RESEARCH ARTICLE

Endothelial-specific depletion of TGF-β signaling affects lymphatic function

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Abstract

Background: Transforming growth factor (TGF)-β is a multifunctional cytokine involved in cell differentiation, cell proliferation, and tissue homeostasis. Although TGF-β signaling is essential for maintaining blood vessel functions, little is known about the role of TGF-β in lymphatic homeostasis.

Methods: To delineate the role of TGF-β signaling in lymphatic vessels, TβRIΙfl/fl mice were crossed with Prox1-CreERT2 mice to generate TβRIΙfl/fl; Prox1-Cre ERT2 mice. The TβRIΙ gene in the lymphatic endothelial cells (LECs) of the conditional knockout TβRIΙΔLEC mice was selectively deleted using tamoxifen. The effects of TβRIΙ gene deletion on embryonic lymphangiogenesis, postnatal lymphatic structure and drainage function, tumor lymphangiogenesis, and lymphatic tumor metastasis were investigated.

Results: Deficiency of LEC-specific TGF-β signaling in embryos, where lymphangiogenesis is active, caused dorsal edema with dilated lymphatic vessels at E13.5. Postnatal mice in which lymphatic vessels had already been formed displayed dilation and increased bifurcator of lymphatic vessels after tamoxifen administration.Similar dilation was also observed in tumor lymphatic vessels. The drainage of FITC-dextran, which was subcutaneously injected into the soles of the feet of the mice, was reduced in TβRIΙΔLEC mice. Furthermore, Lewis lung carcinoma cells constitutively expressing GFP (LLC-GFP) transplanted into the footpads of the mice showed reduced patellar lymph node metastasis.

Conclusion: These data suggest that TGF-β signaling in LECs maintains the structure of lymphatic vessels and lymphatic homeostasis, in addition to promoting tumor lymphatic metastasis. Therefore, suppression of TGF-β signaling in LECs might be effective in inhibiting cancer metastasis.

Keywords: TGF-β, Lymphatic vessel, Endothelial cell, Prox1, Tumor metastasis

Background

Transforming growth factor-β (TGF-β) is a secreted dimeric protein that has pleiotropic effects and plays a key role in many cellular processes during both embryogenesis and tissue homeostasis in adults [1]. Therefore, abnormal TGF-β signaling has also been associated with various diseases, including cancer, fibrotic disorders, and cardiovascular diseases [2] [3]. The TGF-β signaling pathway is initiated through two different serine/threonine kinase receptors: type II (TβRIΙ) and type I (TβRI; also termed activin receptor-like kinase-5 [ALK5]). In canonical TGF-β signaling, the activated receptor complex phosphorylates receptor-regulated Smads (R-Smads; i.e., Smad2 and Smad3) at two serine residues at their C-terminal to permit the phosphorylation of two R-Smads to form ternary complexes with the common partner Smad (Co-Smad), Smad4. The R-Smad/Co-Smad complex then enters the nucleus where it acts as a transcriptional factor to regulate the expression of TGF-β target genes.
genes in cell type-specific and context-dependent manners [4].

Lymphatic vessels maintain homeostasis by balancing tissue fluid throughout the body, regulating inflammation via the immune system, forming new lymphatic vessels from preexisting lymphatic vessels upon environmental stimulus, and contributing to tumor metastasis [5]. Lymphatic progenitor cells originate from the venous endothelial cells during embryonic development. The expression of the transcriptional factor Prox-1 is imperative for the development and maintenance of lymphatic endothelial cells (LECs). Prox1, a master regulator of LECs, induces the expression of vascular endothelial growth factor receptor 3 (VEGFR3) in the lymphatic progenitor cells that bud from the cardinal vein. Subsequently, the lymphatic progenitor cells form the lymph sac in a VEGF-C/VEGFR3 dependent manner. The VEGF-C/VEGFR3 pathway plays a central role in spreading the lymphatic network throughout the body [6, 7]. Although the molecular mechanisms that control lymphangiogenesis have been elucidated, little is known about the mechanisms that maintain the plasticity and stability of lymphatic vessels.

