indicated pretreatment affects VEGF-induced phosphorylation of VE-cadherin at Y685. Conbercept (VEGF-trap) was used as a positive control, the pretreatment time was 5 min. 30 μg total protein from each sample was loaded. (B, D, and F) Quantification of blots shown in (A), (C), and (E); n = 3 - 5 independent experiments. The data represent as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001, one-way ANOVA followed by Tukey's multiple comparisons test in (B), (D), (F).
EXOKV11 inhibits VEGF-induced EC proliferation, migration, and sprouting in vitro

Our data indicated that EXOKV11 suppressed VEGF-downstream signaling in cultured ECs. To further characterize the EC angiogenesis phenotype after EXOKV11 treatment, we performed a set of in vitro angiogenesis assays. EXOKV11-treated mouse retinas in the OIR model showed reduced proliferation (Figure 3E). To test whether EXOKV11 has a direct inhibitory effect on EC proliferation, we performed a 5-Bromo-2-deoxyuridine (BrdU) incorporation assay in HUVECs. In ECs without VEGF stimulation, EXOKV11 already showed an inhibitory effect on proliferation, while KV11 alone had no obvious effect. After VEGF stimulation, EXOKV11 pretreatment showed a more pronounced inhibitory effect, and the difference between KV11 and EXOKV11 became apparent, suggesting that EXOKV11 already has an inhibitory effect in the basal state, and this effect was more marked after VEGF stimulation (Figure 9A,B).

Figure 9. EXOKV11 suppresses VEGF-induced angiogenic effects. (A) Starved HUVECs were pretreated with EXO, KV11, or EXOKV11 and stimulated with VEGF for 24 h. Staining for BrdU incorporation shows proliferating HUVECs. (B) Quantification of BrdU+ nuclei in (A). The proportion of BrdU+ nuclei among total DAPI+ nuclei was determined and then normalized to the control condition. n > 8 biological repeats were analyzed. (C) Representative bright field images of the scratch migration assay of HUVECs treated as in (A) after 12 h of VEGF stimulation. (D) Quantification of the gap closure from (C). n = 8 biological repeats were quantified, and the results were normalized to the control condition. (E) Representative images of the bead-sprouting assay using HUVECs pretreated with EXO, KV11, or EXOKV11 and treated with 50 ng/mL VEGF for 24 h. (F) Quantitative analysis the number of sprouts and total sprouting length in (E). Approximately 20 beads per condition were quantified. The data represent as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001, one-way ANOVA followed by Tukey’s multiple comparisons test in (B), (D), (F). Scale bars, 50 µm in (A and C), 200 µm in (E).
Expansion of vascular networks also requires EC migration. Therefore, we also tested whether EXO$_{KV11}$ could affect EC migration. For this, we performed a wound scratch assay. In the basal state without VEGF stimulation, while EXOs or KV11 alone did not have a significant effect on EC migration, EXO$_{KV11}$ significantly inhibited wound closure. After VEGF stimulation, KV11 alone showed a significant inhibitory effect on wound closure, while EXO$_{KV11}$ showed a more robust inhibitory effect and the difference was highly significant (Figure 9C,D).

We also analyzed EC sprouting in vitro using the beads sprouting assay [2]. VEGF stimulation significantly increased the number of sprouts and the total sprouting length. EXO$_{KV11}$ treatment successfully reduced both parameters in this assay system (Figure 9E,F).

Collectively, these results indicate a strong anti-angiogenic effect of EXO$_{KV11}$ in ECs, especially in response to VEGF stimulus.

Discussion

In this study, we constructed an EXO-based anti-angiogenic peptide delivery system and assessed its feasibility, safety, and efficacy. We showed that this EXO$_{KV11}$ system targeted the retina blood vessels, and reduced pathological angiogenesis as well as vascular leakage. Importantly, EXO$_{KV11}$ can be efficiently delivered into the retina vasculature by retro-orbital injection, a less invasive procedure than intravitreal injection, which is a commonly used method in the clinic [33-35]. Our findings suggest that EXO$_{KV11}$ will work as an effective nanotherapeutic for treating pathological angiogenesis in retinopathy.

KV11 is an 11-amino acid peptide from the KV domain of apolipoprotein(a), a known anti-angiogenic factor [31]. KV11 peptide shows anti-angiogenic activity not only in vitro, but also in vivo in several animal models, including the chicken chorioallantoic membrane model, the mouse corneal micropocket angiogenesis model, and also the mouse xenograft tumor model [31]. In the retina, KV11 also shows an anti-angiogenesis effect in the mouse OIR model. However, to achieve this, KV11 must be injected intravitreally and the treatment requires repeated injections [32]. In this study, therefore, we sought a better solution.

