

ROLE of TGF- β SIGNALING IN THE PATHOGENESIS OF VASCULAR DISEASES

By

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To my loving wife, parents, and brother

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Abstract of Dissertation Presented to the Graduate School
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Genetic mutations causing diverse genetic vascular diseases have been found in the genes that encode components of the transforming growth factor- β (TGF- β) signaling pathway, suggesting that TGF- β signaling is essential for homeostasis of the vascular system. To explore the role of TGF- β signaling in the pathogenesis of a couple of vascular diseases, our laboratory developed genetic mouse models for two major vascular diseases, pulmonary arterial hypertension (PAH) and hereditary hemorrhagic telangiectasia (HHT). PAH is a rare but fatal disease that increases pulmonary pressure due to thickening of pulmonary arterial walls. Although deficiency of bone morphogenetic protein type II receptor (BMP2) in TGF- β signaling is known as a genetic contributor, whether dysfunction of SMAD1, one of the canonical transducers of BMP2 is critical for PAH development remains unknown. We hypothesized that deficiency of SMAD1 would lead to PAH and tested the hypothesis using *Smad1*-conditional knockout (cKO) mice. The *Smad1* gene was deleted in endothelial cells or smooth muscle cells using L1cre and *Tagln*-cre lines respectively. We discovered that these *Smad1*-cKO mice develop PAH, suggesting that SMAD1 could be a critical downstream mediator of BMP2 in PAH. HHT2 is a vascular disease with arteriovenous

malformations (AVMs) and visceral hemorrhages due to *ALK1* deficiency. Our previous studies showed that wounding plays an essential role in AVM development in the *Alk1*-deficient context. We investigated the involvement of two major wound-healing responses: inflammation and angiogenesis. We found that either LPS (for inflammation) or VEGF (for angiogenesis) could recapitulate the wound-induced AVM formation in *Alk1*-deficient skin. The inhibition of angiogenesis with VEGF-neutralizing antibody (Ab) significantly inhibited both LPS- and wound-induced AVMs and ameliorated internal bleedings in *Alk1*-deficient mice, suggesting a critical role for angiogenic stimulation in AVM development. These data provide a better scientific basis for the therapeutic effect of a VEGF blockade (Bevacizumab) for epistaxis, GI bleeding, and liver AVMs in HHT patients. Our data also demonstrated that the skin AVM model by *Alk1*-conditional knockout mice is reliable for preclinical screening of drug candidates for epistaxis and GI bleedings.

CHAPTER 1
SMAD1-DEFICIENCY IN EITHER ENDOTHELIAL OR SMOOTH MUSCLE CELLS
RESULTS IN PULMONARY ARTERIAL HYPERTENSION

Introduction

Pulmonary Arterial Hypertension (PAH)

Pulmonary Arterial Hypertension (PAH) (formerly “primary pulmonary hypertension-PPH”) is rare but fatal lung vascular disease and is characterized by sustained elevation of mean pulmonary arterial pressure (PAP) and increased pulmonary vascular resistance leading to right heart failure. PAH is one of 5 types of pulmonary hypertension (PH), which is caused by chronic thrombosis, embolization, elevated left ventricular end-diastolic pressure, lung disease/hypoxemia or valve disease. While PH is diagnosed by the sole criterion being a resting mean PAP > 25 mmHg, diagnosis of PAH is required two additional criteria: more than 3 Wood units of pulmonary vascular resistance (PVR), and less than 15 mmHg of pulmonary capillary wedge pressure without other causes of PH. PAH is subclassified into idiopathic PAH (IPAH), heritable PAH (HPAH), and PAH associated with other diseases (APAH). Other diseases include congenital heart defect, portal hypertension, HIV infection, connective tissue disease, appetite suppressant drug use.¹

Pathology of Pulmonary Arterial Hypertension

The basic pathological phenotype of PAH is a narrowing and thickening of small pulmonary vessels. All PAH patients exhibit pulmonary vascular remodeling of all layers of the vessel: intimal thickening, smooth muscle cell hypertrophy, adventitial fibrosis and occluded vessels by *in situ* thrombosis.² The intimal layer is a single-layered lining of endothelial cells (ECs) between the internal elastic lamina and lumen. Normally, ECs are quiescent but in PAH lungs, they are activated, rapidly proliferating and form

neointima layers. Myofibroblast is one of major cell types in the neointima and expresses smooth muscle cell (SMC) markers such as smooth muscle α -actin (α SMA) and vimentin instead of EC makers such as CD31 and the von Willebrand factor (vWF).³ Although the origin of the myofibroblasts remains unknown, *in vitro* data suggest that myofibroblasts are likely originated from vascular SMC or transdifferentiated from EC.^{4, 5} Plexiform lesions are commonly observed in the severe form of PAH showing multiple capillary-like channels in a pulmonary artery.⁶ They contain several cell types including ECs, myofibroblast, and connective tissues.⁷ ECs are the most responsible cell types for initiation of plexiform lesions through monoclonal proliferation of tumorlet-like clusters of ECs. A study showed that ECs with *BMPR2* mutation are more susceptible to apoptotic stimulus and after repeated apoptosis, the surviving ECs are apoptosis-resistant and undergo considerable proliferation and develop plexiform lesions.⁸ Vascular SMCs are a predominant cell type of the medial layer and normally unresponsive to mitogens.⁹ Hyperplastic SMCs are a common phenotype in all the different forms of PAH. In precapillary vessels, cells inside internal elastic lamina appear to be involved in differentiation into SMCs¹⁰ and in distal vessels lacking elastic lamina, pericyte and interstitial fibroblast surrounding lung parenchyma seem to contribute to muscularization.¹¹ Remodeling of adventitia layer by increase of fibroblasts is associated with induction of many proinflammatory cytokines including monocyte chemoattractant protein (MCP)-1, macrophage inflammatory protein (MIP)-2, interleukin (IL)-1 β , IL-6.¹² Endothelial level of prostacyclin (prostaglandin-I₂), an endogenous vasodilator, an inhibitor of platelet aggregation and a suppressor of vascular SMCs proliferation, was shown to be decreased in patients with PAH.¹³ The mechanism

involved in this observation could be that lowered expression of prostacyclin synthase impairs balance between prostacyclin and thromboxane A₂, a vasoconstrictor and a potent stimulator of platelet aggregation.¹⁴ Thrombotic lesions in PAH are associated with imbalanced ratio of prostacyclin and thromboxane A₂ and potential outcome of the serotonin pathway.^{15, 16}

Current Treatment Options for PAH

The incidence of PAH is 2-3 cases per million people every year.¹⁷ There is age variability for the initial onset of PAH.^{18, 19} Without proper treatments, mortality is predicted within 3 years after diagnosis. Up to now, the following treatments are available and have been used widely.

