Tie2-Dependent Neovascularization of the Ischemic Hindlimb Is Mediated by Angiopoietin-2

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Abstract

The angiopoietins (ANGPT) are ligands for the endothelial cell (EC) receptor tyrosine kinase, Tie2. Angpt-1 is a Tie2 agonist that promotes vascular maturation and stabilization, whereas Angpt-2 is a partial agonist/antagonist involved in the initiation of postnatal angiogenesis. Therefore, we hypothesized that overexpression of Angpt-2 would be more effective than Angpt-1 for enhancing the perfusion recovery in the ischemic hindlimb. Perfusion recovery was markedly impaired in Tie2-deficient animals at day 35 in a model of chronic hindlimb ischemia. Injections of Angpt-2 or VEGFA plasmid at 7 days post femoral artery resection enhanced recovery and improved arteriogenesis as assessed by angiographic scores, whereas Angpt-1 or null plasmid had no effect. In addition, Angpt-2 together with VEGF resulted in greater improvement in perfusion and collateral vessel formation than VEGF alone. Similarly, conditional overexpression of Angpt-2 in mice improved ischemic hind limb blood flow recovery, while Angpt-1 overexpression was ineffective. These data from Tie2 heterozygote deficient mice demonstrate, for the first time, the importance of the Tie2 pathway in spontaneous neovascularization in response to chronic hindlimb ischemia. Moreover, they show that overexpression of the partial agonist, Angpt-2, but not Angpt-1, enhanced ischemic hind limb perfusion recovery and collateralization, suggesting that a coordinated sequence antagonist and agonist activity is required for effective therapeutic revascularization.

Introduction

The endothelial-selective receptor tyrosine kinase (RTK), Tie2, plays an essential role in blood vessel formation during embryonic development [1]. Targeted deletion of Tie2 [2] or its major agonist ligand, angiopoietin 1 (Angpt-1) [3], results in embryonic lethality in mice characterized by defects in blood vessel maturation, lack of recruitment of supporting pericytes and impaired basement membrane formation [3], and embryonic loss occurs at a slightly later stage than for mice deficient in vascular endothelial growth factor-A (VEGF) or its receptor, VEGFR2 [4]. Thus, VEGF and Angpt-1 appear to function in a temporally segregated yet complimentary manner in the blood vessel formation in the developing embryo [5,6,7]; however, the role of the angiopoietin system in postnatal angiogenesis is less clear.

Angiopoietin-2 (Angpt-2) is another major Tie2 ligand. While both Angpt-1 and Angpt-2 bind to Tie-2 with equal affinity [8], Angpt-2 has been characterized as a functional antagonist of Tie2 [8], blocking the effects of Angpt1 on Tie2 activity. The finding of increased Angpt-2 expression at the leading edge of tumour neovessels [9] has led to the concept that Angpt-2 is required to release endothelial cells (EC) from the tonic inhibitory effect of Angpt1 and facilitate EC activation in response to VEGF [10]. Moreover, in the absence of VEGF, Angpt-2 has been shown to promote EC apoptosis [11] and has been implicated in mediating vascular regression in the involuting corpus luteum [12]. However, it has recently been recognized that Angpt-2 exhibits context-dependent agonist activity, inducing activation of Tie-2 in a time-dependent manner to levels similar to Angpt-1 at high concentrations [13] or during prolonged (i.e. 12 to 24 hours) exposure [14], which corresponds to the time course of capillary-like network formation in cultured ECs [14]. These findings point to a possible role for Angpt-2, not only as an inhibitor of Tie2 in the initiation of the angiogenic response, but also as an agonist in the later stages of blood vessel formation and maturation that are dependent on Tie-2 activation [7].