EXOs have attracted considerable attention as drug delivery vehicles due to their low immunogenicity and high safety profile. They are also relatively stable with a longer half-life in the blood circulation, as they avoid opsonins, coagulation factors, complement, and phagocytosis by circulating monocytes and macrophages [48]. Furthermore, the membrane structure of EXOs enables them to cross the cell membrane system and to be delivered into the recipient cell. Although the mechanisms of EXO uptake and of components delivery into acceptor cells, whether they involve membrane fusion or endocytosis and whether the process has cell type targeting specificity, are still incompletely characterized [49], several studies have suggested that the surface components on EXOs and acceptor cells, such as integrins, tetraspanins, and phosphatidylinositol 4,5-bisphosphate [50-53], probably influence EXO target recognition and cellular uptake by target cells. Physiologically relevant specific targeting or homing remains to be achieved. Notably, however, some studies have already indicated that EXOs have the ability to target their parent cells and to serve as selective vehicles. For example, it has been shown that the U251 glioblastoma cells derived EXOs exhibit higher internalization efficiency in U251 cells than astrocyte derived EXOs [54]. It was also shown that human placental mesenchymal stem cells (MSCs) derived EXOs preferentially go to the cell type of origin [55]. Other studies also indicated that EXOs derived from certain tumor cells have the ability to home to their parent tumors [56, 57]. Thus, considering EXOs natural homing properties, we chose ECs derived EXOs as carriers for blood vessel delivery in this study, as they would be more effective at homing to where they were secreted from than to other cells.

Using an anchor peptide that we had previously identified [30], therapeutic amounts of KV11 were loaded onto EXOs and elicited a functional inhibitory effect on pathological retinal angiogenesis. Importantly, in the OIR model, instead of two intravitreal injections of KV11 as previously reported [32], a single retro-orbital injection of EXO$_{KV11}$ efficiently rescued the vascular defects without any detectable toxicity, indicating that the EXO is an efficient delivery vehicle for treating vascular dysfunction in the retina.

Functioning as carriers of small bioactive molecules, EXOs have emerged as important players in cell-cell communication. It has been shown that EXOs derived from different cell types can deliver pro- and anti-angiogenic factors including proteins, mRNA, and microRNAs into ECs and can thus regulate angiogenesis. For instance, during tumor progression, EXOs secreted from tumor cells interact with ECs and stimulate angiogenesis in multiple tumor models [58-60]; this may be a critical step for tumor angiogenesis and metastasis. ECs also secrete EXOs. Interestingly, under different patho/physiological conditions, ECs secrete EXOs that have different contents and that exert different biological functions. For example, Delta-like 4 is incorporated
into endothelium-derived EXOs, which then signal to neighboring ECs, inhibit Notch signaling, and increase capillary formation [61]. It was also shown that extracellular vesicles released by ECs contain matrix metalloproteinases that stimulate ECs to form capillary-like structures [62]. On the other hand, another study revealed an anti-angiogenic effect of extracellular vesicle(s), namely that endothelium-derived microparticles can also impair angiogenesis by inhibiting EC proliferation and increasing apoptosis [63]. Notably, under pro-angiogenic stimulation by factors such as VEGF, FGF2, or high glucose, EC-secreted EXOs may contain different amounts of particular contents, which will further alter their angiogenic properties [62, 64]. In our study, we observed that HUVEC-derived EXOs alone slightly reduced EC proliferation and migration in vitro, but this effect was not observed in in vivo models. The EXOs that we used were derived from HUVECs cultured in normal endothelial growth medium without any stimulation; therefore, if we use VEGF- or pro-inflammatory factor-treated HUVECs, we may obtain different results as such cells could secrete EXOs with different angiogenic properties.