Prostacyclin (prostaglandin I₂)

One of the most extensively and successfully used therapies for PAH patients is to increase the level of prostacyclin with exogenous prostanoids. Fatty acid cyclooxygenase metabolizes arachidonic acid to prostaglandin H₂, a substrate for both prostacyclin (prostaglandin I₂) synthase and thromboxane synthase. Prostacyclin is expressed by prostacyclin synthase in endothelial cells and works as a vasodilator through stimulation of cyclic AMP (cAMP) and inhibits proliferation of SMCs.²⁰ On the other hand, thromboxane A₂ (TxA₂) is produced by thromboxane synthase in platelet and endothelial cells and stimulates vasoconstriction and platelet aggregation. PAH patients showed decreased prostacyclin metabolites and increase of TxA₂ production,¹⁴ indicating that endothelial dysfunction and platelet activation in PAH might impair the balance between vasodilators and vasoconstrictors. Continuous intravenous infusion of epoprostenol (Floran) started in the early 1980s, has shown to decrease pulmonary vascular resistance (PVR), increased cardiac output, and improved exercise capacity

and overall survival rates.²¹⁻²³ Due to poor stability, high expense, and side effects from intravenous treatment of epoprostenol,²⁴ more stable analogs and alternative delivery of prostacyclin were developed: subcutaneous (treprostinil), oral (beraprost), or inhaled (iloprost) delivery of prostacyclin analogs.²⁵⁻²⁷ Recent combination therapy of prostanoid and a PDE5 inhibitor was effective in improving pulmonary hemodynamic change and exercise tolerance in PAH.^{28, 29}

Endothelin-1 receptor antagonists

Endothelin-1 (ET-1) works as a vasoconstrictor via 2 types of receptors (ETA and ETB) and has a mitogenic effect on SMCs.^{30, 31} Both receptors are expressed in SMCs and mediate vasoconstriction while ETB in ECs promotes vasodilation through NO and prostacyclin production.^{32, 33} PAH patients exhibited high levels of lung and circulating ET-1.^{34, 35} Endothelin receptor antagonist such as Bosentan (a nonselective ET receptor antagonist), sitaxsentan and ambrisentan (selective ETA receptor blocker) were FDA-approved and showed significant but moderate improvement in pulmonary hemodynamics and 6-minute walk distance.³⁶⁻³⁹ However, there is no clear evidence showing advantage of selective ETA antagonism over combined antagonism for both ETA and ETB. Liver toxicity and teratogenicity are classic side effects. Despite a report that bosentan monotherapy increased survival,⁴⁰ more robust data for survival are necessary for clinical trials. Since the endothelins are produced from pro-endothelin by endothelin-converting enzyme, inhibitors for endothelin-converting enzyme could be an alternative approach to blocking overproduction of endothelin in PAH. Studies with this inhibitor (eg, daglutril) have been conducted in systemic hypertension and heart failure⁴¹ and application for PAH patients are being investigated.

Nitric oxide (NO) and cyclic GMP (cGMP)

NO produced by endothelial NO synthase (eNOS) dilates blood vessel in the pulmonary circulation. Endothelium-derived NO stimulates soluble guanylate cyclase (sGC) to produce intracellular cGMP in SMCs.⁴² Increased cGMP activates cGMP-dependent protein kinase (cGKI) and decreases the sensitivity of myosin to calcium-induced contraction and lowers concentration of calcium released from the sarcoplasmic reticulum.⁴² Phosphodiesterase type 5 (PDE5), the target of sildenafil opposes NO-dependent vessel dilation by suppressing the rise of NO-induced cGMP.⁴³ NOS expression and NO bioavailability were shown to be reduced in the lungs of PAH patients.⁴⁴⁻⁴⁶ Furthermore, PDE5 is upregulated in the hypoxia-mediated animal model for PAH, which worsens cGMP availability.^{47,48} Endogenous NOS inhibitors, asymmetrical and symmetrical dimethylarginines (ADMA and SDMA) appeared to be more abundant in PAH.^{49, 50} Thus, Inhaled NO gas dilates pulmonary arteries, which lowers vascular resistance, PA pressure, and RV afterload.^{51, 52} The oral delivery of sildenafil citrate, an inhibitor of PDE5, improved exercise capacity and hemodynamics in PAH patients.⁵³

Calcium channel blocker

Intracellular calcium and calcium-mediated signals play a crucial role in smooth muscle contraction.^{54, 55} A study has shown that a high dose of calcium channel blockers have a beneficial effect on the survival of some PAH patients.⁵⁶ However, it should be used very carefully because calcium channel blockers are only convincingly effective in the 5% of patients who showed an acute vasoactive response to vasoreactivity testing.^{56, 57}

In summary, current treatments for this disease alleviate symptoms and improve heart functions, but are not a cure of the disease. Better understanding on the pathogenetic mechanisms of disease would allow development of drugs targeting the cure.

Pathologic Mechanism of PAH

Involvement of BMP signaling based on genetic studies

Genetic studies have found the linkage of a locus for the gene, named PPH1, to chromosome 2q31–32 in 1997^{58, 59} and later showed that Bone Morphogenetic Protein type 2 Receptor (BMP2), one of the receptors in transforming growth factor β (TGF- β) superfamily signaling, is responsible for heritable PAH in an autosomal dominant manner.⁶⁰⁻⁶³ A heterozygous BMP2 mutation was found in nearly 70% of HPAH patients and also in 25% of sporadic IPAH patients.^{63, 65-67} The levels of *BMP2* mRNA and proteins are markedly reduced in the lungs of PAH patients with heterozygous *BMP2* mutations,⁶⁴ indicating that BMP2 mutations are associated with haploinsufficiency. Penetrance of disease is low and disease expressivity varies even within members of a family. Estimates showed that only about 20% of individuals with *BMP2* mutation develops PAH during their entire life. This low penetrance in PAH suggests that additional factors such as inflammation may be necessary for clinical manifestations of PAH in addition to the genetic predisposition. While *Bmpr2*^{+/-} mice displayed mild PAH phenotype, adenovirus-mediated pulmonary overexpression of 5-lipoxygenase (5-LO), a mediator of inflammation, or a chronic infusion of serotonin (5HT) developed full-blown PAH in *Bmpr2*^{+/-} mice.^{68, 69} Interestingly, BMP2 is downregulated in the lung tissues and cells from idiopathic PAH patients without *Bmpr2* mutation,^{64, 70} implying a wide-range of influence of BMP2 deficiency to other forms of

PAH. Furthermore, BMPR2 signaling plays an important role in the proliferation of local endothelial cells and the migration and local proliferation of smooth muscle cells.^{8, 71}

Taken together, impaired BMP signaling due to BMPR2 deficiency would be a considerable contributor for PAH development.

TGF- β superfamily signaling

TGF- β is a large cytokine family that contributes to diverse cellular processes including cell proliferation, migration, apoptosis, pattern formation, and immunosuppression. TGF- β family members include TGF- β s, bone morphogenetic proteins (BMPs), growth and differentiation factors (GDFs), activins/inhibins, and müllerian inhibiting substance (MIS)/ anti-Müllerian hormone (AMH).⁷²⁻⁷⁴ TGF- β signal transduction is initiated by binding of ligands to heteromeric complex of transmembrane serine/threonine type 2 and type 1 receptors. Once ligands bind to a type 2 receptors, the ligand-type 2 receptor complex recruits and trans-phosphorylates type 1 receptor, which, in turn, activates receptor-regulated SMADs (R-SMADs): SMAD2/3 for TGF- β s and SMAD1/5/8 for BMPs. R-SMADs then form a complex with a common partner, SMAD4 (Co-SMAD) and enter the nucleus and initiate transcription of target genes. On the other hand, there is mounting evidence demonstrating that independently of SMADs, TGF- β signaling can be transduced through mediators other than SMADs such as the mitogen-activated protein kinases (MAPKs), including p38MAPK, p42/44MAPK (ERK1/2), and c-Jun-N-terminal kinase/stress-activated protein kinase (JNK/SAPK).^{75, 76} For instance, exogenous BMP ligands stimulate phosphorylation of p38MAPK and p42/44MAPK, and affect proliferation of SMCs.^{70, 77}