Previously, there have been conflicting reports on the role of the angiopoietins in postnatal angiogenesis and neovascularization. In the corneal implant model, Angpt-1 was shown to enhance neovessel density in combination with VEGF, but had no effect by itself, whereas Angpt-2 increased length but not the density of neovessels [15]. Similarly, synergistic effects of Angpt-1 and VEGF were seen in the ischemic hindlimb model [16], whereas, Angpt-2 was reported to impair angiogenesis in the same model [17]. However, others have reported that Angpt-2 is highly expressed in vascular regions undergoing active angiogenesis [18] and plays a requisite role in postnatal angiogenic vascular remodeling [19]. Moreover, it was recently shown that the selective inhibition of Angpt-2 activity impaired recovery of blood flow in the ischemic hind limb [20], consistent with an important role for the
endogenous ligand in angiogenesis and collateral vessel formation in this model.

In the present study, we show for the first time that Tie2 deficiency resulted in exacerbation of limb loss and impaired spontaneous perfusion recovery in the context of hindlimb ischemia. Moreover, overexpression of Angpt-2, but not Angpt-1, markedly enhanced collateral formation in the rat hindlimb ischemia model, which was further augmented by combination with VEGF. As well, induction of Angpt-2 in conditional transgenic mice also increased circulating levels of progenitor cells. These data strongly support the predominant role of Angpt-2 in postnatal angiogenesis and collateral vessel formation.

Results

Tie2 Deficiency Results in Increased Limb Necrosis and Impaired Perfusion Recovery

Tie2 protein and activity was decreased by 40–50% in Tie2+/− versus Tie2+/+ mice (Figure 1a and b). Interestingly, eNOS protein expression was also reduced in Tie2-deficient animals (Figure 1c). Using the critical ischemia model, we tested the functional importance of Tie2 deficiency on limb survival. Wide surgical excision of the femoral artery produced immediate and profound reduction of hindlimb perfusion at day 0 and critical limb ischemia in both Tie2+/+ and Tie2+/− animals (Figure 2a), with early (day 7) signs of tissue necrosis and distal forefoot loss were observed in Tie 2 deficient mice, associated with significantly reduced perfusion by LDPI. Tie2+/− animals also exhibited a greater incidence of limb loss compared to Tie2+/+ animals at 42 days (75% versus 33%, p<0.05).

Femoral ligation with limited excision resulted in a stable model of chronic hindlimb ischemia without evidence of limb necrosis and tissue loss in either Tie2-deficient or WT mice. Hindlimb perfusion, assessed by serial LDPI measurements over 5 weeks, was reduced immediately after surgery in both Tie2+/+ and Tie2+/− mice (0.14±0.03 and 0.20±0.02, respectively) (Figure 2b). Recovery of perfusion in the ischemic (I) vs. the nonoperated (NI) limb was reduced in Tie2 deficient compared to WT mice at day 35 (I/NI 0.42±0.04 vs. 0.79±0.06, respectively; p<0.001) (Figure 2c). Similarly, the active use of the ischemic limb was impaired after surgery, as reflected by an increase in the Movement Score [21,22] (Figure 1d), which improved more rapidly and completely in the WT compared to Tie2 deficient animals (day 28: 0.8±0.20 vs. 1.67±0.21, respectively; p<0.05).

Interestingly, intramuscular injection (day 7) of plasmid-encoded eNOS into the ischemic hindlimb of Tie2 deficient mice enhanced perfusion recovery in comparison to control (Tie2+/−) mice treated with null plasmid (0.66±0.04 vs. 0.44±0.03, respectively, p<0.05) (Figure 3) to levels similar to WT (0.79±0.05).

Tie2 Deficient Mice Display Diminished Microvascular Perfusion

Fluorescent microangiography revealed similar high levels of microvascular perfusion in the non-ischemic limb in WT and Tie2-deficient animals (Figure 2e). In contrast, there was a profound reduction in microvascular perfusion 14 days after surgery in ischemic limbs, which was more pronounced in the Tie2-deficient animals. Vessel score was determined using morphometric software as a composite of vessel length, nodal branching points and vessel segment end numbers (Figure 2f). Fourteen days after surgery, the hindlimb vessel score was profoundly decreased in both Tie2+/+ and Tie2+/− mice; however, this score was greater in Tie2+/+ than in Tie2+/− mice after 14 days (217.2±29.4 vs 69±18.5, respectively; p<0.01) indicative of a less recovery of deep muscle perfusion in the Tie2 deficient model.

