Slc39a5-mediated zinc homeostasis plays an essential role in venous angiogenesis in zebrafish

Zhidan Xia, Xinying Bi, Jia Lian, Wei Dai, Xuyan He, Lu Zhao, Junxia Min and Fudi Wang

The First Affiliated Hospital, School of Public Health, Institute of Translational Medicine, Zhejiang University School of Medicine, Hangzhou, People’s Republic of China

Angiogenesis is a precise process mediated by a variety of signals and the environmental niche. Although the essential trace element zinc and its homeostasis are essential for maintaining proper cellular functions, whether zinc plays a role in angiogenesis is currently unknown. Using zebrafish embryos as a model system, we found that zinc treatment significantly increased the expression of the slc39a5 gene, which encodes the zinc transporter Slc39a5. Moreover, knocking down slc39a5 expression using either a morpholino or CRISPR/Cas9-mediated gene editing led to cardiac ischaemia and an accumulation of red blood cells in the caudal vein plexus (CVP), as well as delayed venous sprouting and fewer vascular loops in the CVP region during early development. Further analysis revealed significantly reduced proliferation and delayed cell migration in the caudal vein of slc39a5 morphants. At the mechanistic level, we found increased levels of systemic zinc in slc39a5-deficient embryos, and chelating zinc restored CVP development. In addition, we found that zinc overload in wild-type embryos leads to impaired CVP formation. Taken together, these results indicate that Slc39a5 plays a critical role in endothelial sprouting and migration in venous angiogenesis by regulating zinc homeostasis.

1. Introduction

Angiogenesis is an essential biological process in which new blood vessels emanate from existing vascular structures, with vascular budding, migration, proliferation and pruning, thereby establishing the vascular network [1,2]. In zebrafish, the dorsal aorta and axial vein form a primitive circulatory loop, and angiogenesis from these vessels plays a key role in generating complex vascular networks. Specifically, the posterior axial vein extends ventrally at 26–32 hours post-fertilization (hpf), to eventually form a ‘honeycomb-like’ network at 38 hpf; this network, called the caudal vein plexus (CVP), is composed of a dorsal vein and a ventral vein, with interconnected vessels [3].

Angiogenesis is necessary for the efficient delivery of oxygen and plays a critical role in a wide range of physiological and pathophysiological processes, including tissue development, nutrient transport, wound healing and tumour neovascularization [3–5]. However, the underlying cues and mechanisms that initiate and maintain angiogenesis are poorly understood; moreover, the role of zinc and zinc homeostasis in angiogenesis has not been investigated.

As an essential trace element, zinc is important for maintaining cellular physiology and for the development of various organs and tissues [6–8]. At the cellular level, zinc homeostasis is controlled by two families of transporter proteins, including 14 members of the SLC39A (Zrt- and Irt-like protein, ZIP) family and 10 members of the SLC30A (zinc efflux transporter, ZnT) family [9,10]. Interestingly, the zinc transporter SLC39A12 was reported to serve as a...
major regulator of pulmonary vascular remodelling under low-oxygen conditions, as genetically deleting Slc39a12 expression protected against the development of pulmonary hypertension in rats housed in a hypoxic atmosphere [11]. However, whether other members of the SLC39A family play role in vascular development is an open question.

The zinc transporter SLC39A5, mediates the transport of metal ions across cell membranes, including the plasma membrane and subcellular organelles [12]. In intestinal enterocytes, SLC39A5 is expressed in the basolateral membrane and may play an important role in communicating the body’s zinc status to enterocytes in order to control systemic zinc homeostasis [13,14]. In addition, SLC39A5 has been reported to play important roles under both physiological and pathological conditions. For example, β-cell-specific Sla39a5 knockout mice have impaired insulin secretion and reduced glucose tolerance [15]. In oesophageal squamous cell carcinoma, SLC39A5 is highly upregulated and may play a role in cellular proliferation and migration, promoting disease progression [16]. In addition, SLC39A5 is expressed in high levels in the sclera and retina, and mutations in SLC39A5 have been linked to autosomal dominant non-syndromic high myopia in humans [17]. Nevertheless, whether SLC39A5 and/or SLC39A5-mediated zinc homeostasis plays a role in vascular development or angiogenesis is currently unknown.

Here, we report that SLC39A5 plays an important role in angiogenesis during early development in zebrafish. Specifically, we found that maintaining systemic zinc concentrations is essential for the proliferation and migration of endothelial cells during venous angiogenesis. These results reveal a new function for SLC39A5 in early vertebrate development with respect to zinc homeostasis during angiogenesis, providing important new insights into the signals and mechanisms that underlie this process and suggesting a potential new therapeutic target for treating angiogenesis-related diseases.