Although this study focused on retinal angiogenesis, it will also be interesting to test the efficacy of peptide-loaded EXOs in treating pathological angiogenesis in other disease models. It is worth noting that the wet form of age-related macular degeneration (wet AMD) characterized by choroidal neovascularization, is the leading cause of vision loss among elderly persons in developed countries [65]. The pathogenesis of wet AMD is due to the excessive production of VEGF from hypoxic cells in the retina-choroidal complex, which further leads to the growth and abnormal leakage of immature blood vessels. As our results showed that EXO_{KV11} could inhibit vascular leakage and limit the F4/80 infiltration in VEGF-treated retinas, further studies would be required to test whether EXO_{KV11} could also be the potential treatment for wet AMD. Angiogenesis is also essential for tumor growth and metastasis. During solid tumor progression, tumor vessels are also highly proliferative and functionally abnormal, with high permeability. These features facilitate intravasation and dissemination of tumor cells. As EXO_{KV11} suppressed EC proliferation and repaired the barrier leakiness, thus, it would be interesting to test whether EXO_{KV11} can also be used to suppress tumor angiogenesis and improve vascular integrity, as current anti-angiogenic therapy prolongs the survival of patients by only a limited time due to therapy escape or resistance.

In the OIR retina, two types of angiogenesis occurred after the pups were returned to ambient air: revascularization and neovascular tuft formation. Revascularization is a physiological process for repairing the injured retina in which normal blood vessels are formed in the ischemic area. In this study, we observed no difference in the avascular area, suggesting that revascularization is not improved by EXO_{KV11} treatment. This indicates that this system would not ameliorate ischemic disease. Further studies are required to search for a peptide that will yield benefits for revascularization. Notably, we have successfully loaded two peptides with distinctive functions onto EXOs in our previous work [30]. Thus, by using this ocular delivery system, it will be interesting to test whether we can achieve better therapeutic outcomes if we load more types of peptides with different biological functions.

In this study, we established a system that uses EXOs as carriers to deliver therapeutics into the ocular region for treating proliferative retinopathy. Our study highlights three unique advantages of EXO-based therapy: (i) the EXO therapeutic system is relatively stable with safety; (ii) EXOs exhibit high efficacy for treating pathological angiogenesis; and (iii) instead of intraocular injection, which is a commonly used method for ocular drug delivery, a less invasive route can be used for EXO-based therapy. Mechanistically, we demonstrated that EXO_{KV11} did not alter the anti-angiogenic property of KV11, but improved the delivery efficiency and achieved better therapeutic efficacy. A limitation of this study is that although our study clearly demonstrated the protective effect of EXO_{KV11} in OIR model and VEGF-induced retinal vascular leakage model, a long-term test could not be applied in such models. Further studies by using animal models, which are suitable for long-term treatment, would be more insightful. Furthermore, as this study used mouse as an experimental model, more extensive study will need to be performed prior to clinical deployment. Overall, our study provides a novel approach and has proved the feasibility of this approach for the future development of clinical translation for therapy.

**Methods Details**

**Animals**

All study protocols involving the use of animals were approved by the Institutional Animal Care and Use Committee of Tianjin Medical University. Adult wild type C57BL/6j and CD-1 mice, neonatal C57BL/6j mice with mothers were purchased from Model Animal Research Center of Nanjing University. For IVIS imaging and subsequent biodistribution testing, albino CD-1 mice were used. For Miles assay,
Sprague Dawley (SD) rats were used. For other experiments, C57BL/6j mice were used.

**Cell Culture**

Human Umbilical Vascular Endothelial Cells (HUVECs) were cultured as previously described [66] and were used from passage 2 to passage 6. For exosomes isolation, all cells were cultured in exosome-free normal growth medium (removed possible FBS-contained exosomes by centrifugation at 100,000×g for 16 h at 4 °C). For stimulation experiments, cells were starved overnight in growth factor free M199 medium (Gibco®, Life Technologies, US) supplemented with 2% FBS, 1% endothelial cell growth supplements and 1% antibiotic solution (P/S). All cells were used from passage 2 to passage 6.

**Exosome isolation**

Exosomes were isolated as previously described [67]. Briefly, cell culture medium was sequentially centrifuged at 1000×g for 10 min, followed by 10,000×g for 30 min. The supernatant was collected and filtered with a 0.22 µm filter (Millex, Germany), followed by ultracentrifugation at 100,000×g for 1 h. Exosome pellets were washed in a large volume of PBS and recovered by centrifugation at 100,000×g for 1 h and was resuspended in sterile PBS for the following experiments. The protein concentration of isolated exosomes was quantified by using the Bradford assay kit (Sangon Biotech, China).