Endothelial dysfunction in PAH

The endothelium is the cells lining the interior surface of blood vessels in the entire circulatory system. Endothelial cells function in many aspects of vascular biology: vasoconstriction and vasodilation, and hence the control of blood pressure, blood clotting (thrombosis & fibrinolysis), atherosclerosis, formation of new blood vessels (angiogenesis), inflammation and swelling (oedema) and also control of the passages of materials and blood cells (permeability). Thus, endothelial dysfunction may result in increased coagulation, proliferation, and vasoconstriction.⁷⁸⁻⁸¹ Endothelial dysfunction seems to play an integral role in mediating the structural changes in the pulmonary vasculature. Disordered endothelial cell proliferation along with concurrent neoangiogenesis results in the formation of glomeruloid structures known as the plexiform lesions, which were found in the pulmonary vessels of patients with severe PAH.⁸² *In vitro* studies with endothelial cells suggest a plausible mechanism of angioproliferative change of ECs in PAH. Loss of BMPR2 function cannot protect ECs from apoptosis, representing a possible initiating step and increased apoptosis of ECs promoted appearance of the apoptosis-resistant cells.^{8, 83} In addition, an altered production of various endothelial vasoactive mediators such as Nitric oxide (NO), prostacyclin, endothelin-1 (ET-1), serotonin, and thromboxane, has been increasingly recognized in patients with PAH.^{14, 34, 84-86} Since change of these mediators can affect the growth of the smooth muscle cells, alteration in their production may facilitate the development of pulmonary vascular hypertrophy. Thus, it is conceivable that the beneficial effects of currently available treatments for PAH, such as prostacyclin, NO, and ET antagonists, result in part from restoring the balance between these mediators. Furthermore, endothelial dysfunction may bring about a change in the EC permeability

and allow direct contact of serum proliferative mediators with the subendothelium, leading to cell proliferation in the medial and adventitial layers.⁸⁷⁻⁸⁹

Compromised immune response and inflammation

Pulmonary hypertension has been associated with connective tissue disease,⁹⁰ human immunodeficiency virus (HIV),⁹¹ and auto-antibodies.⁹² Mononuclear cell infiltration was often observed in the PAH vascular lesions.⁶⁴ Inflammatory cells including macrophages and lymphocytes are highly accumulated in the plexiform lesions of hypertensive pulmonary vessels.⁷ Induction of many proinflammatory cytokines and chemokines, such as monocyte chemoattractant protein (MCP)-1, macrophage inflammatory protein (MIP)-1 α , interleukin (IL)-1 β , IL-6, RANTES, fractalkine has been implicated in PAH.^{12, 93-97} In the classical PH animal model, Monocrotalline (MCT, a pyrrolizidine alkaloid plant toxin)-injected rats elevated pulmonary pressure with apoptosis of ECs and SMCs and massive mononuclear infiltration into the perivascular regions of pulmonary arterioles.^{98, 99} Therefore, the observations from a variety of human and animal studies suggest a compelling conclusion that intact immune system is required to maintain homeostasis of pulmonary circulation. In other words, dysregulated immunity could be an environmental second hit for clinical manifestation of PAH in individuals with *Bmpr2* mutation.

Role of TGF- β /BMP Balance in Pulmonary Vascular Homeostasis

Only 20% disease penetrance in individuals with *Bmpr2* mutation implies that the *Bmpr2* mutation is necessary but not sufficient alone for clinical manifestation of PAH. This fact suggests that other genetic or environmental modifiers are necessary for initiation and progression of PAH. TGF- β could be a possible candidate of modifiers in PAH.

BMP and TGF- β signaling form an opposing balance in pulmonary vascular homeostasis.¹⁰⁰ Thus, BMPR2 deficiency may result in predominant TGF- β signaling which favors fibrosis and smooth muscle cell growth. Recent studies suggest that an imbalance between TGF- β and BMP signaling leads to PAH development. Activation of BMP signaling suppresses proliferation of pulmonary vascular smooth muscle cells.⁶⁴ Furthermore, TGF- β polymorphisms that increase TGF- β activity are associated with early onset of PAH and increased penetrance of heritable PAH.¹⁰¹ Zaiman *et al.* reported that increased TGF- β signaling participates in the pathogenesis of monocrotaline-induced PH.¹⁰² Long *et al.* showed that suppression of TGF- β signaling via activin receptor-like kinase-5 inhibitor prevents development and progression of PAH in the monocrotaline model.¹⁰³ Impaired BMP signaling from BMPR2 deficiency would not be sufficient to develop PAH and when in conjunction with predominance of TGF- β , finally would elevate pulmonary pressure.

Is SMAD Important Signaling Mediator for PAH Caused by BMPR2-Deficiency?

Human genetic studies have identified various *BMPR2* mutations throughout the exons coding for the BMPR2 protein.¹⁰⁴ These data imply that deficiency of BMPR2 is a crucial genetic factor in the PAH development. However, which downstream signaling molecules of BMPR2 such as MAPKs or SMADs contribute to PAH is still unknown. Mutations in the kinase domain (about 50% of total mutations) suggested that downstream SMAD signaling play a role for PAH development because of its reduced SMAD-dependent transcriptional activity.¹⁰⁵ However, mutations in cytoplasmic tail domain (about 20% of total mutations) such as R899X leave SMAD signaling intact, indicating that the cytoplasmic tail of BMPR2 may not be essential for transduction of BMP signals through SMADs.^{106, 107} In other words, SMAD-deficiency may not be

associated with the pathogenesis of PAH. James West *et al.* addressed this question using transgenic mice expressing BMPR2^{R899X} transgene in smooth muscle.¹⁰⁸ In these mice, truncated BMPR2 proteins with a tail domain mutation were overexpressed in smooth muscle cells by doxycycline treatment from 4 weeks of age. They elevated right ventricular systolic pressure, associated with extensive pruning, muscularization of pulmonary arterioles and perivascular infiltration of immune cells, not affecting SMAD activity. These *in vivo* results suggest that the SMAD-deficiency may not be associated with PAH caused by BMPR2-deficiency.

Endothelial BMPR2-Deficient PAH Mouse Model

Previously we produced *Bmpr2*-conditional knockout (cKO) mice [L1cre(+);*Bmpr2*^{fl/fl} or ^{+/fl}] by deleting the *Bmpr2* gene in the pulmonary endothelium using the novel L1cre line.¹⁰⁹ A subset of mice exhibited elevated right ventricular systolic pressure (RVSP) and right ventricular hypertrophy (RVH) which are the representative phenotypes of pulmonary hypertensive (PH) mice¹¹⁰ and these PH mice showed increased muscularization of pulmonary arterioles and thickening of vessel wall compared to non-PH mice and controls.

Deletion of *Smad1* Gene in ECs or SMCs

SMAD1 is a canonical signal transducer of BMPR2 and its reduced activity has been associated with PAH. Yang *et al.* showed that phosphorylation of SMAD1 was reduced in the pulmonary arterial SMCs of PAH patients with *BMPR2* mutation.⁷⁰ Thus, we hypothesized that SMAD1 is a contributing downstream mediator of BMPR2 in the pathogenesis of PAH. To test this hypothesis, we produced *Smad1* cKO mice by deleting *Smad1* gene in the pulmonary endothelial cells (pECs) or smooth muscle cells (SMCs) using L1cre and *Tagln*-cre lines, respectively and found that *Smad1* deletion in

pECs or SMCs can develop PAH in mice with muscularization of pulmonary vessels, suggesting that SMAD1 may be a critical downstream molecule in PAH. Recent studies reported that enhanced TGF- β activity is associated with PAH pathology as well as impaired BMP signaling.^{103, 111-113} In *Bmpr2*-deleted pulmonary endothelial cells, we found that a reduced BMP response resulted in a predominant TGF- β response, supporting the role of a balance between TGF- β and BMP signaling in the homeostasis of pulmonary blood pressure.¹⁰⁰

Results

SMAD1/5/8 Phosphorylation was Impaired in the Lungs of Pulmonary Hypertensive L1cre(+);*Bmpr2*^{2f/2f} Mice.