Spatio-temporal expression profile of Angiopoietins in the ischemic hindlimb

Endogenous mRNA expression of Angpt-2 was found to be increased in the ischemic gastrocnemius muscle at day 1 (Figure 4a and b), returning to baseline expression levels by day 28 as the endogenous blood flow recovery reached a plateau. In contrast, there was a reciprocal change in regulation of Angpt-1 expression,
which was unchanged or even reduced initially and increased at later time periods, particularly in more proximal regions of the limb (i.e. adductor muscle) (Figure 4c). Despite these differences in expression profile, there was a parallel change in the ratio of Angpt-2/Angpt-1, which was increased by more than 2-fold at day 1, and reduced by about 50% at day 28, in both the ischemic gastrocnemius and the more proximal adductor muscles (Figure 4d). However, in the adductor muscle the early increase in ration was due mainly to a reduced expression of Angpt1, whereas in the gastrocnemius it was driven mainly by an increase in Angpt-2.

Angpt-2 gene transfer enhances perfusion in response to ischemia

Figure 5a shows representative images of hindlimb perfusion before (day 0), just prior to intramuscular delivery of null (control)
plasmid or Angpt-1, Angpt-2, and VEGF plasmids alone or in combination (day 7), and 35 after surgical induction of ischemia (4 weeks after gene delivery). Immediately after surgery, perfusion was decreased in the ischemic vs. the non-operated limb in all groups to a similar extent (I/NI: 0.25–0.32, Figure 5b). A modest recovery of ischemic hindlimb perfusion observed at day 7, prior to randomization to the various treatment groups. Rats receiving Angpt-2 transfer alone exhibited enhanced hindlimb perfusion by day 35, (Figure 5b; I/NI: 0.69±0.04 and 0.63±0.05, respectively, p<0.05 compared with control; 0.50±0.02). However, no significant improvement was seen within the group treated with the Angpt-1 plasmid (0.55±0.05), which was not different from the null-plasmid control group. Perfusion recovery in the ischemic limb was greater with Angpt-1 combined with VEGF (Figure 5c; 0.66±0.02) compared to Angpt-1 alone (p<0.05) but not different from VEGF alone (Figure 5c). In contrast, the combination Angpt-2 and VEGF gene transfer resulted in further improvement in ischemic limb perfusion (0.75±0.02, Figure 5c and d), significantly better than VEGF or Angpt-2 alone (p<0.05).

Effect of Angiopoietins on collateral vessel formation

Immediately after operative excision of femoral vasculature, no collaterals were detected in the thigh (adductor and vastus muscle groups) and there was only minimal distal leg filling. At day 35, some bridging collaterals originating from the internal iliac artery were visible in the thigh of null plasmid control animals with minimal distal filling, whereas collateral vessels and distal filling was much more pronounced in the animals treated with Angpt-2 and VEGF alone or in combination (Figure 6a). Collateralization was expressed as an angiographic score, defined as the ratio of vessel counts intersecting a grid in the ischemic versus non-ischemic hindlimb (Figure 6b). Treatment with VEGF or Angpt-2 alone or their combination resulted in a significant increase in collateral vessel formation compared with control animals, whereas no significant effect was seen with Angpt-1 alone. However, the combination of VEGF and Angpt-2 was not superior to VEGF or Angpt-2 alone.