**Exosome characterization**

To detect the expression of exosomes associated markers, exosomes pellets and cultured cells were lysed in RIPA lysis buffer containing 1X protease inhibitor. Proteins (50 µg for both exosome and cell lysate) were subjected to 12% SDS-polyacrylamide gel electrophoresis and transferred to a polyvinylidene difluoride membrane (Millipore, Germany). Specific antibodies used for detect exosome associated markers include Alix (3A9) (1:1000, 2171, Cell Signaling Technology, US), CD63 (1:200, sc-5275, Santa Cruz, US), CD81 (1:200, sc-166029, Santa Cruz, US), Cytochrome C (D18C7) (1:1000, 11940, Cell Signaling Technology, US) was used to determine mitochondria contamination.

The size distribution of HUVECs derived exosomes was measured by Nanosight NS300 (Malvern, UK). The morphology of exosomes was visualized by a high-resolution transmission electron microscope (TEM, Hitachi HT7700, Japan). The procedure of exosome staining prepared for TEM was performed as previously described [68].

**Peptides conjugation**

The KV11 sequence was YTMNPRKLFDYK; the KV11-CP05 sequence was YTMNPRKLFDYKCRHSQMTVSRL. KV11 and KV11-CP05 were synthesized as a single fusion peptide via peptide bond without spacer by ChinaPeptides (Shanghai, China) with 98% of purity.

**Peptide binding and cellular uptake in vitro**

To test the binding affinity of KV11-CP05 on EXO, 5 µg exosomes were preincubated with FITC-labeled KV11-CP05 (20 µg) in a 1.5 mL eppendorf tube under rotation at 4 °C for 6 h, followed by washing with PBS at 7500×g for 30 min for five times in 2 mL ultracentrifuge tubes and filtration with 100-kDa diafiltration tubes (Millipore, German) to remove unbound peptides. Final solution was collected by reversing the tubes and centrifugating at 1500×g for 30 min and analyzed with flow cytometry.

To test the cellular uptake of exosomes by HUVECs and HRMECs, isolated exosomes were labeled with DiR (Invitrogen, US) following the manufacturer’s instruction. 10 µg/mL labeled exosomes were incubated with HUVECs or HRMECs for 24 h in exosomes free medium. Cells were then washed with cold PBS and fixed for 30 min at RT with 4° PFA and nuclei were counterstained with DAPI (Invitrogen, US). DiR and DAPI were imaged on a confocal microscope (LSM800, Carl Zeiss, Germany).

To determine the delivery efficiency of EXO_KV11 in HUVECs and HRMECs, DiR-labeled exosomes (5 µg) were incubated with FITC-labeled KV11-CP05 (20 µg) or FITC-labeled KV11 (20 µg) at 4 °C for 6 h. Subsequently, peptides/exosomes mixture or peptide-exosome complexes were incubated with HUVECs or HRMECs in 24-well plate for 24 h or 48 h. Cells were washed with cold PBS and fixed for 30 min at RT with 4° PFA and nuclei were counterstained with DAPI. Images were obtained by confocal microscope. To compare peptide delivery efficiency in HUVECs and HRMECs, cells of each group were harvested and was analyzed with flow cytometry (FACSVerso, BD, US).

**Ocular distribution**

FITC-labeled KV11 peptide and KV11-CP05 were synthesized by ChinaPeptides (Shanghai, China). To examine the ocular distribution of peptide, KV11 or EXO_KV11 were administered retro-orbitally or intravitreally into CD1 mice. 50 µg FITC-KV11 and EXO_KV11 (50 µg FITC-KV11-CP05 and 25 µg EXO were incubated for 6 h at 4 °C before injection) were dissolved in saline solution (0.9% NaCl) and injected
retro-orbitally. 2 μg FITC-KV11 and EXO_{KV11} (2 μg KV11-CF05 and 1 μg EXO were incubated for 6 h at 4 °C before injection) in saline solution (0.9% NaCl) and injected intravitreally. After indicated time points (6 h, 12 h for retro-orbital injection or intravitreal injection), the mice were perfused with cold PBS to wash out free peptides and exosomes in the circulation. Eyes were then harvested for imaging with IVIS spectrum (PE, Waltham, MA, US). The fluorescence intensity was measured by using the ROI tool of Living Image software (V4.2). Briefly, the region of the eyes was selected by using the ROI tool and the fluorescence intensity was measured. After subtracting the background fluorescence, the data were applied for analysis.