We have previously shown that about 40% of mice in which *Bmpr2* gene was specifically deleted in endothelial cells by L1cre exhibited pulmonary hypertensive phenotype.¹¹⁰ To assess the extent to which the conditional *Bmpr2* deletion affects on its downstream SMAD1/5/8 signaling, lungs of cre-negative controls, Non-PH L1cre(+);*Bmpr2*^{2f/2f}, PH L1cre(+);*Bmpr2*^{2f/2f} mice were immunostained with anti-phospho-SMAD1/5/8 antibodies (Fig. 1-1A through 1-1F). While pSMAD1/5/8 positive cells were readily detected in the cre-negative controls (Fig. 1-1A through 1-1B) and Non-PH *Bmpr2* mutants (Fig. 1-1C), they appeared to be much less in the PH *Bmpr2* mutants (Fig. 1-1D through 1-1F). Consistent with this immunostaining result, the levels of SMAD1/5/8 phosphorylation in the whole lung samples were significantly reduced in the PH *Bmpr2* mutants compared to the cre-negative controls and Non-PH *Bmpr2* mutants (Fig. 1-1G, H). These data suggested to us that impaired SMAD1 signaling may play a pivotal role in the pathogenesis of pulmonary arterial hypertension associated with *Bmpr2* deficiency.

***Smad1* Deletion in Pulmonary ECs or SMCs by L1cre or *Tagln*-cre**

To investigate the role of SMAD1 in the pathogenesis of PAH, we exploited conditional knockout approaches for deleting the *Smad1* gene in endothelial cells or smooth muscle cells by L1cre or *Tagln*-cre lines, respectively, because *Smad1*-null mice were embryonic lethal.¹¹⁴ Both L1cre(+);*Smad1*^{fl/fl} and *Tagln*-cre(+);*Smad1*^{2fl/2fl} mice were viable and normal compared to their cre-negative littermates. We analyzed the Cre-activity by detecting the *Smad1*-null allele in several organs of 2 month-old mice, including the lung, heart, liver, kidney, and spleen. Consistent with our expectations based on our previous reports,^{109, 110} the Cre-mediated *Smad1* deletion was detected primarily in the lungs of L1cre(+);*Smad1*^{fl/fl} mice, and found in most organs of *Tagln*-cre(+); *Smad1*^{fl/fl} mice (Fig 1-2).

Some Mice with the *Smad1* Deletion in Pulmonary ECs or SMCs Exhibited Elevated Pulmonary Pressure and Right Ventricular Hypertrophy.

To assess the pulmonary pressure, we measured the right ventricular systolic pressure (RVSP) of L1cre(+);*Smad1*^{fl/fl}, *Tagln*-cre(+);*Smad1*^{fl/fl}, and their age-matched cre(-) control mice. While RVSPs of controls were clustered in a 20-27 mmHg range, those of L1cre(+);*Smad1*^{fl/fl} and *Tagln*-cre(+);*Smad1*^{fl/fl} mice were scattered in a wide range (Fig. 2A) from 22 to 45 mmHg. About 40% (14/35) of L1cre (+);*Smad1*^{fl/fl} mice and 12% (4/33) of *Tagln*-cre(+);*Smad1*^{fl/fl} mice had their RVSPs greater than 30 mmHg, and we designated them as pulmonary hypertensive (PH) group (Fig. 1-3A). The mean RVSPs of L1cre(+);*Smad1*^{fl/fl} (28.1 mmHg) was significantly higher than that of cre(-);*Smad1*^{2fl/2fl} (23.9 mmHg). Fulton index, the ratio of RV free wall weight over septum plus left ventricular free wall weight, was used to estimate RV hypertrophy. The Fulton index of L1cre(+);*Smad1*^{fl/fl} and *Tagln*-cre(+);*Smad1*^{fl/fl} mice was significantly greater

than that of *cre(-);Smad1^{ff}* controls (Fig. 1-3B). It was greater in the PH mice than that in N-PH mice, indicating that sustained elevation of pulmonary pressure might have resulted in RV hypertrophy in the PH groups (Fig. 1-3C). There was no difference in systemic blood pressure among three groups (Fig. 1-3D).

The Number of α SMA-Positive Distal Arteries and Medial Wall Thickness Were Increased in the *Smad1* Mutant Mice.

To examine whether the elevated RVSP and RV hypertrophy in the *Smad1* mutants is associated with pulmonary vascular remodeling, anti-smooth muscle α -actin (α SMA)-positive pulmonary arteries ranging from 30-70 μ m in diameter were counted and the wall thickness was measured. The PH group *L1cre(+);Smad1^{ff}* mice showed higher number of α SMA-positive pulmonary arterioles and thicker arterial walls compared to the N-PH group and the *cre(-);Smad1^{ff}* (Fig. 1-4B, E, F). In *Tagln-cre(+);Smad1^{ff}* mice, however, both PH and N-PH groups showed a higher number of muscularized vessels and thicker walls compared to the *cre(-);Smad1^{ff}* controls.

Isolation of Pulmonary Endothelial Cells Carrying *R26^{creER/+};Bmpr2^{2f/2f}* Allele

It has been hypothesized that an opposing balance between TGF- β and BMP signalings is critical for homeostasis of pulmonary vasculature, and that imbalance of TGF- β /BMP signalings may contribute to the pathogenesis of PAH.¹⁰⁰ In order to investigate this hypothesis and to examine the extent to which *Bmpr2* deficiency impact on this balance, we established an in vitro model as follows. We isolated pulmonary ECs (pECs) from the lung of *R26^{creER/+};Bmpr2^{2f/2f}* mouse, in which the conditional “2f” allele can be converted to null “1f” allele by tamoxifen treatment that activates the Cre recombinase function. Three days of culture of the pECs with medium containing 1 μ M 4-hydroxy-tamoxifen (OH-TM) efficiently deleted exons 4 and 5 of the *Bmpr2* gene (Fig.

1-5A). When the *Bmpr2*-deleted cells were subsequently cultured with OH-TM-free media for 10 days, no overgrowth of undeleted cell populations were observed (Fig. 1-5B). Henceforth, *Bmpr2*-intact and -deleted cells will be designated as *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells, respectively.

To examine whether OH-TM treatment and *Bmpr2* deletion affected the expression of other genes involved in TGF- β pathways or altered general EC characteristics, semi-quantitative RT-PCR analyses were performed. Expression of EC-specific markers including *Nos3*, *Tie2*, *Eng*, and *Flk1* were maintained in *Bmpr2*^{1f/1f} pECs (Fig. 1-5D). *Bmpr2* transcript level was undetectable while transcripts for other TGF- β /BMP receptors were unchanged in *Bmpr2*^{1f/1f} pECs (Fig. 1-5C).

Deletion of *Bmpr2* Gene and Impaired BMP Signaling

In order to assess the extent to which *Bmpr2*-deletion affects BMP signaling, BMP4 or BMP7 was added to *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells at various doses (0, 5, 25, 50 ng/ml) after 15 hr serum starvation. First, we examined the protein level of BMPR2 in these cells. While BMPR2 was readily detected in *Bmpr2*^{2f/2f} cells, BMPR2 protein was undetectable in *Bmpr2*^{1f/1f} cells (Fig 1-6A, B). Level of phosphorylation of SMAD1/5/8 was augmented in a dose-dependent manner in *Bmpr2*^{2f/2f} cells whereas it was suppressed at all BMP4 doses in *Bmpr2*^{1f/1f} cells (Fig. 1-6A, C), indicating that the BMP4 signaling is impaired in *Bmpr2*^{1f/1f} pECs. BMP7 signaling can be compensated by ACVR2A in BMPR2-depleted pulmonary artery SMCs (PASMCs).⁷⁷ As shown in Fig. 1-6D, phosphorylation of SMAD1/5/8 was elevated in a dose-dependent manner in both *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells, suggesting that BMP7 signaling can be compensated in *Bmpr2*^{1f/1f} pECs.