2. Results

2.1. Knocking down slc39a5 leads to blood accumulation in the zebrafish caudal vein plexus

To determine whether members of the SLC39A family play a role in regulating systemic zinc homeostasis in zebrafish, we exposed wild-type (WT) embryos to 1 mM zinc and measured the expression of slc39a1 through slc39a14. We found that zinc treatment significantly and selectively increased the expression of slc39a5 (figure 1a). Using whole-mount in situ hybridization, we also found that slc39a5 is expressed uniformly through WT zebrafish embryos at 1–2 days post-fertilization (dpf), and its expression becomes concentrated at the head and gut region by 3 dpf (figure 1b).

To analyse the functional role of Slc39a5 in zebrafish, we knocked down slc39a5 expression using an ATG-based morpholino to prevent translation. We found that slc39a5 morphants develop normally until 48 hpf, when they develop cardiac ischaemia (figure 1c). By 72 hpf, slc39a5 morphants have severely reduced blood flow in the heart compared with control embryos (figure 1c). Using o-Dianisidine to stain haem, we confirmed the presence of cardiac ischaemia in slc39a5 morphants at 48–72 hpf (figure 1d).

To test whether the cardiac ischaemia in slc39a5 morphants was due to impaired cardiovascular function, we measured heart rate and cardiac morphology in control and morphant embryos. We found no significant difference between control and morphant embryos with respect to either heart rate (electronic supplementary material, figure S1a) or cardiac morphology (electronic supplementary material, figure S1b). To determine whether the slc39a5 morphants develop anaemia, we measured red blood cells (RBCs) in the transgenic zebrafish line Tg(globinLCR:eGFP), in which RBCs are fluorescent [18]. We found no significant difference in the number of RBCs between control and morphant embryos measured at 26–72 hpf (electronic supplementary material, figure S1c–e). Interestingly, however, we found that RBCs accumulated in the CVP of morphants but not in control embryos (figure 1e). This result was confirmed by performing in situ hybridization of hhba3, a marker for RBCs (figure 1f). Taken together, these results show that knocking down slc39a5 in zebrafish embryos causes RBCs to accumulate in the CVP and reduces cardiac blood flow, resulting in cardiac ischaemia.

2.2. Slc39a5 morphants have delayed angiogenesis in the caudal vein plexus

Next, we examined the vascular structure and blood distribution in the CVP using a dual-fluorescent zebrafish expressing the endothelial cell marker kdr:mCherry [19] and globinLCR:eGFP. Compared with control embryos, slc39a5 morphants had significantly fewer filopodia sprouting from exiting endothelial cells at 32 hpf (figure 2a,b), the age at which the caudal vasculature begins budding in the CVP [3]. Similarly, the slc39a5 morphants had significantly fewer capillary loops compared with control embryos at 32 hpf (figure 2a,c). At 38 hpf, when the honeycomb-like network finally forms in the CVP of control embryos [3], the vascular plexus were considerably less developed in slc39a5 morphants, with significantly fewer capillary loops (figure 2d–f). In addition, we found an increased number of RBCs in the caudal veins of slc39a5 morphants compared with control embryos (figure 2d, green fluorescence), suggesting that the accumulation of RBCs in the CVP of morphant embryos is associated with the reduced development of caudal veins. At 42 hpf, control embryos no longer had CVP sprouts, whereas a significantly higher number of sprouts were present in the caudal veins of slc39a5 morphants (figure 2g–i), indicating delayed venous angiogenesis in the CVP region.

To determine whether impaired venous angiogenesis in slc39a5 morphants is tissue-specific, we analysed the development of the sub-intestinal vein plexus (SIVP), which sprouts from the ventral axial vein and mediates the absorption of nutrients from the yolk [20]. Similar to angiogenesis in the CVP region, we found that endothelial sprouting was significantly delayed in the SIVP in slc39a5 morphants (electronic supplementary material, figure S2a–e), suggesting that the delayed venous angiogenesis in slc39a5 morphants is a general phenomenon.

To confirm that the effects of the slc39a5 morpholino were due specifically to knocking down slc39a5, we performed a rescue experiment in which we co-injected single-cell embryos with both the slc39a5 morpholino and an ATG-MO-insensitive slc39a5 mRNA containing a 5-nucleotide mismatch at the 5′ end; for these experiments, we used the transgenic Tg(fli1a: eGFP) zebrafish line in which endothelial cells express eGFP [4]. We found that expressing the ATG-MO-insensitive slc39a5 mRNA restored the vascular structure (figure 2j),
preventing both the reduction in endothelial sprout formation (figure 2k) and reduced capillary loop formation (figure 2l).

**2.3. Reduced caudal vein plexus angiogenesis in slc39a5 morphants is due to impaired proliferation and migration of endothelial cells**

During angiogenesis, sprout elongation is driven primarily by the proliferation and migration of endothelial cells [21]. Given the reduction in endothelial sprouting in slc39a5 morphants, we measured endothelial cell proliferation in the CVP region using phosphohistone H3 (PH3) and BrdU immunostaining. We found that both PH3 and BrdU staining were significantly reduced in the caudal veins in slc39a5 morphants (figure 3a–c), consistent with reduced cell proliferation in the CVP.