Enhanced perfusion recovery in transgenic mice overexpressing Angpt-2

All binary transgenic (BT) animals were maintained on Dox in order to suppress transgene expression until the time of surgical induction of hindlimb ischemia. Measurement of human angiopoietin levels in plasma samples derived from binary transgenic (BT) mice confirmed conditional expression of hAngpt-1 and hAngpt-2 one week after Dox withdrawal (Figure 7a and b). Induction of transgene expression by withdrawal of Dox at the time of surgery significantly increased recovery of perfusion of the ischemic leg in hAngpt-2 BT mice at 4 weeks (I/NI: 0.78±0.07) compared with their NBT littermate controls (0.52±0.06, p<0.01), whereas there was no improvement seen in Angpt-1 BT mice (0.44±0.05, Figure 7c and d; Angpt-2 (but not Angpt-1) overexpression also increased the mobilization of c-kit/VEGFR2 double positive circulating progenitor cells in the peripheral blood (Figure 7c). Similarly, fluorescent microangiography (FMA) revealed improvement in functional vascularity of ischemic hindlimb muscle in the hAngpt-2 BT animals at 4 weeks post induction of ischemia (Figure 8a and b) compared to the hAngpt-1 BT and NBT mice.

Discussion

To our knowledge, this is the first demonstration in the adult of the central role of the Tie2 receptor in the spontaneous neovascularization response to ischemia. We also show that the context-dependent partial Tie2 agonist/antagonist, Angpt-2, is a more potent proangiogenic and arteriogenic factor than Angpt-1 both in both rat and mouse hindlimb ischemia models, and may be a potentially useful therapeutic agent for the treatment of severe chronic ischemic syndromes.

Angpt-1 activates Tie2 to promote EC survival and quiescence and thus contributes to neovessel stabilization and maturation during embryonic development, as well as the maintenance of postnatal vascular homeostasis [7]. Conversely, Angpt-2 is believed to function as an antagonist of Tie2 [8], releasing ECs from the tonic stabilizing effects of Angpt-1, and thus permitting efficient EC activation and plasticity in the response to classical angiogenic mediators, such as VEGF. However, recent reports suggest that Angpt-2 may acquire agonist activity under specific conditions, in particular during more prolonged exposure to Tie2 [13,14]. Thus, this bimodal activity of Angpt-2 may be crucial in promoting efficient neovascularization by contributing to both the “activation” and “consolidation” phases of angiogenesis.
In order to better define the relevance of these changes for neovascularization, we used two complementary strategies to overexpress Angpt-1 or Angpt-2 in the ischemic hindlimb: plasmid-based gene transfer and transgenic conditional overexpression. A consistent finding was the superior efficacy of Angpt-2 in restoring perfusion in the ischemic hindlimb compared with Angpt-1 using both approaches. Moreover, the effect of Angpt-2 on neovascularization was at least additive with VEGF in the rat model. The greater efficacy of Angpt-2 may be related to its bimodal functional activity, with an ability to both inhibit and activate Tie2 in a defined temporal sequence that closely corresponds to the time course of angiogenesis [14].

The lack of ability of Angpt-1 to induce neovascularization is consistent with a previous study showing that gene transfer of Angpt-1 alone or combined with VEGF using an adenovirus vector failed to enhance neovascularization in the rat [23]. Furthermore, myocardial overexpression of Angpt-1 has been shown to inhibit VEGF-induced angiogenesis in a cardiac specific transgenic model [24]. However, several other studies have reported increased neovascularization in response to Angpt-1 [16,25] but not Angpt-2. The reasons for the discrepancy between our study and the one by Shyu et al. may relate to species utilized and the “dose” of plasmid DNA used, which was very different in the two studies [16]. Shyu et al. injected 125 µg of plasmid-encoding DNA into the larger volume of the rabbit hindlimb compared to our study utilizing 500 µg in the smaller rat hindlimb. Finally, it is likely that there were differences in the degree of induced ischemia between the two models given the greater size of rabbit hindlimb and species differences in pre-existing collaterals.