Enhanced TGF- β Signaling in BMPR2-Deficient pECs

To investigate whether impaired BMP signaling affects TGF- β signaling, we first examined the basal level of SMAD2 phosphorylation in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells cultured in medium containing 10% fetal bovine serum (Fig. 1-7A). The level of SMAD2 phosphorylated in *Bmpr2*^{1f/1f} cells was much higher than that in *Bmpr2*^{2f/2f} cells. To test if *Bmpr2*-deficient cells are more sensitive to TGF- β , we examined SMAD2 phosphorylation as a response to TGF- β 1 (0, 1, and 2 ng/ml) (Fig. 1-7B). While both *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells showed a dose-dependent augmentation of pSMAD2, the level of SMAD2 phosphorylation was significantly higher in *Bmpr2*^{1f/1f} cells at 1 and 2 ng/ml TGF- β 1 compared to *Bmpr2*^{2f/2f} cells, suggesting that impaired BMP signaling may potentiate the TGF- β signaling.

Opposing Balance Between TGF- β and BMP Signalings in pECs

To investigate whether BMP and TGF- β signaling form an opposing balance in pECs, we examined whether TGF- β -induced SMAD2 phosphorylation is suppressed by BMP treatment in pECs. TGF- β 1 (0, 0.1, 1, and 2 ng/ml) and BMP4 (25 ng/ml) or BMP7 (25 ng/ml) were treated for 30 min after serum starvation in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells. The level of SMAD2 phosphorylation by 1 and 2 ng/ml of TGF- β 1 treatment in *Bmpr2*^{2f/2f} cells was decreased by either BMP4 or BMP7, implying that BMP and TGF- β have a competitive relationship by opposing each other (Fig. 1-8). However, TGF- β -mediated SMAD2 phosphorylation in *Bmpr2*^{1f/1f} cells was not affected by BMP4 treatment, suggesting that BMP4 signaling is mediated mainly through BMPR2 (Fig. 1-8A). Interestingly, BMP7 could suppress overactivated TGF- β signaling in *Bmpr2*-deficient pECs as well as *Bmpr2*-intact pECs, suggesting that BMP7 signaling is not mediated

mainly through BMPR2 and also mediated by other type 2 receptors such as ACVR2a (Fig. 1-8B)

Discussion

In this study, we showed that genetic ablation of *Smad1* in pECs or SMCs predisposes mice to PAH, suggesting that impaired SMAD1 signaling is relevant to the pathogenesis of PAH. Furthermore, the findings of reduced SMAD1/5/8 activation and the overactivation of TGF- β signaling in the BMPR2-deficient pECs suggests that a predominance of TGF- β signaling, gained from the genetically-disturbed balance between TGF- β and BMP signaling, may be an important factor triggering PAH development.

SMAD1 is a Downstream Mediator of BMP Signaling in Pathogenesis.

SMAD1 is a canonical downstream mediator of the BMP pathway that functions in a variety of cellular and developmental events. Therefore, reduced BMP activity in association with the reduced expression of its receptors is accompanied by decreased phosphorylation of SMAD1/5/8 in the development of diseases. Liu *et al.* found that pSMAD1/5/8 and BMPR1b were decreased in malignant glioma tissues compared with normal brain tissues.¹¹⁵ Yang *et al.* reported that the activated form of SMAD1 is deficient in the pulmonary vasculature of patients with a *BMPR2* mutation, suggesting that inactivation of SMAD1 plays a role in the PAH pathogenesis.⁷⁰ Our data demonstrate that *Smad1* deletion in either ECs or SMCs can induce PH in mice, implying that SMAD1 may be an important downstream signaling molecule of BMPR2 in PAH development.

SMAD1 Involvement in the Pathogenesis of PAH

Our data showing the role of SMAD1 in PAH may appear to conflict with the previous data from James West's group demonstrated that overexpression of tail domain *BMPR2* mutant (R899X) not affecting the SMAD1 activation resulted in PAH.¹⁰⁸ However, frame shift and nonsense mutations consist of 70% of total mutations of *BMPR2* gene and interestingly, approximately one-quarter of this type mutations are collected in the cytoplasmic tail domain. Nonsense mutation decay (NMD) is a cellular mechanism to destroy defective RNA transcripts with nonsense mutation to block the production of truncated proteins.¹¹⁶ Thus, most of *BMPR2* mRNAs with the cytoplasmic tail mutation are destroyed through NMD-mediated processing, leading to *BMPR2* deficiency.^{104, 117} In James West's studies, truncated *BMPR2* was not destroyed by NMD and showed normal SMAD1 activation. Thus, their mutant mice may not represent human PAH patients carrying a cytoplasmic tail mutation of *BMPR2*. Otherwise, this discrepancy suggests that some *BMPR2*-associated PAH might be SMAD-dependent (mutation in the kinase domain) and some are not (mutation in the tail domain). Mutations in the tail domain have been shown to interrupt interactions between *BMPR2* and the dynein light chain Tctex-1 as well as LIMK1, a key regulator of actin dynamics.^{118, 119} Thus, the cytoplasmic tail mutation may contribute to PAH development by disrupting these interactions with potentially critical signaling molecules in SMAD-independent manner. As an alternative interpretation, SMAD1 signaling is critical in ECs, not in SMCs for PAH development. Hansmann *et al.* presented a novel anti-proliferative axis – BMP2/*BMPR2*/*PPAR γ* /*ApoE* in PAH.¹²⁰ Mice with targeted deletion of *PPAR γ* in SMCs spontaneously developed PAH with elevated RVSP, RVH, and increased muscularization of the distal pulmonary arteries. This was independent of SMAD1/5/8

phosphorylation. Li *et al.* suggested a novel role of NOTCH3 in controlling proliferation of SMCs and in maintaining SMCs in an undifferentiated state.¹²¹ They found that the severity of disease in human PAH patients and rodent PH models correlated with the amount of NOTCH3 protein in vascular SMCs of the lung, suggesting that NOTCH3 signaling pathway in SMCs is crucial for the development of PAH. Even though it still remains unknown whether NOTCH3 and PPAR γ signaling in ECs are also critical contributors to PAH pathogenesis, up to now, our data with *Smad1* cKO mice demonstrated that SMAD1 deficiency in ECs is an important contributing factor for spontaneous PAH development.

SMAD1 Deficiency in ECs has a Greater Impact on PAH Than That in SMCs

Mice with endothelial *Smad1* deletion showed 40% penetrance while mice with *Smad1*-deficiency in SMCs displayed 13% penetrance. Why did endothelial deletion make a greater influence on PAH pathogenesis compared to deletion in SMCs? Malfunctioning of SMCs usually increase their growth rate, leading to thickening of vessel wall, whereas endothelial dysfunction contributes to PAH pathogenesis in various ways. Disordered endothelial proliferation forms neointimal layers⁸² and disrupted balance of vasoactive mediators exacerbates vasoconstriction^{14, 34, 84-86} and increased cell permeability allows serum growth factors to affect SMC proliferation and release of mitogen such as serotonin from ECs induces smooth muscle hyperplasia¹²² and the number of occluded vessels are increased by *in situ* thrombosis.² Therefore, because of these various effects of ECs, *Smad1* deletion in ECs may accelerate PAH development compared to *Smad1* deletion in SMCs.