To detect the distribution of FITC-labeled peptides in retinal sections, eyes from each group were enucleated in PBS after mice were sacrificed. The eyes were fixed in 4% PFA for 1 h at 4 °C. Afterwards, they were transferred to 30% sucrose/PBS at 4 °C overnight and subsequently embedded in optimal cutting temperature compound (OCT) (Sakura, Japan) and frozen at -80 °C. Serial 15 μm-thick sections were cut using a cryostat (CM1950, Leica, Germany). Cryosections were washed in PBS for 10 min, permeabilized in 0.3% TritonX-100 PBS for 15 min and then blocked in 2% BSA, 0.3% TritonX-100 PBS for 1 h. Cryosections were then incubated with IsolectinGS-IB4 (Alexa Fluor 568-conjugated, 1:200, I21413, Invitrogen, US) for 2 h, that was used for labeling the blood vessels. Images were collected on a confocal microscope (LSM800, Carl Zeiss, Germany). To quantify the abundance of FITC labeled peptides delivered to the blood vessels, the fluorescence intensity of FITC was analyzed in isolectinB4 positive area by using image J software.

**Oxygen induced retinopathy (OIR) model**

Oxygen induced retinopathy model was performed as previously described [54]. Briefly, neonatal C57BL/6j mice with the nursing mother at day P7 were exposed to hyperoxia (75% O2) for 5 days. Mice returned to room air at P12. For applying treatment, 25 μg EXO, 50 μg KV11 and EXO_{KV11} (50 μg KV11-CF05 and 25 μg EXO were incubated for 6 h at 4 °C before injection) were dissolved in saline solution (0.9% NaCl) and injected retro-orbitally at P12. The retinas were collected at P17 for analysis.

**VEGF induced retina vascular leakage model**

In pre-treatment model, adult wild type C57BL/6j mice were pretreated retro-orbitally with 50 μg KV11, 25 μg EXOs or EXO_{KV11} (50 μg KV11-CF05 and 25 μg EXO were incubated for 6 h at 4 °C before injection) in 50 μL saline solution (0.9% NaCl). In VEGF-trap treated group, 15 μg VEGF-trap (10 mg/mL Conbercept Ophthalmic Injection, KANGHONG, China) was pretreated intravitreally by a 33-gauge beveled needle. 1 μL VEGF164 protein (100 ng/μL) was injected intravitreally 24 h after pretreatment using a 33-gauge beveled needle. Mice were sacrificed and retinas were collected for analysis after another 24 h.

In post-treatment model, 1 μL VEGF164 protein (100 ng/μL) was injected intravitreally. After 24 h, mice were treated retro-orbitally with 50 μg KV11, 25 μg EXO or EXO_{KV11} (50 μg KV11-CF05 and 25 μg EXO were incubated for 6 h at 4 °C before injection) in 50 μL saline solution (0.9% NaCl). After 48 h post-treatment, the retinas were collected for analysis.

**Immunofluorescence**

For flat-mounted retinas, eyes were enucleated from mice and fixed in 4% PFA for 1 h at 4 °C. Retinas were dissected, washed with PBS and permeabilized with PBS containing 1% TritonX-100 overnight at 4°C then blocked in PBS containing 2% BSA, 0.3% TritonX-100 for 12 h at 4 °C. After blocking, for visualization of retinal vasculature in OIR models, flat-mounted retinas were stained with isolectinGS-IB4 (1:100) for 2 h at RT. For IF staining of red blood cells, blood vessels, macrophage and microglia, flat-mounted retinas were incubated in blocking solution with following primary antibodies at 4 °C overnight: rat anti-TER119 monoclonal (1:200, MAB1125, RD, US); rat anti-CD105 monoclonal (1:200, MAB1320, RD, US); rabbit anti-F4/80 monoclonal (1:200, 30325S, Cell Signaling Technology, US); mouse anti-Claudin-5 monoclonal (1:200, 35-2500, Invitrogen, US) and rabbit anti-ZO-1 polyclonal (1:300, 61-7300, Invitrogen, US). After washing, the retinas were incubated with corresponding secondary antibodies (1:300, Jackson Immunoresearch, US) for 2 h at RT. Flat-mounted retinas were analyzed using a confocal fluorescence microscope (LSM 800, Carl Zeiss, Germany).