Mouse genetic studies regarding the deficiency of TGF-β signaling in fetal LECs have revealed a significant reduction in lymphatic vessel germination and remodeling in the absence of TGF-β [8]. Contrastingly, TGF-β inhibited LEC differentiation through suppression of Prox1 and LYVE-1 expression in cultured LECs. Consistently, the blockage of endogenous TGF-β signaling by a chemical inhibitor enhanced lymphangiogenesis in a mouse model of chronic peritonitis [9]. Thus, the nature of involvement of TGF-β signaling in lymphangiogenesis is still unclear.

In this study, we showed that TGF-β signaling plays a key role in maintaining the structure and function of lymphatic vessels. LEC-specific deletion of the TβRII gene results in dilation of the lymphatic lumen and impaired lymphatic drainage function. When Lewis lung carcinoma cells constitutively expressing green fluorescent protein (LLC-GFP) were implanted into mice lacking the TβRII gene in their LECs, tumor lymphatic vessels were dilated and lymphatic metastasis was suppressed. These results indicate that TGF-β signaling acts as a tumor malignant factor via lymphatic endothelial cells.

Materials and methods

Mice
TbβRIIfl/fl mice [10] were crossed with Prox1-CreERT2 mice [11] to generate TbβRIIfl/fl, Prox1-CreERT2 mice. Tamoxifen (Tx) (Sigma-Aldrich, St. Louis, MO T5648) dissolved in corn oil (20 mg/mL) was intraperitoneally administrated to mice (40 mg/kg/day) for consecutive 5 days into control (TbβRIIfl/fl) mice and TbβRIIfl/fl, Prox1-CreERT2 mice, where TbβRII gene in LECs is deficient (TbβRIIfl/LEC). For embryonic analysis, pregnant mice were administrated with Tx (40 mg/kg/day) 10.5 and 11.5 days after mating. Then, mice were sacrificed to analyze embryos at E13.5. The lymphatic structures of adult mouse ear and tail were examined 4 weeks after the first injection of Tx. The mice were housed in the animal facilities of the Laboratory Animal Resource Center at the Tokyo University of Pharmacy and Life Sciences under specific pathogen-free (SPF) conditions at constant temperature and humidity and fed a standard diet. Treatment of the mice was in accordance with the institutional guidelines of the Animal Care and Use Program of the Tokyo University of Pharmacy and Life Sciences (L18-3, L19-18, L20-5).

Establishment LLC cells expressing eGFP
LLC cells were cultured in Dulbecco’s modified Eagle’s medium (DMEM; Nakalai Tesque, Kyoto, Japan) containing 10% fetal calf serum (FCS; Invitrogen, Carlsbad, CA, USA), 1× MEM nonessential amino acids (NEAA; Sigma-Aldrich) and 100 U/mL penicillin/streptomycin (FUJIFILM Wako, Osaka, Japan) [12]. To obtain the eGFP stable transfectants, LLC cells were co-transfected with pEGFP-N1 (GenBank Accession #U55762) and CS-CDF-CG-PRE (RDB04379) including the Zeocin-resistant gene. For selection of stable transfectants, LLC cells were maintained in DMEM containing 600 mg/mL of Zeocin (Invitrogen).

In vivo lymphatic drainage of FITC-dextran
Four weeks after Tx administration, male TbβRIIfl/fl and TbβRIIfl/LEC mice (7-12 weeks old) were subcutaneously injected with 50 μL of 10 mg/mL FITC-dextran (MW 2,000,000, Sigma-Aldrich, FD2000S) into their rear footpads. Five minutes later, the transport of FITC-dextran was visualized with MVX10 fluorescence stereomicroscope (Olympus, Tokyo, Japan).