Incomplete Penetrance of PAH in *Smad1* Conditional KO Mice

All the mice with endothelial or smooth muscle-*Smad1* deletion did not develop PAH. Only a subset of mice showed PH phenotypes including elevation of RVSP and thickening of smooth muscle layer. First, SMAD1/5/8 could be functionally redundant and compensate for the absence of other molecules. Cre-mediated genetic ablation of either *Smad1* or *Smad5* in ovarian granulosa cells results in normal reproductive function but that combined loss of *Smad1* and *Smad5* results in fertility defects and granulosa cell tumors.¹²³ SMAD1, SMAD5, and SMAD8 function redundantly in activating regression of the Müllerian duct mesoepithelium.¹²⁴ Second, additional environmental insult would be required for the manifestation of PAH with genetic predisposition. Inflammation has been associated with PAH as an influencing environmental factor. Inflammatory cells including macrophages and lymphocytes were accumulated and many proinflammatory cytokines were induced in the PAH vascular lesions.^{7, 12, 64, 93-97} Thus, if treated with inflammatory stimulants such as IL-6¹²⁵ and 5-lipoxygenase (5-LO)⁶⁸, *Smad1* cKO mice may show increased penetrance. Conversely, blocking inflammatory activation could be therapeutically beneficial for PAH patients. Studies with animal models support this. Platelet-activating factor (PAF) antagonists, an anti-inflammatory drug inhibited pulmonary vascular remodeling induced by hypobaric hypoxia in rats.¹²⁶ Inhibition of 5-lipoxygenase-activating protein (FLAP) suppressed hypoxia-induced pulmonary vasoconstriction *in vitro* and the development of chronic hypoxic pulmonary hypertension in rats.¹²⁷

Imbalance in TGF- β and BMP Signalings in PAH Pathogenesis

Many studies have been focused on impaired BMP signaling to explain PAH pathogenesis. However, as dysfunction of BMP pathway alone is not sufficient to

develop PAH, recent studies suggest a role of imbalance in TGF- β /BMP signaling for PAH development because BMP and TGF- β signaling form an opposing balance in pulmonary vascular homeostasis.¹²⁸ BMPR2-deficient pulmonary vessels lose control of overactivation of TGF- β signaling leading to predominance of TGF- β signaling.^{101-103, 129} Based on these reports, we hypothesized that BMPR2 deficiency results in overactivated TGF- β signaling in pECs. However, it is difficult to demonstrate that BMPR2 deficiency directly elevates TGF- β signaling *in vivo* system because even though our *Bmpr2* or *Smad1* cKO mice exhibit TGF- β activation such as increased pSMAD2/3, it could be a secondary effect of PAH. Hence, we isolated and immortalized pECs from a *R26^{creER+/-};Bmpr2^{2f/2f}* mouse, in which tamoxifen treatment induces *Bmpr2* deletion. This inducible system is advantageous because we can exclude strain and individual difference using the same parental cells. *Bmpr2*-deleted pECs showed the low level of pSMAD1/5/8 by BMP4 treatment and enhanced phosphorylation of SMAD2 not only at the basal level but also by TGF- β 1 treatment compared with *Bmpr2*-intact pECs, supporting our hypothesis that BMPR2 deficiency overactivates TGF- β signaling in pECs. In addition, we found a marked reduction of total SMAD2 with an accompanying high level of pSMAD2. This may have arisen because activated SMAD2 is multi-ubiquitinated and is degraded by the proteasome.¹³⁰⁻¹³²

These results from pECs present the possibility of the future therapeutic options to restore TGF- β /BMP imbalance. Recent reports support our results and suggest potential candidates for inhibitor of TGF- β pathway. Inhibition of TGF- β signaling by ALK5 inhibitor, IN-1233 prevented PAH in the monocrotalin-treated rat, which showed increased TGF- β activity.¹⁰³ The angiotensin II type 1 receptor (AT1) blocker, losartan

averted aortic aneurysm in a mouse model of Marfan syndrome (MFS), which is associated with increased TGF- β signaling.¹³³ In the aspect of BMP activation, Irrespective of BMPR2 deficiency, pECs activated SMAD1/5/8 in response to BMP7. This intact BMP7 response was also shown in SMCs with *Bmpr2* deletion.⁷⁷ Therefore, restoration of pSMAD1/5/8 by BMP7 could be beneficial to PAH patients with BMPR2 deficiency.

Thickening of the Smooth Muscle Layer Might not be Sufficient to Lead to High Pulmonary Pressure.

The PH group in *L1cre(+);Smad1^{2f/2f}* mice showed thickened vessel walls, implying that high pulmonary pressure was associated with the muscularization of pulmonary arterioles. However, *Tagln-cre(+);Smad1^{2f/2f}* mice including Non-PH and PH groups showed thickening of the vascular smooth muscle layer, suggesting that SMAD1 dysfunction in SMCs directly affected proliferation of vascular smooth muscles. Yang *et al.* reported that SMAD1 dysfunction promoted MAPK signaling leading to aggressive proliferation of PASMCs.⁷⁰ More importantly, our result suggests that thickening of the smooth muscle layer might not be sufficient to lead to high pulmonary pressure in the absence of related endothelial dysfunction. We speculate that SMAD1 deficiency in pECs results in endothelial dysfunction. Pulmonary endothelial dysfunction induces sustained constriction of blood vessels due to an increase of vasoconstrictors such as endothelin or decrease of vasodilators such as eNOS or prostacyclin. These alterations of vasoactive molecules were readily observed in PAH patients^{14, 44, 134, 135} and PH animal models.¹³⁶⁻¹³⁸ Continued vascular constriction leads to high pulmonary pressure and results in thickening of the vascular smooth muscle layer. However, although *Smad1* deletion in SMCs induces thickening of smooth muscle layer, unaffected

endothelial cells may enable vessels to dilate more in response to the high blood pressure gained from thickened blood vessels. Therefore, thickening of the smooth muscle layer might not be sufficient to sustain high pulmonary blood pressure.

***Smad1* cKO Mice and Inducible *Bmpr2* KO pECs are Useful Resources for Future Mechanism Research, Drug Screening and Preclinical Study.**

In this study, we developed two useful resources for PAH studies; *Smad1* cKO mice and inducible *Bmpr2* KO pECs. In the future, these mice and pECs will be usefully utilized for mechanistic research to find downstream targets of BMPR2/SMAD1 pathway for novel PAH therapy and when candidate drugs are developed or preexisting medicines with potential effect are not tried to PAH patients, we can get valuable information about efficacy and effectiveness of candidate drugs for PAH using these materials.