Next, we monitored the migration of endothelial cells in the CVP region from 32 hpf through 35 hpf using high-resolution spinning disk confocal microscopy. In control embryos, two venous endothelial cells sprouted at 33 hpf and gradually

---

**Figure 1.** slc39a5 morphant zebrafish embryos develop cardiac ischaemia and blood cell accumulation in the caudal vein plexus. (a) Quantitative PCR analysis of the 12 indicated slc39a genes measured in wild-type (WT) embryos and embryos treated with 1 mM zinc (+Zn). Note zebrafish lacking slc39a2 and slc39a12. (b) In situ hybridization of WT zebrafish embryos using the antisense and sense (control) slc39a5 probes. Note the concentrated expression in the head and gut region at 3 dpf (red arrow). (c–d) Representative images (c) and o-Dianisidine-stained (d) control and slc39a5 morphant embryos at the indicated developmental stages. The red arrows indicate the heart region. (e) Representative images of GFP fluorescence measured in the caudal region of control and slc39a5 morphant Tg(globinLCR:eGFP) embryos. Note the accumulation of RBCs in the CVP of slc39a5 morphants. (f) Whole-mount in situ hybridization of the RBC marker hbae3 in control and slc39a5 morphants. Note the increased signal in the CVP of the slc39a5 morphant (arrow).
both mutant lines (figure 4a). Using dual-fluorescence microscopy at 32 hpf (b–c) and 42 hpf (h–i), (j–l) Representative images of the CVP in control Tg(fli1a:eGFP) embryos and Tg(fli1a:eGFP) embryos injected with the slc39a5 morpholino alone or co-injected with the slc39a5 morpholino and a morpholino-resistant slc39a5 mRNA. The numbers of endothelial sprouts (k) and capillary loops (l) in the CVP were measured at 32 hpf. *p < 0.05, ***p < 0.001 and n.s., not significant.

We also measured cell death using TUNEL staining and analysed DNA damage by measuring the accumulation of H2AX (H2A histone family member X) in embryos during angiogenesis. We found that neither cell death nor DNA damage was significantly different between the control embryos and the slc39a5 morphants (figure 3f–h).

2.4. Knocking out slc39a5 in zebrafish produces a similar phenotype, including impaired angiogenesis

To confirm the role of Slc39a5 in venous angiogenesis using an independent approach, we generated slc39a5 knockout mutants using CRISPR/Cas9 and obtained two separate lines with specific genetic mutations in the target gene; both mutations introduce a frame-shift, resulting in a premature stop codon (figure 4a). Quantitative real-time PCR confirmed that slc39a5 expression is virtually abolished in both mutant lines (figure 4b). Consistent with our knockdown strategy, both mutant lines develop a phenotype similar to slc39a5 morphants, with reduced blood flow in the heart (figure 4c), impaired angiogenesis in the CVP, delayed endothelial sprouting and fewer capillary vascular loops (figure 4d–g) compared with control embryos. Also consistent with our previous results, endothelial cell proliferation was significantly lower in mutant embryos compared with controls (figure 4h,i), while cell death was unaffected (electronic supplementary material, figure S3). Thus, both knocking down and knocking out slc39a5 revealed that it plays a central role in venous angiogenesis.

2.5. Slc39a5-mediated zinc homeostasis regulates angiogenesis in the caudal vein plexus

Given that Slc39a5 is a putative zinc transporter that regulates systemic zinc concentrations [13,14], we used a fluorescent zinc probe to measure the effect of deleting Slc39a5 on systemic zinc levels in zebrafish embryos. We found that compared with control embryos, both the slc39a5 mutants and the slc39a5 knockout embryos had higher levels of zinc (figure 5a,b), consistent with zinc accumulation in absence of Slc39a5. Next, we examined the role of zinc homeostasis in venous angiogenesis by treating control embryos and slc39a5 morphants with a high zinc solution (to increase systemic zinc levels) and/or the cell-permeable zinc chelator TPEN.
We found that zinc-treated control embryos had reduced filopodia sprouting and fewer capillary loops (figure 5c–d,i,j), mimicking the phenotype observed in slc39a5 morphants in which zinc was accumulated; moreover, these effects were absent in control embryos co-treated with both zinc and TPEN (figure 5c,e,i,j). Finally, treating slc39a5 morphants with TPEN restored angiogenesis, with increased filopodia sprouting and capillary loop formation (figure 5f,g,i,j), while co-treatment with TPEN and zinc neutralized the ability of TPEN to rescue angiogenesis in slc39a5 morphants (figure 5f, h,i,j). Taken together, these results indicate that loss of Slc39a5 in zebrafish embryos disrupts zinc homeostasis and impairs venous angiogenesis.

3. Discussion

Angiogenesis is orchestrated by a close interaction between endothelial cells and their niche [22]. Although a variety of signals and processes have been investigated with respect to their role in angiogenesis, how zinc and zinc homeostasis play a role in angiogenesis is poorly understood. Here, we report that the zinc transporter Slc39a5 plays an essential role in venous angiogenesis in zebrafish, and we report that zinc homeostasis is required for endothelial budding and the proliferation and migration of endothelial cells.