There have been relatively few studies that have examined the ability of Angpt-2 either alone or in combination with VEGF to induce angiogenesis, and these have produced conflicting results. A previous report showed that increased expression of VEGF and Angpt-2 in transgenic mice had a synergistic effect on enhancing capillary number in the myocardium [24], though this was in the absence of a physiologically relevant ischemic stimulus. In contrast, a report by Reiss et al. [17] using a binary transgenic model system similar to ours showed that targeted vascular overexpression of Angpt-2 reduced perfusion recovery in the ischemic hindlimb, a finding seemingly opposite of that described in the present report. However, different targeting strategies were used in these two studies, which likely resulted in marked differences in local concentrations of Angpt-2 in the ischemic limb. In the present study, the transgene was targeted to the liver using a hepatic-specific promoter, and delivery of Angpt-2 to the ischemic hindlimb was entirely dependent on hepatic release and transport via the circulation, thus mimicking the effects of systemic administration of a therapeutic protein. In contrast, the earlier report targeted the Angpt-2 transgene directly to the vascular endothelium-specific Tie1 promoter, thus Angpt-2 was directly overexpressed locally within the vascular endothelium of the hindlimb. Although both studies reported similar increases in circulating Angpt-2 levels, endothelial-targeted expression would...
result in far greater local vascular levels, and these very high concentrations of Angpt-2 have been shown to have potentially deleterious effects on vascular integrity and function [26,27]. In contrast, in our transgenic system, the site of Angpt-2 generation and its action were segregated, and thus any effects on the ischemic limb are solely attributable to the effects of increased circulating protein. Similarly, the gene transfer model utilized direct plasmid DNA injection that results in a relatively modest degree of muscle transfection [28,29], and which does not result detectable increases in circulating levels [30].

In support of the possibility that differences in local vascular Angpt-2 levels could be responsible for the discrepant findings between our study and that of Reiss et al, this group showed a clear “dose-dependent” effect such that mice exhibiting highest

Figure 5. Laser doppler perfusion imaging serial assessment of hindlimb perfusion. (a) Representative laser doppler perfusion imaging scans of rats treated with intramuscular injections of plasmids encoding for human Angpt-1 alone (- □, n = 7), Angpt-2 alone (- ○, n = 8), VEGF alone (- Δ, n = 9) or combination of Angpt-1+VEGF (- △, n = 8), Angpt-2+VEGF (- ○, n = 7). Controls received null plasmid (- △, n = 9). Animals were imaged on days 0, 7 and 35 after surgically-induced left hindlimb ischemia. Blood flow is displayed by color-coded pixels where blue indicates low perfusion and red-white denotes high perfusion. Assessment of perfusion was performed on the lower leg and foot. (b) A serial quantitative evaluation of hindlimb blood perfusion expressed as a ratio of ischemic leg perfusion to that in the non-ischemic contralateral limb (I/NI) is shown for all single gene transfer groups. Combination gene transfer groups are shown in (c). (d) The increment in perfusion recovery (%) from day 7 (time of gene injection) to day 35. * denotes p<0.05 versus controls and Angpt-1 treated. # denotes p<0.05 versus VEGF and Angpt-2 treated animals. doi:10.1371/journal.pone.0043568.g005
Angiopoietin-2 and Ischemic Neovascularization

In this context, the reduction in basal eNOS expression in Tie2- 
[32], which is also an important pathway in angiogenesis 
[33,34]. Moreover, Angpt-2 has recently been shown to 
reduce atherogenesis by an NO/eNOS-dependent mechanism 
in the ischemic hindlimb model when expressed at levels that are 
physiologically relevant. The superiority of Angpt-2 may be due to 
its bifunctional role as a modulator of Tie2 activation, incorpo-
rating both antagonist and agonist activities and thus contributing 
to the initial increase in vessel plasticity and sprouting, while 
ensuring maturation in the later phases of neovascularization 
[37]. These data support further exploration of the use of Angpt-2 
as a potential proangiogenesis agent.

In conclusion, these data demonstrate that Angpt-2, but not 
Angpt-1, is a potent mediator of angiogenesis and collateralization 
in the ischemic hindlimb model when expressed at levels that are 
physiologically relevant. The superior angiogenic activity of Angpt-2 
may be due to its bi-functional role as a modulator of Tie2 activation, incorpo-
rating both antagonist and agonist activities and thus contributing 
to the initial increase in vessel plasticity and sprouting, while 
ensuring maturation in the later phases of neovascularization 
[37]. These data support further exploration of the use of Angpt-2 
as a potential proangiogenesis agent.