Table 1-1. Primers for genomic and semi-quantitative PCR reaction

| Genomic PCR | | | |
|-------------|---------------------------|-------------|--------------------------|
| Smad1-A | CACCTGTGCCCCCTCCAAGT | Smad1-B | GAGCTCTGCTCCGCCACTCA |
| Alk1-F | CAGCACCTACATCTTGGGTGGAGA | Alk1-R | ACTGTTCTTCTCGGAGCCTTGTC |
| Bmpr2-2A | CACACCAGCCTTATACTCTAGATAC | Bmpr2-6R | CACATATCTGTTATGAAACTTGAG |
| Bmpr2-2C | TTATTGTAAGTACACTGTTGCTGTC | | |
| L1cre-F | GTTTTCCCTTTGAAAAACACGATGA | L1cre-R | ATCAGGTTCTTGCGAACCTCATCA |
| Tagln-cre-F | CTCCTTCCAGTCCACAAACGAGC | Tagln-cre-R | GGGCGATCCCTGAACATGTCC |
| R26R-F | GTCGTTTTACAACGTCGTGACT | R26R-R | GATGGGCGCATCGTAACCGTGC |
| RT-PCR | | | |
| Gapdh-F | CAATGCATCCTGCACCACCAA | Gapdh-R | GTCATTGAGAGCAATGCCAGC |
| Bmpr2-F | GTTGACAGGAGACCGGAAACAG | Bmpr2-R | GGAGACTCAGATATTTGCACAG |
| Tgfr2-F | TTGCCTGTGTGACTTCGGGCT | Tgfr2-R | CTATTTGGTAGTGTTCCAGCGA |
| Acvr2a-F | CGTTCGCCGCTTTTCTTATC | Acvr2a-R | AGGATTTGAAGTGGGCTGTG |
| Alk1-F | TCATGGTGCACAGTGGTGCTG | Alk1-R | CAAATCCCGCTGCTTCTCCTG |
| Alk2-F | AGTCATGGTTCAGGGAGACG | Alk2-R | TGCAGCACTGTCCATTCTTC |
| Alk3-F | TAAAGGCCGCTATGGAGAAG | Alk3-R | CCAGGTCAGCAATAAGCAA |
| Alk6-F | CACTCCATTCTCATCAA | Alk6-R | TTCCAATCTGCTTCACCATC |
| Tgfβ1-F | CGGAAGCGCATCGAAGCCATCC | Tgfβ1-R | GCAAGCGCAGCTCTGCACGG |
| Nos3-F | TTCCGGCTGCCACCTGATCCTAA | Nos3-R | AACATATGTCCTTGCTCAAGGCA |
| Tie2-F | CTCATCTGTGGACGCTGGATG | Tie2-R | GGCACTGAGTGGATGAAGGAG |
| Eng-F | TGCACTCTGGTACATCTATTC | Eng-R | TGGATTGGGCAGTTCTGTAAA |
| Flk1-F | AGAACACCAAAAAGAGAGGAACG | Flk1-R | GCACACAGGCAGAAACCAGTAG |

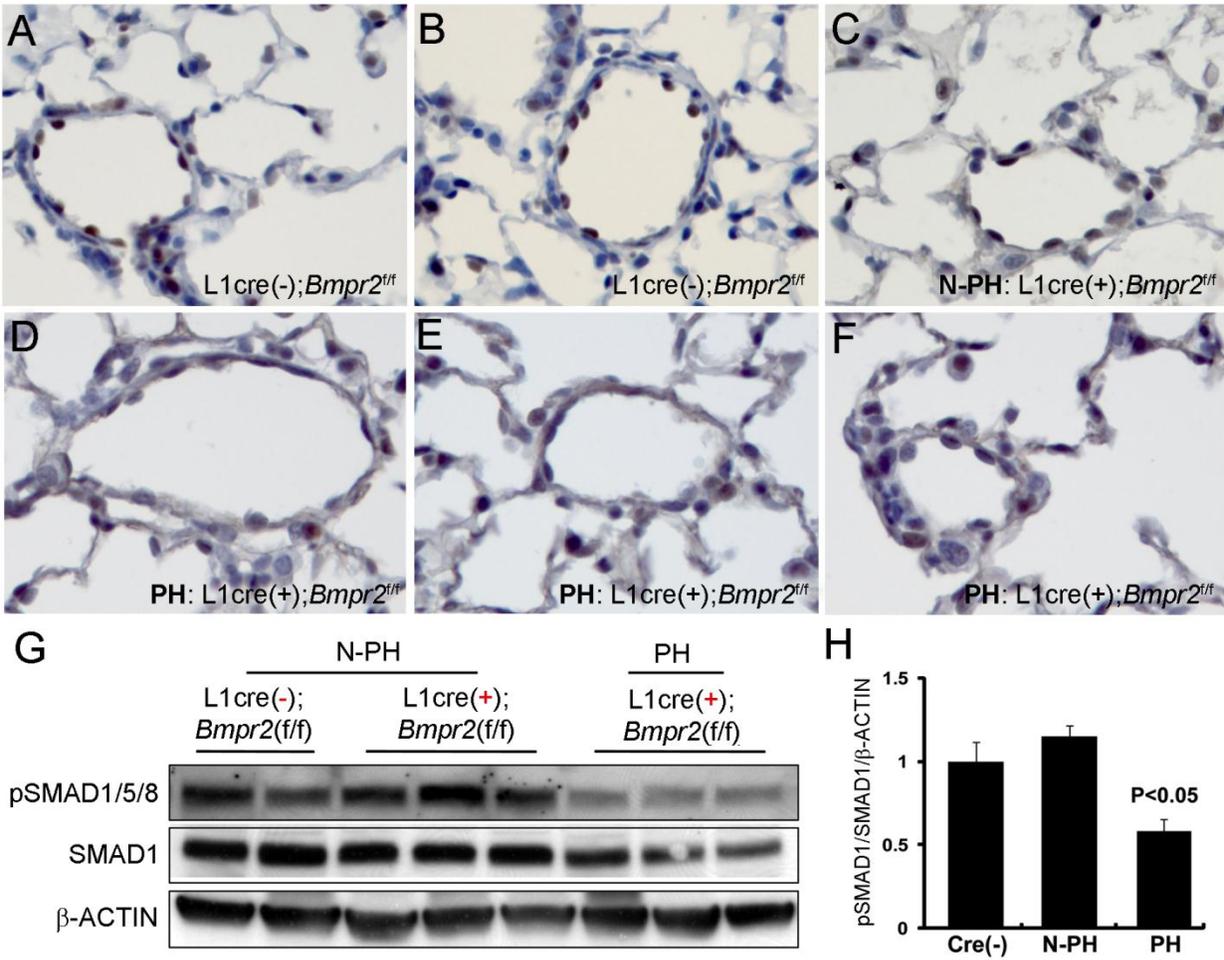


Figure 1-1. BMP response was severely impaired in the lungs of PH mice [L1cre(+);*Bmpr2*^{fl/fl}]. (A-F) The levels of pSMAD1/5/8 and SMAD1 were examined to compare BMP response among lungs of cre-negative controls (A,B), N-PH (C), and PH mice (D-F). (G-H) The level of pSMAD1/5/8 was significantly decreased in the PH lungs compared to controls and N-PH lungs.