**Analyses of vascular leakage and perfusion**

Vascular leakage was analyzed by intracardial injection performed as previously described [69]. In brief, 31-G needle was positioned above the heart 2 mm parasternal to the left at a virtual line connecting both armpits, in caudal and lateral angles of 30° and 10°, 50 μL, warm PBS containing FITC-conjugated dextran (25 mg/mL, 70 kDa, FD705, Sigma-Aldrich, US) were injected. After 10 min of circulation, eyes were enucleated and fixed in 4% PFA for 1 h at 4 °C. In OIR model, retinas were then disected, washed with PBS and permeabilized with PBS containing 1% TritonX-100 overnight at 4 °C and stained with
isolateG5S-IB4 (1:100) for 2 h at RT, and flat-mounted in fluorescent mounting medium (Fluoromount-G®, 0100-01, SouthernBiotech, US) after several washes. In VEGF induced retina vascular leakage model, retinas were incubated in blocking solution with the primary antibody rat anti-CD105 monoclonal (1:200, MAB1320, RD, US) at 4 °C overnight after permeabilization and blocking. Then corresponding secondary antibody was used for 2 h at RT. Leakage was observed and imaged on a confocal fluorescence microscope (LSM800, Carl Zeiss, Germany). FITC-conjugated dextran was measured as dextran area outside the vessels divided by total measured area of retina by using Image J software.

**Morphometric analyses**

Morphometric analyses of the retinas were performed using Image J software and Adobe Photoshop software. To determine the amount of regression and neovascularization, the number of pixels in the avascular regression area and neovascular area was measured using the Lasso tool of Adobe Photoshop software as described [54], and divided by the number of pixels in the total retinal area and presented as a percentage. RBC leakage was measured as area outside the vessels divided by total measured area of retina as previously described [70]. Macrophage infiltration was measured as the total number of F4/80+ macrophage in the whole retina.

**EdU injection in pups**

EdU (E6032, US EVERBRIGHT INC., China) was injected at a concentration of 50 μg/g body weight into pups at P17 in OIR model 2.5 h before sacrifice. Eyes were enucleated from pups and fixed in 4% PFA for 1 h at 4 °C, then were transferred to 30% sucrose/PBS at 4 °C overnight and subsequently embedded in optimal cutting temperature compound (OCT) (Sakura, Japan) and frozen at -80 °C. Afterwards, sagittal cryosections of the eyes were prepared. EdU+ cells in the cryosections of retinas were detected by using YF® 488 Click-iT EdU Stain Kits (C6033, US EVERBRIGHT INC., China), according to the manufacturer’s instructions. ECs were counterstained with ERG (1:400, ab92513, Abcam, UK) overnight at 4 °C, isolateG5S-IB4 (1:200) and corresponding secondary antibody (1:300, Jackson ImmunoResearch, US) for 2 h at RT. The numbers of EdU+ERG+ cells were counted from five sagittal eye sections. To ensure that similar areas were used for doing quantification for each eye, we always use sections intersect the optic nerve area to maintain consistency.

**RNA extraction and quantitative real-time PCR analysis**

Eyes were enucleated from pups at P17 in OIR model. Retinas were dissected and homogenized in TRIzol® reagent (Invitrogen, US). RNA samples were reverse-transcribed to complementary DNA (cDNA) using TransScript One-Step gDNA Removal and cDNA Synthesis SuperMix (TransGen, China). qPCR was performed using PerfectStart™ Green qPCR SuperMix (TransGen, China) and was processed with QuantStudio 5 Real-Time PCR system (Applied Biosystems, US). β-actin was used as internal control. All qPCR results were obtained from at least 5 biological repeats. The primers used in this study were: Hif1a Forward, 5'-ACCTTCATCGGAAAC TCCAAAG-3'; Hif1a reverse, 5'-CTGTTAGGCTGG GAAAAGTTAGG-3'; Vegfa forward, 5'-GCACAT AGAGAGAATGAGCITCC-3'; Vegfa reverse, 5'-CTC CGCTCTGAAAGGCT-3'; Il6 forward, 5'-TAG TCCCTCCTACCCAATTTCC-3'; Il6 reverse, 5'-TTG GTCTTACTGCCACTCTTC-3'. Vcam1 forward, 5'-AGTG GGGATTTCGTTTCT-3'; Vcam1 reverse, 5'- CGCTGTATTTCCCTCATCG-3'; Actb forward, 5'- GGCTGTATTTCCCTCATCG-3'; Actb reverse, 5'- CCAGTTGGTAAATGCCATGT-3'.