Materials and Methods

Animals: Rodent Hindlimb Ischemia Model

All animal protocols were approved by the Institutional Animal 
Care and Use Committee at St. Michael’s Hospital. Male 
Sprague-Dawley rats (500–650 gm) were anesthetized with 
intraperitoneal Ketamine (200 mg/kg) and Xylazine (100 mg/ 
kg). Surgery was performed to induce a model of chronic unilateral 
hindlimb ischemia. After a lower abdominal midline incision, the entire 
 femoral artery was exposed and excised after ligation of the 
 superficial femoral, profundus, circumflex, and superficial 
epigastric arteries. Details for the preparation of the mouse 
hindlimb ischemia model are provided in detail in Materials and 
Methods S1.

Gene Transfer Studies

Rats underwent surgery (day 0) to induce hindlimb ischemia. 
On day 7, animals were divided into six groups that received 
intramuscular injections of plasmid DNA encoding human Angpt-
1 (n = 7), Angpt-2 (n = 6), VEGF (n = 9) or combination of Angpt-
1+VEGF (n = 8) and Angpt-2+VEGF (n = 7). Control animals 
received null (empty) plasmid (n = 9). Serial, non-invasive assess-
ment of perfusion by laser doppler perfusion imaging was 
performed on day 0, 7, 14 and 35. Animals were sacrificed at 
day 35. At sacrifice, angiography was done to assess for degree of 
collateralization. In a separate groups of rats, the gene expression 
profile of Angpt-1, Angpt-2, VEGF following 1 (n = 5) and 28 (n = 5) 
days of hindlimb ischemia was assessed from muscles taken from 
ischemic and contralateral non-ischemic limbs for real-time reverse 
transcriptase polymerase chain reaction (RT-PCR) anal-
ysis.
Intramuscular Gene Transfer

Human Angpt-1, Angpt-2 and VEGF_164 cDNA were subcloned into cytomegalovirus (CMV) promoter-driven and ampicillin-resistance gene containing, mammalian cell expression vectors pFLAG, pCRUni3.1 and pVax respectively. On day 7, plasmid DNA was delivered by five injections in the ischemic leg, namely, two into the adductor, one into vastus medialis and two sites in the gastrocnemius muscle using a 1-mL syringe and a 30-gauge needle. A total of 500 μg (single gene transfer) or 1000 μg (combination gene transfer) of plasmid DNA in a constant volume of 1000 μL of saline was injected. In mouse studies, human eNOS was subcloned into the plasmid pVax. On day 7 plasmid DNA (150 μg in 100 μL saline) was delivered via injection (BD Ultrafine Insulin syringe) similar to gene transfer studies in the rats.

Laser Doppler Pefusion Imaging

Both rats and mice were anesthetized with Ketamine (200 mg/kg) and Xylazine (10 mg/kg). Hair was removed using an electric shaver. Serial, non-invasive assessment of ischemic limb microvascular perfusion was performed in triplicate and in a blinded manner using the LDPI system (Moor instruments, United Kingdom) after placing all mice on a homeothermic heating pad maintained at 37°C. Identical regions equal in area encompassing the distal leg (entire foot) of both ischemic and contralateral non-

Figure 7. Confirmation of Angpt-1 and Angpt-2 overexpression in BT mice and its effects on ischemic neovascularization. ELISA quantitation of conditional hAngpt-1 (a) and hAngpt-2 (b) expression in binary transgenic (BT) and in control, non-binary transgenic (NBT) mice one week after withdrawal from doxycycline. (c) Representative laser doppler perfusion images performed serially immediately after femoral vasculature surgical excision (Day 0) and after four weeks are shown. (d) Quantitative serial evaluation of hindlimb perfusion in hAngpt-1 BT (n = 6), hAngpt-2 BT (n = 10) and NBT (n = 8) mice withdrawn from doxycycline on the same day as induction of hindlimb ischemia by femoral artery excision. * denotes p<0.01 versus Angpt-1 BT and NBT. (e) Mobilization of bone marrow-derived endothelial progenitor cells into the peripheral blood (VEGFR-2+/c-kit+) seven days after doxycycline withdrawal in NBT (n = 6), Angpt-1 BT (n = 4) and Angpt-2 BT (n = 5) mice. * denotes p<0.05 vs. NBT.