In vivo xenografts
Male TbβRIIfl/fl and TbβRIIfl/LEC mice (7-12 weeks old) were subcutaneously implanted with LLCs (2.5 × 10^5 cells) in 100 μL of PBS. Tumor volumes (V) were calculated using the following formula: length × width × width × 0.5 [13]. The tumors were surgically removed and embedded into a frozen section compound (Leica). Lymphatic metastasis was measured by the expression of eGFP when LLC-GFP cells implanted into mouse footpad were metastasized. In brief, the legs were clarified with CUBIC 1 week after the LLC-GFP transplantation [14] and tumor metastasis to the popliteal lymph nodes (PLN) was evaluated with the fluorescent stereomicroscope (Olympus).
Immunofluorescence analysis and quantification
Antibodies were obtained from the following sources: Rabbit polyclonal anti-LYVE-1 (ab14917) and anti-Ki-67 (ab15580) antibodies from Abcam (Cambridge, UK); rat monoclonal anti-PECAM-1 (550274) and anti-V-VEcadherin (550548) antibodies from BD Transduction Laboratories (Franklin Lakes, NJ); goat anti-VEGFR3 antibody (35917) from R&D systems (Minneapolis, MN). Ear skins from adult mice were dissected, fixed in 4% paraformaldehyde (PFA) in PBS for overnight at 4 °C, and washed three times with PBS. After connective tissues and hairs were removed, the samples were permeabilized in 0.1% Triton X-100 (Sigma-Aldrich) and incubated in blocking reagent (Dako, Glostrup, Denmark) before staining. The tumors and 1 cm from buttock of the tail were surgically removed and embedded into a frozen section compound (Leica). Fresh frozen sections (5 μm) were cut with a CM1850 cryostat (Leica Camera AG, Wetzlar, Germany), mounted on Cryofilm (Leica), and fixed in 100% ethanol and 4% PFA. The films were washed three times with PBS, permeabilized with 0.1% Triton X-100 for 5 min, and blocked with Blocking Reagent for 1 h at 37 °C. First antibodies in Blocking Reagent were added and incubated overnight at 4 °C. The films were washed three times with PBS and then incubated with Alexa488-conjugated donkey anti-rabbit IgG (A21206, Thermo Fisher Scientific, Waltham, MA), Alexa488-conjugated goat anti-rabbit IgG (A11008, Thermo Fisher Scientific), Alexa594-conjugated goat anti-rat IgG (A21208, Thermo Fisher Scientific) antibody, Alexa594-conjugated donkey anti-rat IgG (A21209, Thermo Fisher Scientific) antibody, or Alexa647-conjugated donkey anti-goat IgG (A11058, Thermo Fisher Scientific) at 1:200 for 1 h at room temperature [13]. After the nuclei as needed were stained with 2 μg/mL DAPI (Dojindo Laboratories, Kumamoto, Japan) for 5 min, the samples were washed three times with PBS, and the fluorescence signals were visualized by BZ-9000 fluorescence microscope (Keyence, Osaka, Japan). For mouse ear skin, the LYVE-1-positive area and the number of branches were analyzed with 12 μm² image from each of the 4 mice used. For tumor sections, the immunostaining-positive area was measured using three independent images at 10x magnification from each mouse (3 control and 6 TβRIIΔLEC mice). For tail sections, the immunostaining-positive area was measured using one image including tissues around the artery at × 10 magnification from each of the 4 mice. These analyses were carried out using BZ-X analyzer imaging software (Keyence) or ImageJ. The analysis of proliferative lymphatic endothelial cells from each mouse (n = 3) were measured the rate of Ki-67-positive cells from at least 100 VEGFR3-positive LECs per mouse.