Our in vivo results show that slc39a5 is widely expressed in zebrafish embryos during early development, subsequently becoming concentrated in the head and gut region. Previous studies showed that SLC39A5 localizes to the basolateral surface of enterocytes in mice fed a zinc adequate diet, suggesting that this protein may remove zinc from the blood via the intestine [13,14]. These reports support our finding that knocking down slc39a5 leads to systemic zinc accumulation in zebrafish embryos, indicating that Slc39a5 plays an essential role in maintaining systemic zinc homeostasis and functions to remove excess zinc from the body.

Our finding that slc39a5 morphant embryos develop a clear phenotype that includes cardiac ischaemia and the accumulation of blood in the CVP suggests the possibility of toxicity and/or an off-target effect of the morpholino [23,24]. We believe these effects are unlikely, however, as

---

**Figure 3.** slc39a5 morphants have impaired endothelial cell proliferation and migration in the CVP region. (a–b) PH3 (a) and BrdU (b) immunostaining of control and slc39a5 morphant Tg(fli1a:eGFP) embryos, with magnified views of the CVP showing co-localization of GFP fluorescence and PH3 immunoactivity (arrows). (c) Summary of BrdU-positive cells in the CVP of control and slc39a5 morphants. ***p < 0.001. (d) Representative sequential images of the CVP in control and slc39a5 morphants. Tg(fli1a:eGFP) embryos. For each embryo, two separate endothelial cells are indicated with a white asterisk and a yellow asterisk. Note that 0 h corresponds to 32 hpf. (e) Summary of the migration distance of endothelial cells from 32 to 35 hpf in the CVP of control and slc39a5 morphant Tg(fli1a:eGFP) embryos, with magnified views of the CVP showing co-localization of GFP fluorescence and PH3 immunoreactivity (arrows). (f) Representative images of TUNEL staining in the CVP of control and slc39a5 morphant Tg(fli1a:eGFP) embryos. (g–h) Representative images (g) and summary (h) of H2AX immunostaining in the CVP of control and slc39a5 morphant Tg(fli1a:eGFP) embryos.
co-injecting single-cell embryos with a morpholino-resistant slc39a5 transcript together with the slc39a5 morpholino restored normal development, thereby preventing the phenotype induced by the slc39a5 morpholino. In addition, we generated two separate slc39a5 knockout lines using CRISPR/Cas9 and found that the resulting phenotype is similar to the slc39a5 morphants, confirming that both cardiac ischaemia and blood accumulation in the CVP are due to the loss of Slc39a5.

In the condition of slc39a5 deficiency, although the endothelial sprouting and migration in CVP and SIVP were delayed, the intersegmental vessel (ISV) showed a relative normal development (figure 2j; electronic supplementary material, figure S4e). As CVP and SIVP are formed by angiogenic sprouts from the axial vein, while ISV is formed by angiogenic sprouts from the dorsal aorta [3], we speculate that Slc39a5-mediated zinc homeostasis affects venous angiogenesis in zebrafish.

Since the CVP overlaps with caudal haematopoietic tissue (CHT), an important tissue for definitive haematopoiesis [25,26], we evaluated the definitive haematopoiesis development by analysis the expression of cmyb, a well-known marker gene for definitive haematopoiesis in zebrafish [27,28]. By using a transgenic fish line Tg(cmyb:eGFP), we found that the

---

**Figure 4.** Targeting strategy and characterization of slc39a5 knockout zebrafish. (a) DNA and corresponding amino acid sequences of the wild-type (WT) slc39a5 allele and the slc39a5 allele after inserting one nucleotide (+1) or deleting one nucleotide (−1) using CRISPR/Cas9-based editing. Both the insertion and the deletion introduce a premature stop codon. (b) Summary of slc39a5 mRNA measured using qPRC in the WT, +1, and −1 slc39a5 mutant lines (n = 3 sets of 50 pooled embryos/group). (c) Representative images of a wild-type and mutant embryo at 3 dpf. Note the significantly smaller heart with reduced cardiac blood flow in the mutant embryo (arrow). (d–e) Representative images of the CVP in wild-type (left) and mutant (right) Tg(fli1a:eGFP) (d) and summary of CVP sprouting in these embryos (e). (f–g) Representative images of the CVP in wild-type (left) and mutant Tg(kdrl:mCherry) (f) and summary of capillary loops in the CVP (g) in these embryos. (h) Representative images of PH3 immunostaining in the CVP of wild-type and mutant Tg(fli1a:eGFP) embryos. (i) Summary of PH3-positive cells in the CVP of wild-type and mutant Tg(fli1a:eGFP) embryos at 32 hpf. ***p < 0.001.
formation and distribution pattern of haematopoietic stem cells were not significantly changed at 2–3 dpf, while the expression of cmyb showed a slight increase at 4–5 dpf in slc39a5 morphants (electronic supplementary material, figure S4c). In situ hybridization data also showed an increased cmyb expression at 5 dpf, which suggests more haematopoietic stem cell formation in slc39a5 morphants (electronic supplementary material, figure S4c). One hypothesis to explain this phenomenon is that, since lots of red cells are accumulated in CVP, which leads to a lack of red cells in circulation, the increased haematopoietic cells in CHT could be a compensation for the insufficient blood in circulation. However, a detailed mechanism to illustrate the complex network between CVP formation and CHT haematopoiesis in the condition of slc39a5 deficiency should be explored in the future.