**In vitro BrdU incorporation assay**

**In vitro** BrdU incorporation was performed as described [71]. Briefly, 2 × 10⁴ HUVECs were cultured in collagen coated coverslips in 24-well plates and pretreated with EXO, KV11 or EX0KV11 (KV11-CP05 and EXO were incubated for 6 h at 4 °C before treatment) at concentration of 12.5 μg/mL EXO and 25 μg/mL peptide for 24 h in starvation medium. Afterwards, cells were treated with or without VEGF (50 ng/mL) for 24 h. BrdU (10 μM, B5002, Sigma-Aldrich, US) was subsequently added and incubated for 4 h at 37 °C. Cells were fixed in 4% PFA/PBS for 20 min and permeabilized and blocked in PBS containing 2% BSA, 0.3% TritonX-100 for 30 min at RT. Nuclei were counterstained with DAPI (1:1000, D1306, Invitrogen, US). Images were obtained using a fluorescence microscope (BX51, Olympus, Japan). N > 8 biological repeats were analyzed. Quantification was done blind to the experimental condition.

**In vivo** BrdU incorporation was performed as described [71]. Briefly, 2 × 10⁴ HUVECs were cultured in collagen coated coverslips in 24-well plates and pretreated with EXO, KV11 or EX0KV11 (KV11-CP05 and EXO were incubated for 6 h at 4 °C before treatment) at concentration of 12.5 μg/mL EXO and 25 μg/mL peptide for 24 h in starvation medium. Afterwards, cells were treated with or without VEGF (50 ng/mL) for 24 h. BrdU (10 μM, B5002, Sigma-Aldrich, US) was subsequently added and incubated for 4 h at 37 °C. Cells were fixed in 4% PFA/PBS for 20 min and permeabilized and blocked in PBS containing 2% BSA, 0.3% TritonX-100 for 30 min at RT. Nuclei were counterstained with DAPI (1:1000, D1306, Invitrogen, US). Images were obtained using a fluorescence microscope (BX51, Olympus, Japan). N > 8 biological repeats were analyzed. Quantification was done blind to the experimental condition.
Scratch assay

To analyze the EC migration, 2×10⁵ HUVECs were cultured in collagen coated 6-well plates and pretreated with EXO, KV11 or EXOₓ₁₁ (KV11-CP₀⁵ and EXO were incubated for 6 h at 4 °C before treatment) at concentration of 12.5 µg/mL EXO and 25 µg/mL peptide for 24 h in starvation medium. A wound was made by scraping the cell monolayer with a 10 µL pipette tip, and cells were stimulated with or without VEGF (50 ng/mL). Pictures were acquired at time-point zero and 12 h after incubation at 37 °C. The percentage of wound closure between 0 h and 12 h was analyzed with Image J software. N = 8 biological repeats were quantified.

Fibrin gel bead sprouting assay

Fibrin gel bead sprouting assay was performed as previously described [2]. Briefly 1×10⁵ HUVECs were seeded in collagen coated 6-well plates and pretreated with EXO, KV11 or EXOₓ₁₁ (KV11-CP₀⁵ and EXO were incubated for 6 h at 4 °C before treatment) at concentration of 12.5 µg/mL EXO and 25 µg/mL peptide for 24 h in starvation medium. Cytodex 3 microcarrier beads (GE Healthcare) were coated with pretreated HUVECs (mixed at 200 cells per bead), and embedded in 2 mg/mL fibrin gels in 24-well plates by mixing 2 mg/mL fibrinogen (Sigma-Aldrich, US) in PBS, 0.625 units/mL thrombin (Sigma-Aldrich, US) and 0.15 units/mL aprotinin (Sigma-Aldrich, US). Beads were cultured in starvation medium in the presence or absence of 50 ng/mL VEGF. After 24 h, the culture beads were fixed with 4% PFA for 15 min at RT. Images were taken with an inverted fluorescence microscope (Ti2-U, Nikon, Japan) and sprout number and length were analyzed by Image J software. Approximately 20 beads per condition were quantified.

Auricular miles assay

Auricular miles assay was performed as previously described [72]. Briefly, Evans Blue dye (45 mg/kg) was injected intravenously. After 10 min, rats were anesthetized and intradermally injected with 200 ng recombinant VEGF in 20 µL PBS (vehicle) in the left ear, and with recombinant VEGF and KV11(12.5 µg) or EXOₓ₁₁ (12.5 µg KV11-CP₀⁵ and 6.25 µg EXO were incubated for 6 h at 4 °C before treatment) in 20 µL PBS in the contralateral ear. Evans blue extravasation was photographed 3 h after EB injection.