RNA isolation and RT-PCR
Total RNA was extracted using a NucleoSpin® RNA Plus kit (Takara Bio Inc., Shiga, Japan). Reverse transcription was performed with a PrimeScript II 1st strand cDNA Synthesis Kit (Takara). qPCR was performed with a KAPA SYBR Fast qPCR kit (Nippon Genetics, Tokyo, Japan). All reactions were carried out on a LightCycler®96 (Roche). Each sample was analyzed in triplicate at least twice for each PCR measurement. Melting curves were checked to ensure specificity. Relative quantification of mRNA expression was calculated using the standard curve method with the GAPDH level. Before qPCR, the DNA fragment amplified using each primer set was detected to be a single band with the correct size by agarose gel electrophoresis. The following primer sets were used to amplify cDNAs; 5′-CAGCCCCCCTCTAA TACCCAG-3′ and 5′-AGAAGGTGTGGGCTGCT-3′ for mouse VEGF-C [15], 5′-TGCAGTGCGAAA GTGGAGATT-3′ and 5′-TGCCGTGAAATGGCC GT-3′ for mouse GAPDH [16].

Statistical analysis
Numerical results were expressed as means ± standard deviation. Significance was assessed using the unequal variances t test and the chi-square test. Probability values below 0.05 were considered significant.

Results
TGF-β regulates lymphatic network development
In mouse development, the initiation of lymphatic vessel differentiation is observed at embryonic day 10–10.5 (E10–10.5) in the anterior cardinal vein with a subpopulation of endothelial cells expressing LYVE-1, Sox18, Prox1, and VEGFR3 [6] [7, 17]. To verify the effect of TGF-β signal deficiency on active lymphangiogenesis of LECs, Tx was administered to pregnant mice 10.5 and 11.5 days after mating. Next, we analyzed the phenotype of embryos at E13.5 (Fig. 1a). Mild edema and blood clots were found on the back of TβRIIΔLEC embryos harvested from Tx-treated pregnant mice (Fig. 1b). Therefore, we visualized lymphatic vessels in the skin from their backs using the anti-LYVE-1 antibody (Fig. 1c). Interestingly, lymphatic networks with abnormally dilated lymphatic vessels were observed in the TβRIIΔLEC embryos. These results are consistent with previous findings in which TGF-β signaling acts as an active regulator of lymphangiogenesis [9].

TβRIIΔLEC mice injected with Tx showed expanded lymphatic vessel network
The deficiency of TGF-β signaling in lymphatic vessels was studied using the ear skin and tail 4 weeks after postnatal mice were administered with Tx (Fig. 2a). To visualize the lymphatic vessels of the mouse ear [18], the
lymphatic capillary maker, LYVE-1 was detected with fluorescence. Remarkable differences in the composition of the lymphatic vessel network were observed between control (TβRIIfl/fl mice treated with Tx) and TβRIIΔLEC mice (Fig. 2b). In adult TβRIIΔLEC mice, the lymphatic networks exhibited distorted structures with dilated and narrowed lymphatic vessels compared to the control mice. Their dilated structures were not as remarkable as those seen during embryogenesis. Furthermore, the LYVE-1-positive area (Fig. 2c) and the number of lymphatic branches in TβRIIΔLEC mice (Fig. 2d) were significantly increased compared to those in the control mice.

The transverse sequential sections of mouse tails were stained with antibodies against LYVE-1 and PECAM-1, which is a marker of endothelial cells, or against LYVE-1 and VE-cadherin, which is an endothelial-specific adhesion molecule. There were no differences in the vessel structure stained with anti-PECAM-1 (Fig. 2e) or anti-VE-cadherin antibodies (Fig. 2g) of mouse tails between control and TβRIIΔLEC mice, whereas the structures of the lymphatic vessels were increased in TβRIIΔLEC mice compared with the control mice (Fig. 2e). When we measured the area of LYVE-1-positive lymphatic capillary, LYVE-1-positive LECs in TβRIIΔLEC mice were significantly increased compared to those in the control mice (Fig. 2g). These results suggest that TGF-β signaling might control the number of LYVE-1-positive lymphatic endothelial cells.