Zinc is a dietary micronutrient essential for proper cellular function, and its dynamics are closely associated with cell proliferation and migration in a variety of systems and tissues, including the immune system, central nervous system and skin [7,29–31]. Moreover, in vitro studies showed that the proliferation and migration of human lens epithelial B3 cells are inhibited by ZnCl2 treatment [32], whereas the migration of human microvascular endothelial cells increases under zinc-deficient conditions [33]. To further determine whether zinc homeostasis plays a role in endothelial cell migration in vitro, we performed a scratch assay to detect the cell migration speed under zinc and/or TPEN treatment by using human umbilical vein endothelial cells (HUVEC). The results showed that the cell migration speed was slowed down under zinc exposure, with significantly decreased expression of migration marker genes, including PECAM1 and MMP2 [34,35] (electronic supplementary material, figure S5a–d). Based on these data, we think that zinc homeostasis could affect the endothelium via an endothelial cell-specific manner.

In addition, a growing number of in vivo and in vitro studies suggest that zinc transporters play a role in regulating cell migration. For example, Taylor et al. [36] reported that knocking down Slc39a10 delayed epiboly in the developing zebrafish embryo. Other studies using cultured cell lines found that over-expressing SLC39A9 causes increased migration of glioblastoma cells [37], and increased expression of SLC39A6 has been associated with increased intracellular levels of zinc and increased tumour invasiveness, while knocking down SLC39A6 reduces the proliferation of oesophageal squamous cell carcinoma cells [38]. Although both cell proliferation and migration—two processes that have been studied extensively in angiogenesis—are...
triggered by a multitude of signals and guidance cues leading to proper endothelial cell path finding [39–41], the precise role of zinc and zinc transporters in regulating angiogenesis remains poorly understood.

In summary, our results support a model in which Slc39a5-mediated zinc homeostasis is essential for the normal development of venous angiogenesis, as loss of Slc39a5 results in systemic zinc accumulation, delayed CVP sprouting and impaired endothelial cell migration (figure 6). These findings provide insights into the way in which zinc regulates the microenvironment during angiogenesis, suggesting potential new molecular targets for the study and treatment of angiogenesis-related physiological and pathological conditions.

4. Material and methods

4.1. Zebrafish

Zebrafish were raised and maintained in accordance with guidelines established by the Laboratory Animal Center, Zhejiang University. Transgenic zebrafish lines were purchased from the Chinese Zebrafish Resource Center in Wuhan, China.

4.2. Whole-mount in situ hybridization

The 1-kb antisense probe was cloned into pEASY-T3 (TransGen Biotech), and the sense and antisense probes were synthesized using T7 or SP6 RNA polymerase (TR101-02, Vazyme; AM1340, mMESSAGE mMACHINE). In situ hybridization was performed essentially as previously described [42], and images were captured using a Nikon SMZ18 stereo microscope.

4.3. Microinjection of morpholinos and mRNA

Morpholinos (0.5 mM, GeneTools) and/or mRNA (200–300 pg/embryo) were injected into embryos in the single-cell stage. The sequence for the morpholinos were: slc39a5 5'-AGACTGTCTGTCCACCCA TGATTTT-3'; control 5'-CCTC- TTACCTCAGTTACAAITTTA-3'. mRNA was synthesized using the mMESSAGE mMACHINE SP6 transcription kit (AM1344, Ambion) and extracted and purified using the Quick-RNA MicroPrep kit (R1050, Zymo Research).

4.4. Generation of knockout zebrafish lines using CRISPR/Cas9

The Cas9/gRNA target site was designed and selected from the CHOPCHOP website (https://chopchop.cbu.uib.no/).
Primers containing the T7 or SP6 promoter sequences were synthesized and used for amplification from a gRNA template (pMD19-gRNA scaffold plasmid) [43]. gRNAs were transcribed in vitro using T7 or SP6 RNA polymerase. Cas9 protein (M0646T, New England Biolabs) and gRNA were mixed and then injected into single-cell zebrafish embryos, and T7 endonuclease 1 enzyme was used to evaluate the efficiency of target gene disruption. F0 embryos with the highest editing efficiency were raised to sexual maturity, and heterozygous F1 fish were obtained and identified from the offspring of F0 fishes using DNA sequencing. F1 fish carrying homozygous F1 fish were obtained and identified from the offspring of F0 fishes using DNA sequencing. F1 fish carrying homozygous mutant fish.