VEGF downstream signaling analysis

To detect the VEGF downstream signaling, 2×10⁵ HUVECs were pretreated with EXO, KV11 or EXOₓ₁₁ (KV11-CP₀⁵ and EXO were incubated for 6 h at 4 °C before treatment) at concentration of 12.5 µg/mL EXO and 25 µg/mL peptide for 24 h in starvation medium. In VEGF-trap (conbercept) treated wells, cells were pretreated with 10 µg conbercept for 5 min before VEGF stimulation. HUVECs were then stimulated with VEGF (25 ng/mL, R&D systems, US). Cells were washed with ice cold PBS and lysed in cold RIPA lysis buffer supplemented with 1X protease inhibitor cocktail and 1X phosphatase inhibitor (Roche, Switzerland). Proteins (30 µg for each sample) were subjected to 10% SDS-polyacrylamide gel electrophoresis and transferred to a polyvinylidene difluoride membrane (Millipore, Germany). Target proteins were detected by specific primary antibodies including Src (36D10) (1:1000, 2109, Cell Signaling Technology, US), Src-Phospho (Y416) (1:1000, AP0452, ABclonal, China), Erk1/2 p44/42 (1:1000, 9102, Cell Signaling Technology, US), Erk1/2 phospho-p44/42 (T202/Y204) (1:1000, 9106, Cell Signaling Technology, US), β-actin (1:1000, 3700, Cell Signaling Technology, US) or β-tubulin (1:1000, UM4003, Utibody, China)-were used as loading control.

The bound primary antibody was detected by horseradish peroxidase-conjugated goat anti-mouse IgG (1:3000, 7076, Cell Signaling Technology, US) and goat anti-rabbit IgG (1:3000, 7074, Cell Signaling Technology, US), respectively. The ECL western blotting analysis system (Millipore, US) was applied. Quantification of WB signal was done by Image J software. Proteins expression values were divided by their β-actin or β-tubulin, respectively. Each experiment was performed at least three times.

Toxicity assessment

Toxicity assessment was performed at P17 in OIR model and endpoint in pre-treated VEGF induced vascular leakage model.

Terminal deoxynucleotidyl transferase mediated dUTP nick end labeling (TUNEL) assay was used to examine apoptotic cells in retinas. Mice were sacrificed after indicated endpoint in each model and eyes from each group were enucleated in PBS. The eyes were fixed in 4% PFA for 1 h at 4 °C. Afterwards, they were transferred to 30% sucrose/PBS at 4 °C overnight and subsequently in OCT (Sakura, Japan) and frozen at -80 °C. Serial 15 µm-thick sections were cut using a cryostat (CM1950, Leica, Germany). TUNEL assay was performed using a FragEL™ DNA Fragmentation Detection Kit (QIA39, Millipore, US) according to the manufacturer’s instructions.

Routine H&E staining was used to examine the overall major organs morphology, including liver, spleen, lung and kidney from mice in OIR model and VEGF induced retina vascular leakage model. Organs were fixed with 4% formaldehyde overnight at RT,
embedded in paraffin, and sectioned to 6-μm thickness. Images were obtained using a microscope (Ti2-U, Nikon, Japan).

To analyze the serum biochemistry, mouse blood was taken immediately after sacrifice and centrifuged at 1,500×g for 30 min and stored at -80 °C. Analysis of alanine aminotransferase (ALT) and aspartate aminotransferase (AST) were performed using AST/GOT KIT and ALT/GPT KIT (BioSino, China) and read by Microlab 300 (Vital, Holland).

**Statistical analysis**

Results were expressed as the mean ± SEM. To calculate statistical significance, the Mann-Whitney test or student’s t test (two-tailed) (when comparing two groups) or one-way ANOVA followed by Tukey’s multiple comparisons test (when comparing three or more groups) were used. P value < 0.05 were considered significant. All calculations were performed using Prism software.

**Supplementary Material**

Supplementary figures. http://www.thno.org/v11p5107s1.pdf

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**Author Contributions**

X.D., X.W. and H.Y. designed the project; X.D., Y.L., Z.Y., T.W., Y.L., G.H., X.Z., Y.L., Y.S. carried out the experiments; X.D., Y.L. and Z.Y. analyzed the data; H.Y., X.W. and H.Y. supervised the study; H.X., Y.L., Z.Y., T.W., Y.L., G.H., X.Z., Y.L., Y.S. carried out the experiments; X.D., Y.L., X.W. and H.Y. wrote the paper with the input from all authors.

**Competing Interests**

The authors have declared that no competing interest exists.

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