TβRIIΔLEC mice treated with Tx showed expanded lymphatic vessel network

Next, we examined the lymphatic drainage function in TβRIIΔLEC mice. It is known that any substrate injected into the hind footpad is excreted in the popliteal lymph node (PLN). Thus, we injected FITC-dextran into the footpad of TβRIIΔLEC mice [19] 4 weeks after the administration of Tx (Fig. 3a). Five minutes later, FITC-dextran was found to be excreted in the PLN and reached the upper limbs of the control mice. On the other hand, less amount of FITC-dextran fluorescence could be observed in the lymphatic vessels upstream of the PLN around the thigh from TβRIIΔLEC mice (Fig. 3b). FITC-dextran administrated into footpad might not be able to drain into the lymphatic vessels beyond the PLN due to impaired lymphatic function in TβRIIΔLEC mice. We further explored lymphatic vessel dysfunction in TβRIIΔLEC mice using a tumor lymphatic metastasis model. When LLC cells that constitutively express eGFP (LLC-GFP) were inoculated into the footpad of the mice 3 weeks after the administration of Tx (Fig. 3a),
Fig. 2 Lymphatic vessels increase in TβRIIΔLEC mice. a. Experimental schedule. Mice were treated with i.p. tamoxifen (40 mg/kg) for five consecutive days and sacrificed at day 22 (3 weeks). The ears shown using the dotted square and the tail shown using the dotted line were analyzed. b Whole-mount LYVE-1 staining of ear skin from control and TβRIIΔLEC mice. Representable images are shown. Scale bar: 100 μm. c Image-based analysis of LYVE-1-positive lymphatic vessel area. The average of LYVE-1-positive area in TβRIIΔLEC mice (n = 4) was significantly higher than that in control mice (n = 4). d Number of branching points of lymphatic vessels. The average of the branching number in TβRIIΔLEC mice (n = 4) was significantly higher than that in control mice (n = 4). e, f Immunohistostaining of the transverse section of the mice tails with anti-LYVE-1 (green) and anti-PECAM-1 (red)/VE-Cadherin (red) antibodies. Sections were counterstained with DAPI (nucleus, blue). Scale bar: 100 μm. g Image-based analysis of LYVE-1-positive lymphatic vessel area of Fig. 2e from control (n = 3) and TβRIIΔLEC mice (n = 3).
metastasis to PLN was noted 1 week after the transplantation of LLC cells (Fig. 3c). Metastasis of LLC cells was observed in 85.7% of control mice, whereas it was observed in only 27.3% of TβRIIΔLEC mice treated with Tx, which significantly reduced compared to the control (P = 0.013; chi-square test) (Fig. 3d). Additionally, immunofluorescence staining was performed to confirm the presence of LLC-GFP cells in PLNs from wild-type mice (Fig. 3e). GFP-positive LLC cells were detected in the PLN stained with an anti-VEGFR3 antibody, demonstrating that lymphatic metastasis took place in the PLN. These results indicated that TGF-β signaling plays a key role in lymphatic metastasis.

**Fig. 3** TβRIIΔLEC mice exhibit decreased lymphatic drainage and tumor lymphatic metastasis. a. The mice were intraperitoneally administered Tx for five consecutive days. Three or 4 weeks later, LLC-GFP or FITC-dextran respectively were subcutaneously injected into their rear footpads independently. b. Representative images of lymph flow from the lower extremities to the popliteal lymph node (PLN). (n = 7 mice per genotype) Scale bar: 5 mm. c. Representative images of LLC-GFP metastasized to the PLN. Scale bar: 500 μm. d. Quantification of metastasis (black) and no metastasis (white) to the PLN. Scale bar: 200 μm. (control; n = 7, TβRIIΔLEC; n = 11) e. Immunohistostaining of the sagittal section of PLN from control mice with anti-VEGFR3 (red) and anti-PECAM-1 (blue) antibodies. Green fluorescence was derived from LLC-GFP. Scale bar: 100 μm.
role in the maintenance of the lymphatic drainage function and that it promotes tumor lymphatic metastasis.