4.5. o-Dianisidine staining
An o-Dianisidine solution containing 100 mg o-Dianisidine dissolved in 70 ml ethanol was stored at 4°C in the dark. For staining, live embryos were exposed to the o-Dianisidine working solution consisting of 2 ml o-Dianisidine stock solution, 0.1 M sodium acetate (pH 4.5) and 0.65% H2O2 in the dark for 3–5 min at room temperature; images were captured using a Nikon SMZ18 stereo microscope (Nikon, Japan).

4.6. Zinc staining
The fluorescent zinc indicator FluoZin-3 AM (F24195, Invitrogen) was used to measure intracellular zinc concentration as reported previously [44]. Embryos were anesthetized, ground thoroughly in phosphate-buffered saline (PBS), and then centrifuged to remove the PBS. The cells were then suspended in 200 µl detection buffer consisting of 5 mM glucose, 1 mM MgCl2, 1 mM NaH2PO4, 1.3 mM CaCl2, 25 mM HEPES, 120 mM NaCl, 5.4 mM KCl and 1 µM FluoZin-3 AM (pH 7.5), and incubated at 37°C for 30 min. After washing several times with detection buffer, the cells were resuspended in 1 ml detection buffer and divided equally into three sample tubes. Tube 1 was used to measure the average fluorescence value (F); 5 µl zinc sulfate solution (10 mM, Sigma) and 5 µl NaPr cell membrane penetrating solution (Sigma) were added to tube 2 and used to measure maximum fluorescence (Fmax); finally, 7.5 µl TPEN solution (2 mM, Sigma) was added to tube 3 and used to measure minimum fluorescence (Fmin). Each sample was then adjusted to 500 µl with detection buffer, filtered through a 350 µm mesh filter into the flow tube, and analysed using flow cytometry. The following formula was then used to calculate the zinc concentration [Zn]: \[ K_D \times ((F - F_{\text{min}})/(F_{\text{max}} - F)) \], where \( K_D = 15 \) (nM).

4.7. Immunofluorescence and TUNEL assay
The TUNEL assay was performed using the fluorescein-based Roche In Situ Cell Death Detection Kit (12156792910, Roche) in accordance with the manufacturer’s instructions. For PH3 immunostaining, anti-pH3-Ser-10 (sc-8656-R, Santa Cruz) and Cy3-conjugated goat anti-rabbit (A0516, Beyotime) were used as the primary and secondary antibodies, respectively. For H2AX immunostaining, anti-phospho-Histone H2A.X (Ser139) (05-636-I, EMD Millipore) and Cy3-conjugated goat anti-rabbit (A0516, Beyotime) were used as the primary and secondary antibodies, respectively. BrdU immunostaining was performed using the BrdU Labeling and Detection Kit (10280879001, Roche) in accordance with the manufacturer’s instructions.

4.8. Reverse transcription and quantitative PCR
More than 30 embryos were anaesthetized and transferred to TRIzol Reagent (Ambion) for total RNA extraction. the PrimeScript RT Kit (RR047A, Takara Bio) was then used to synthesize cDNA using a standard protocol, and 2x SYBR Green qPCR Master Mix (B21202, Binake) was used for real-time PCR. Target gene expression was normalized to ef1a mRNA.

4.9. Spinning disk confocal live-cell imaging
Live embryos were embedded carefully in soft agarose (5265, TaKaRa), and images were captured in the time-lapse mode using a SpinSR10 spinning disk confocal super-resolution microscope (Olympus).

4.10. Quantification and statistical analysis
All experiments were performed using at least three biological replicates. Statistical analyses were performed with Prism (GraphPad) using a two-tailed Student’s t-test, and differences with a p-value < 0.05 were considered statistically significant.

4.11. Scratch assay
HUVEC were seeded in 12-well plates. After 24 h, a straight scratch wound was made by a sterilized 10 µl pipette tip and then washed by PBS. Then the cells were cultured in fresh medium for 12 h. Cells were washed by PBS and stained by Crystal Violet Staining Solution (0.2% solution dissolved in 10% ethanol) at 0, 6 and 12 h. Images of the scratch area were captured under an inverted microscope (Olympus, Japan).

Acknowledgements. This article has no additional data.

Authors’ contributions. Z.X., X.B., J.L., W.D., X.H. and L.Z. carried out the experiments, participated in data analysis, participated in the design of the study; Z.X., X.B. and F.W. coordinated the study, drafted the manuscript and critically revised the manuscript; J.M. and F.W. conceived of and designed the study; J.L., W.D., X.H. and L.Z. helped draft the manuscript. All authors gave final approval for publication and agree to be held accountable for the work performed therein.

Ethics. All animal studies were approved by the Animal Ethics Committee of Zhejiang University School of Medicine.

Data accessibility. This article has no additional data.