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Intramuscular Gene Transfer

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ischemic limbs were assessed for quantification of perfusion using the LDPI image processing software (v5.0). To minimize data variation possibly secondary to ambient temperature changes all perfusion data were expressed as a ratio of operated ischemic (I) to non-operated, non-ischemic (NI) limb perfusion.

**Angiographic assessment of collateral circulation**

Rats were anesthetized with Ketamine (200 mg/kg) and Xylazine (10 mg/kg). An angiocatheter (24G, Becton-Dickinson, Franklin Lakes, NJ) was inserted into the abdominal aorta, below the renal arteries and above the iliac artery bifurcations as described in detail in Materials and Methods S1.

**Fluorescent microangiography of hindlimb vasculature**

A catheter was inserted into the mouse abdominal aorta and then flushed with heparinized heparinized saline followed by right atrium venting. A warmed (45°C) solution of 1% low melting point agarose (Sigma) containing 0.2 μm yellow-green fluorescent microspheres (305/315 nm excitation-emission, Life Technologies, Carlsbad, CA) was then slowly infused into the aorta at a constant rate resulting in bilateral hindlimb perfusion. The animal was immediately placed on ice and cooled to promote solidification of the agarose solution. Harvested adductor muscle was then placed in OCT on dried ice and subsequently sectioned (100 μm) using an oscillating blade microtome (Leica, Richmond Hill, Canada). Confocal microscopy (Bio-Rad Laboratories, Mississauga, Canada) was used to produce a Z stack of images that were projected in order to quantify vessel number using automated software (IPTK analysis software, Reindeer Graphics Inc., Asheville, NC). This technique has previously been used to quantify pulmonary vasculature density in a rat experimental model of pulmonary hypertension [38].

**Tie2 Deficient Mice**

The generation of Tie2 knockout mice by targeted deletion of the Tie2 gene has been previously reported [39]. Whereas the homozygous null mutation is embryonic lethal, heterozygous Tie2-deficient mice are viable with no overt phenotype. Genotypes of wild-type (WT, Tie2+/+) and heterozygote deficient (Tie2+/−) mice were determined by PCR. Mice were on a CD1 genetic strain background. All animal protocols were approved by the Institutional Animal Care and Use Committee at St. Michael’s Hospital. Heterozygote deficient and wild-type littermates were used at 12–18 weeks of age. Genomic DNA was extracted from mouse ear-notch samples and PCR was performed using the buffers and PCR reaction mix from the REDExtract-N-Amp™ Tissue PCR Kit, following manufacturer’s instructions (Sigma-Aldrich, St. Louis, MO). Briefly, samples were incubated in a 1:4 mixture of the tissue preparation solution and extraction solution for 10 minutes at room temperature, heated to 95°C for 3 minutes, and then mixed with a neutralization solution to deactivate inhibitory substances prior to PCR. A 0.5 μm mixture of the following primers was used in the PCR to detect the transgene-inserted Tie2 (~350 bp) and endogenous Tie2 (~150 bp): Forward (endogenous): 5'- AGC GGC TGG ACC ATG CGA GC - 3', Reverse (endogenous): 5'- CGA TTG CTC AGC GGT GCT GTC CAT - 3', and Forward (transgene): 5'- AGG AGC AAG CTG ACT CGA CAG - 3'. The PCR thermal cycling parameters included an initial denaturation at 94°C for 3 min., followed by 30 cycles consisting of denaturation at 95°C for 30 sec., annealing at 54°C for 40 sec., and extension at 72°C for 30 sec. Following this, a final extension at 72°C for 10 min. was performed and the amplified DNA resolved in a 1.5% agarose gel.