**TβRIiΔLEC** mice treated with Tx showed dilated tumor lymphatic vessels

To investigate whether deletion of the **TβRII** gene in LECs affects tumor growth, **TβRIIΔLEC** mice were treated with tamoxifen consecutively for 5 days before a subcutaneous injection of $2.5 \times 10^5$ LLC cells into their dorsal region. Figure 4a shows a comparison of the tumor volume between **TβRIIΔLEC** and control mice 2 weeks after injection. There were no differences in tumor growth or tumor weight (Fig. 4b) between the two groups. However, accumulation of tissue fluid around tumors from **TβRIIΔLEC** mice could be observed although the gross morphology of tumors between two groups was quite similar (Fig. 4c). Abundant VEGFR3-positive lymphatic vessels were observed in the tumors from **TβRIIΔLEC** mice (Fig. 4d).

To determine the differences in blood and lymphatic vessel structures between control and **TβRIIΔLEC** mice, we analyzed the sections that were stained with anti-PECAM-1 and anti-VEGFR3 antibodies for blood and lymphatic vessel structures, respectively. No differences

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**Fig. 4** LEC-specific **TβRII** deletion does not affect tumor growth. **a** LLC cells were subcutaneously transplanted into Tx-treated control or **TβRIIΔLEC** mice. Tumor size was measured from above the skin at the indicated days. Tumor volumes were calculated using the formula: length $\times$ width $\times$ width $\times 0.5$. The data presented are the means $\pm$ S.D. ($n = 6$). **b** Tumor weight 15 days after LLC transplantation. ($n = 6$). **c** Appearance of xenografted tumors transplanted into control or **TβRIIΔLEC** mice 15 days after LLC transplantation. **TβRIIΔLEC** mice had tissue fluid accumulation around tumor (arrows). Representative images are shown. Scale bar: 5 mm. **d** Immunohistostaining of the tumor sections with anti-PECAM-1 (green) and anti-VEGFR3 (red) antibodies. Sections were counterstained with DAPI (nucleus, blue). Scale bar: 50 μm.
Fig. 5 (See legend on next page.)

(a) PECAM-1 positive area (µm²) for Control and TβRII^ΔLEC.

(b) VEGFR3 positive area (µm²) for Control and TβRII^ΔLEC.

(c) Lymphatic lumen area (µm²) for Control and TβRII^ΔLEC.

(d) Ki-67/PECAM-1/VEGFR3 images for Control and TβRII^ΔLEC.

(e) Ki-67/VEGFR3^+ cells (%) for Control and TβRII^ΔLEC.

(f) Relative VEGF-C expression/GAPDH for Control and TβRII^ΔLEC.
in PECAM-1-positive area between control and TβRIIΔLEC mice were observed in blood vessel structures (Fig. 5a), whereas the VEGFR3-positive areas were remarkably reduced in TβRIIΔLEC mice (Fig. 5b). Additionally, lymphatic lumens in TβRIIΔLEC mice were larger than those in the control mice (Fig. 5c), indicating upregulation of lymphangiogenesis. Inhibition of TGF-β signaling has been reported to promote lymphangiogenesis and LEC proliferation in the presence of VEGF-C. Therefore, we investigated endothelial cell growth and VEGF-C expression in the tumor tissue. Immunofluorescent staining of tumor tissues showed that Ki-67-positive lymphatic endothelial cell in embryos from TβRIIΔLEC mice caused a marked decrease in lymphatic vessel structure. In the mouse model for chronic peritonitis, the TGF-β kinase inhibitor increased LYVE-1-positive area [9]. They also showed lymphangiogenesis could be enhanced by VEGF-C secreted from inflammatory macrophages. The tumors from TβRIIΔLEC mice also increased the area of VEGFR3-positive lymphatic lumen. This phenomenon might be due to the fact that LECs escaping from TGF-β signal proliferation in the presence of VEGF-C from the tumor microenvironment (Fig. 5e) [22]. Since the role of TGF-β signaling in LECs seems to be context-dependent, it would be necessary to analyze the crosstalk of TGF-β signal with other signaling pathways in future. In this study, we used anti-LYVE-1 and anti-VEGFR3 antibodies to detect LECs (Figs. 1c and 2b, e), although there are several available antibodies that recognize lymphatic vessels. The anti-LYVE-1 antibody is useful to detect lymphatic vessel structures in fetal skin and tissue from adult mice. However, the anti-LYVE-1 antibody is not available in some cases because it can also recognize embryonic hematopoietic stem cells [23] and macrophages [24] in tumor microenvironment. Thus, we used the anti-VEGFR3 antibody in tumor tissues (Figs. 4d and 5d, Supplemental Figure 1). Podoplanin is another lymphatic specific maker [25], but it is also used as a cancer biomarker [26] (Supplementary Figure 1).