Competing interests. We declare we have no competing interests.
1. Betz C, Betz C, Lenard A, Belting HG, Affolter M. 2016 Cell behaviors and dynamics during angiogenesis. Development 143, 2249–2260. (doi:10.1242/dev.135616)

2. Schuermann A, Helker CS, Herzog W. 2014 Angiogenesis in zebrafish. Semin. Cell Dev. Biol. 31, 106–114. (doi:10.1016/j.semcdb.2014.04.037)

3. Wiley DM, Kim JD, Hao J, Hong CC, Bautch VL, Jin SW. 2011 Distinct signaling pathways regulate sprouting angiogenesis from the dorsal aorta and the axial vein. Nat. Cell Biol. 13, 686–692. (doi:10.1038/nccb2232)

4. Chavez MN, Aedo G, Fierro FA, Allende ML, Egaña M. 2004 The adaptive response to dietary zinc in mice involves the differential cellular localization and zinc regulation of the zinc transporters ZIP4 and ZIP5. J. Biol. Chem. 279, 49 082–49 090. (doi:10.1074/jbc.M409962200)

5. Noichihi K, Yuge S, Ando K, Wakayama Y, Mochizuki N, Ogawa R, Fukuhara S. 2019 Live imaging of angiogenesis during cutaneous wound healing in adult zebrafish. Angiogenesis 22, 341–354. (doi:10.1007/s10456-018-09660-y)

6. Maywald M, Wessells I, Rink L. 2017 Zinc signals and immunity. Int. J. Mol. Sci. 18, 2222. (doi:10.3390/ijms18102222)

7. Wang F, Kim BE, Petris MJ, Eide DJ. 2004 The effect of low dose iron and zinc intake during the first 1000 days of life: a systematic review and meta-analysis. Nutrients 8, 773.

8. Paty N, Olofin I, Boy E, Donahue Angel M, Rohner F. 2016 The effect of low dose iron and zinc intake on child micronutrient status and development during the first 1000 days of life: a systematic review and meta-analysis. Nutrients 8, 773.

9. Ogawa Y, Kawamura T, Shimada S. 2016 Zinc and skin biology. Arch. Biochem. Biophys. 611, 113–119. (doi:10.1016/j.jabb.2016.06.003)

10. Jeong J, Eide DJ. 2013 The SLB3 family of zinc transporters. Mol. Aspects Med. 34, 548–560. (doi:10.1016/j.mam.2012.05.008)

11. Zhao L et al. 2015 The zinc transporter ZIP12 regulates the pulmonary vascular response to chronic hypoxia. Nature 524, 356–360. (doi:10.1038/nature14620)

12. Takagishi T, Hara T, Fukada T. 2017 Recent advances in the role of SLCA9A/ZIP zinc transporters in vivo. Int. J. Mol. Sci. 18, 2708. (doi:10.3390/ijms18122708)

13. Wang F, Kim BE, Petris MJ, Eide DJ. 2004 The mammalian Zip5 protein is a zinc transporter that localizes to the basolateral surface of polarized cells. J. Biol. Chem. 279, 51 433–51 441. (doi:10.1074/jbc.M408361200)

14. Dufner-Beattie J, Kuo YM, Gitschier J, Andrews GK. 2004 The adaptive response to dietary zinc in mice involves the differential cellular localization and zinc regulation of the zinc transporters ZIP4 and ZIP5. J. Biol. Chem. 279, 49 082–49 090. (doi:10.1074/jbc.M409962200)

15. Wang X et al. 2019 The zinc transporter Slc39a5 controls glucose sensing and insulin secretion in pancreatic beta-cells via Sirt1- and Pgc-1alpha-mediated regulation of Glut2. Protein Cell 10, 436–449. (doi:10.1007/s13328-018-0580-1)

16. Jin J, Li Z, Liu J, Wu Y, Gao X, He Y. 2015 Knockdown of zinc transporter Zips (SLC39A5) expression significantly inhibits human esophageal cancer progression. Oncol. Rep. 34, 1431–1439. (doi:10.5829/orn.2015.4097)

17. Guo H et al. 2014 SLC39A5 mutations interfering with the BMP/TGF-beta pathway in non-syndromic high myopia. J. Med. Genet. 51, 518–525. (doi:10.1136/jmedgenet-2014-102351)

18. Gani J et al. 2012 Zebrafish globin switching occurs in two developmental stages and is controlled by the LCR. Dev. Biol. 366, 185–194. (doi:10.1016/j.ydbio.2012.02.021)

19. Chen J et al. 2016 Efficient extravasation of tumor-{

20. Hogan BM, Schulte-Merker S. 2017 How to plumb the axial vein. Dev. Cell 42, 567–583. (doi:10.1016/j.devcel.2017.08.015)

21. Bedell VM, Westcott SE, Ekker SC. 2011 Lessons from morpholino-based screening in zebrafish. Brief. Funct. Genomics 10, 181–188. (doi:10.1093/bfgp/elr021)