**Conditional Angiopoietin Transgenic Mice**

Two binary transgenic (BT) mouse models in a CD1 background were generated with conditional overexpression of human Angpt-1 (hAngpt-1) or human Angpt-2 (hAngpt-2), respectively, driven by a tetracycline (doxycycline) responsive, liver specific, Lap-1 promoter (Figure 9) [40], as is more fully described in the expanded in Materials and Methods S1. BT (Lap:hAngpt-1 and Lap:hAngpt-2) and single transgenic (ST: Lap, hAngpt-1 or hAngpt-2) mice were maintained on drinking water containing Dox (100 μg/mL) in a 5% drinking solution from conception up to adulthood. Mice (16–24 weeks old) were withdrawn from Dox at the time of surgical induction of unilateral
hindlimb ischemia (day 0), as previously described. Microvascular hindlimb perfusion was assessed by laser doppler perfusion imaging on day 0 as well as 35 and expressed as a ratio of ischemic to non-ischemic limb perfusion. Mice were sacrificed on day 35.

Measurement of human Angpt-1 and Angpt-2

The concentration of hAngpt-1 and hAngpt-2 in the serum of transgenic mice was measured with enzyme-linked immunosorbent assay (ELISA) human Angpt-1 and Angpt-2 kits (R&D Systems, Minneapolis, MN) as per the manufacturer’s instructions. Blood collected by cardiac puncture from BT:hAngpt-1 and BT:hAngpt-2 overexpressing mice was placed into ethylenediaminetetraacetic acid (EDTA)-lined tubes and spun at 1000 g for 15 minutes at 2–5 °C within 30 minutes of collection. Plasma from BT:hAngpt-1 expressing mice was spun again at 10,000 g for 10 minutes at 2–5 °C to ensure complete platelet removal prior to assay and stored at −70 °C.

RNA Extraction and Quantitative Reverse Transcription – Polymerase Chain Reaction (qRT-PCR)

Total RNA was extracted from mouse liver as well as rat hindlimb and contralateral limb adductor and gastrocnemius muscles using Trizol extraction (Life Technologies, Carlsbad, CA) as per manufacturers’ protocols. Extracted RNA (1 μg) and omniscript reverse transcriptase (Qiagen, Hilden, Germany) were used in RT reactions that were diluted 1/20. Each diluted RT was then used in PCRs in duplicates to assess the RNA abundance of rat Angpt-1, Angpt-2, VEGF and Tie2 using the SYBR Green PCR Master Mix and the ABI PRISM 7900HT sequence detection system (Applied Biosystems, Life Technologies, Carlsbad, CA). Further detail including a complete list of the sequences for primers utilized in these studies is located in Materials and Methods S1.

Peripheral Blood Cell Mobilization

The profile of circulating progenitor cells was analyzed in transgenic mice 7 days after doxycycline withdrawal. Briefly, animals were anesthetized and underwent cardiac puncture for blood withdrawal (800–1000 μL) collected in K3-EDTA tubes (BD Biosciences, Mississauga, Canada). Following red blood cell lysis with Lyse buffer (Sigma), flow cytometry analysis for c-kit (CD117) and VEGFR-2 were performed using a FC500 flow cytometer (Beckman Coulter, Brea, CA).

Statistical Analysis

Data was expressed as mean ± SEM. Comparison between two means was done using the Student’s unpaired t-test. For all other comparisons, one-way ANOVA with a post-hoc Neuman-Keuls test for multiple comparisons was used. Significance was set at p<0.05.

Supporting Information

Materials and Methods S1 Expanded list of materials and methods used in this study.

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Author Contributions

Conceived and designed the experiments: ML SHJM DJS. Performed the experiments: ML PL SHJM YD. Analyzed the data: ML SHJM DJS. Contributed reagents/materials/analysis tools: DJD. Wrote the paper: ML SHJM DJS.
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