Cancer metastasis responsible for the death of patients with cancer is a hallmark of malignant tumors [27][28]. Since cancer cells metastasize to distant organs through blood and lymphatic vessels, it is very important to understand how tumor angiogenesis and lymphangiogenesis are regulated in the human body. Tumor lymphatic vessels connect primary tumor cells and lymph nodes. Consequently, cancer cells invade the lymph nodes to move to other organs [29]. In the present study, we found that TGF-β signaling is involved in lymphatic drainage, in addition to confirming structural abnormalities of lymphatic vessels by the loss of LEC-specific TGF-β signaling. The quantitative analysis to evaluate the effect of TGF-β signaling on lymphatic flow might be needed [30], but it is interesting that deletion of TGF-β signaling suppressed tumor lymphatic metastasis by reducing lymphatic drainage [31] (Fig. 3). TGF-β is abundant in the tumor microenvironment and enhances motility and metastasis of cancer cells.
Our results suggest that the abundant TGF-β may act on lymphatic endothelial cells to promote tumor metastasis. Thus, the inhibition of TGF-β signaling by small compounds, antibodies, Fc-chimeric receptors [32], or RNA interference may block tumor metastasis via lymphatic vessels [33] [34].

Conclusions
LEC-specific TβRII knockout mice showed that TGF-β signaling promotes lymphatic drainage. Furthermore, TGF-β enhances tumor metastasis via lymphatic vessels. Therefore, blockade of TGF-β signaling might inhibit tumor metastasis targeting the lymphatic vessels.

Abbreviations
ALK: Activin-like kinase; cdNA: Complementary deoxyribonucleic acid; DMEM: Dulbecco's modified Eagle's medium; GFP: Green fluorescent protein; FCS: Fetal calf serum; FITC-dextran: Fluorescein isothiocyanate-dextran; LEC: Lymphatic endothelial cell; TGF-β: Transforming growth factor β; VEGF: Vascular endothelial growth factor; VEGFR: Vascular endothelial growth factor receptor

Supplementary Information
The online version contains supplementary material available at https://doi.org/10.1186/s41232-021-00185-4.

Additional file 1: Supplementary Figure 1. Antibodies that recognize LECs in tumor tissue. LLC cells were subcutaneously transplanted into Tx-13IF/F mice, and 15 days after LLC injection, fresh frozen sections were prepared. Immunohistostaining with anti-podoplanin (DPDPN, green), anti-LYVE-1 (red) and anti-VEGFR3 (blue) was performed. Scale bar: 100 μm.

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Authors' contributions
KF, KH, KL, MH, TT, and FF conducted the in vivo and in vitro experiments and analyzed the data. MH, SI and TW took part in the interpretation of the data and discussion of the paper. FF and SI wrote the manuscript. FF, MH, SI, and TW revised the manuscript. All authors have read and approved the manuscript.

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Availability of data and materials
All data generated and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations
Ethics approval and consent to participate
All experimental procedures were performed in accordance with the institutional guidelines of the Animal Care and Use Program of the Tokyo University of Pharmacy and Life Sciences (L18-3, L19-18, L20-5).

Competing interests
The authors declare that they have no competing interests.

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