22. Eisen JS, Smith JC. 2008 Controlling morpholino experiments: don’t stop making antisense. Development 135, 1735–1743. (doi:10.1242/dev.001113)

23. Boisset JC, van Cappellen W, Boquet P, Greaves M, Hogstrand C, Schlomann U, Nimmanon T, Ziliotto S, Kille P, Hogstrand C. 2016 Zebrafish globin switching in vivo imaging of haematopoietic cells emerging from the mouse aortic endothelium. Nature 464, 116–120. (doi:10.1038/nature08764)

24. Boisset JC, van Cappellen W, Boquet P, Greaves M, Hogstrand C, Schlomann U, Nimmanon T, Ziliotto S, Kille P, Hogstrand C. 2016 Zebrafish globin switching in vivo imaging of haematopoietic cells emerging from the mouse aortic endothelium. Nature 464, 116–120. (doi:10.1038/nature08764)

25. Bertrand JY, Cisson JL, Stachuda DL, Traver D. 2010 Notch signaling distinguishes 2 waves of definitive hematopoiesis in the zebrafish embryo. Blood 115, 2777–2781. (doi:10.1182/blood-2009-09-244590)

26. Gering M, Patient R. 2005 Hedgehog signaling is required for adult blood stem cell formation in zebrafish embryos. Dev. Cell 8, 389–400. (doi:10.1016/j.devcel.2005.01.010)

27. Morand J, Briantso-Marjellet A, Lemarie E, Gonthier B, Arnaud J, Kornevich I, Godin-Ribault D. 2019 Zinc deficiency promotes endothelin secretion and endothelial cell migration through nuclear hypoxia-inducible factor-1 translocation. Am. J. Physiol. Cell Physiol. 317, C270–C276. (doi:10.1152/ajpcell.00460.2018)

28. Sun X, Huang S, Wang X, Zhang X, Wang X. 2018 CD300A promotes tumor progression by PECAM1, ADCY7 and AKT pathway in acute myeloid leukemia. Oncotarget 9, 2754–2764. (doi:10.18632/oncotarget.24164)

29. Xue F, Si X, Wang J, Yang A, Qin T, Yang Y. 2019 Nectin-3 is a new biomarker that mediates the upregulation of MMP2 and MMP9 in ovarian cancer cells. Biomed. Pharmacother. 110, 139–144. (doi:10.1016/j.biopha.2018.11.020)

30. Taylor KM, Murixina IA, Berthour D, Schmitt-Ulms G, Nimmanon T, Zillotto S, Kille P, Hogstrand C. 2016 Zinc transporter ZIP10 forms a heteromer with ZIP6 which regulates embryonic development and cell migration. Biochem. J. 473, 2531–2544. (doi:10.1042/BCJ20160388)

31. Münnich N, Wernhart S, Hogstrand C, Schloenmann U, Nimsky C, Bartsh JW. 2016 Expression of the zinc importer protein ZIP/ZSLC39A9 in glioblastoma cells affects phosphorylation states of p35 and GSK-3β and causes increased cell migration. BioMetals 29, 995–1004. (doi:10.1007/s10534-016-9971-z)

32. Cheng X et al. 2017 Solute carrier family 39 member 6 gene promotes aggressiveness of esophageal carcinoma cells by increasing intracellular levels of zinc, activating phosphatidylinositol 3-kinase signaling, and up-regulating genes that regulate metastasis. Gastroenterology 152, 1985–1997.e12. (doi:10.1016/j.gast.2017.02.006)

33. Lanniné B, Freitas C, Suchting S, Brunet I, Eichhorn A. 2009 Guidance of vascular development: lessons from the nervous system. Circ. Res. 104, 428–441. (doi:10.1161/CIRCRESAHA.108.188144)
40. Kir D, Schnettler E, Modi S, Ramakrishnan S. 2018 Regulation of angiogenesis by microRNAs in cardiovascular diseases. Angiogenesis 21, 699–710. (doi:10.1007/s10456-018-9632-7)

41. Harrison MR et al. 2015 Chemokine-guided angiogenesis directs coronary vasculature formation in zebrafish. Dev. Cell 33, 442–454. (doi:10.1016/j.devcel.2015.04.001)

42. Thisse C, Thisse B. 2008 High-resolution in situ hybridization to whole-mount zebrafish embryos. Nat. Protoc. 3, 59–69. (doi:10.1038/nprot.2007.514)

43. Chang N, Sun C, Gao L, Zhu D, Xu X, Zhu X, Xiong JW, Xi JJ. 2013 Genome editing with RNA-guided Cas9 nuclease in zebrafish embryos. Cell Res. 23, 465–472. (doi:10.1038/cr.2013.45)

44. Gao H et al. 2017 Metal transporter Slc39a10 regulates susceptibility to inflammatory stimuli by controlling macrophage survival. Proc. Natl Acad. Sci. USA 114, 12 940–12 945. (doi:10.1073/pnas.1708018114)