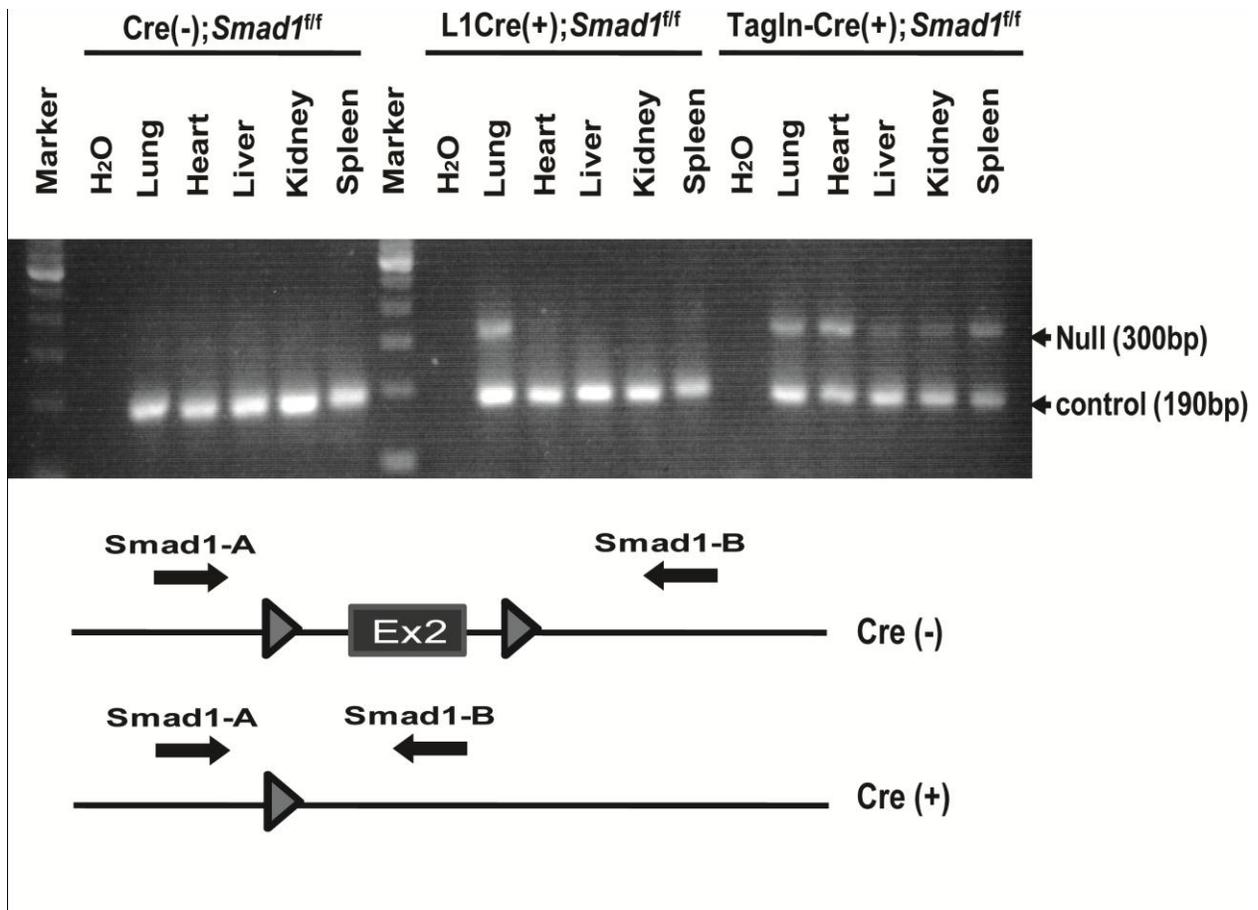


Figure 1-2. *Smad1* deletion in L1cre(+);*Smad1^{fl/fl}* and *Tagln*-cre(+);*Smad1^{fl/fl}* mice. Deleted *Smad1* allele was examined in various organs of *Smad1*-conditional KO (cKO) mice and cre-negative control mice by genomic PCR analysis. Null *Smad1* was detected at 300 bp. A primer set amplifying the *Alk1* locus (190 bp) was used as a control for the PCR amplification. Genomic segment between 2 loxp sites including exon2 was deleted by cre recombinase activated by tamoxifen leading to production of PCR product by a primer set (Smad1-A and Smad1-B).

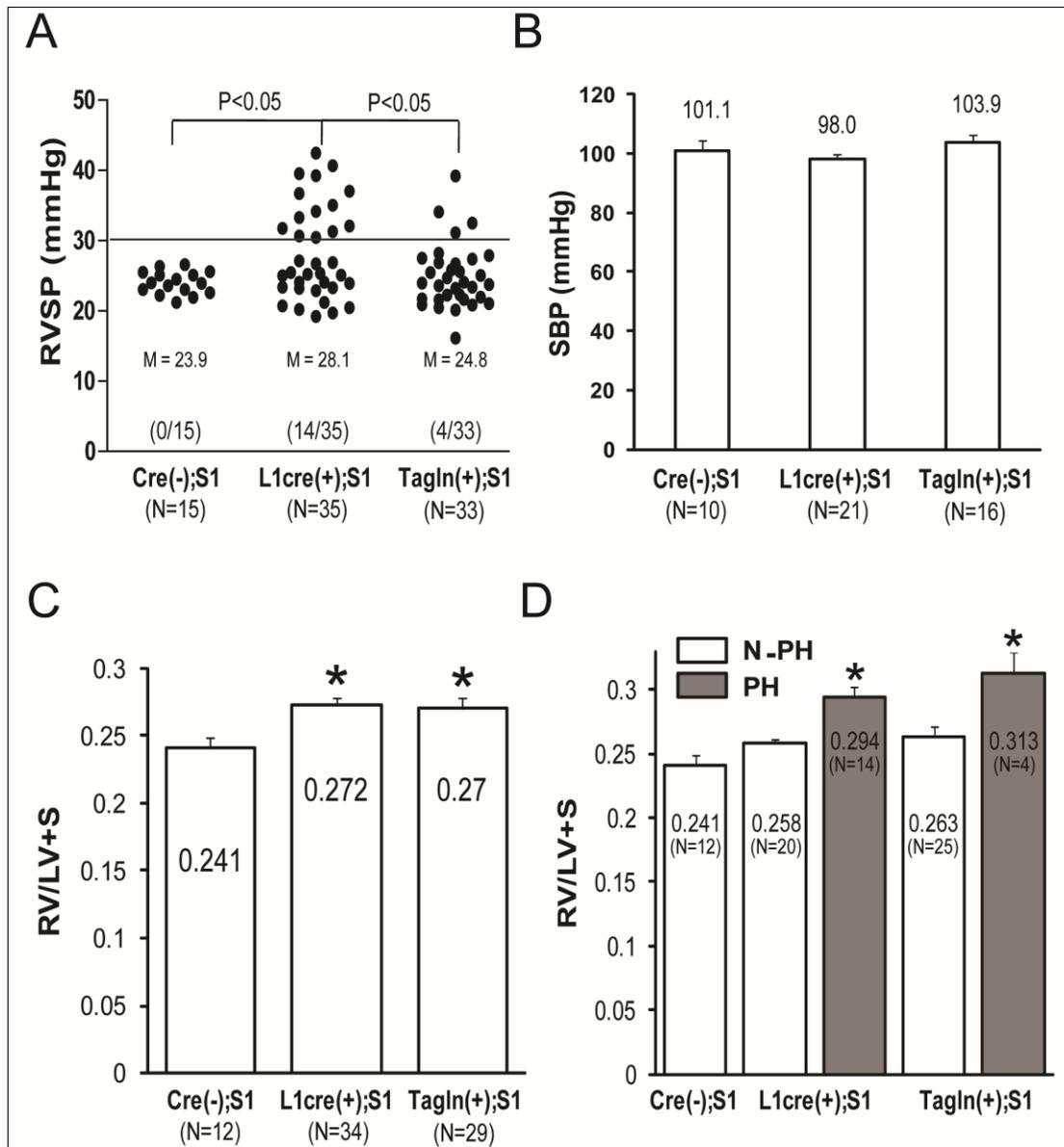


Figure 1-3. *Smad1* deletion in SMCs and ECs resulted in elevation of RVSP and RV hypertrophy. (A) closed circles indicate RVSP of each mouse. More than 30 mmHg of RVSP was designated as pulmonary hypertensive pressure. 40% of L1cre(+);*Smad1*^{f/f} mice and about 12% of Tagln-cre(+);*Smad1*^{f/f} mice were pulmonary hypertensive. RVSP of L1cre(+);*Smad1*^{f/f} was significantly higher compared to either that of controls or that of Tagln-cre(+);*Smad1*^{f/f}. (B) Systemic blood pressure was not significantly different by genotypes of mice. (C) Both *Smad1*-cKO mouse lines showed significant RV hypertrophy compared to cre-negative controls. (D) When mice were divided into non-pulmonary hypertensive (N-PH) and pulmonary hypertensive (PH) groups, RV hypertrophy of PH mice was significantly higher than cre-negative controls and N-PH mice of each cKO group. Statistical differences ($p < 0.05$) were indicated by asterisks above each bar.

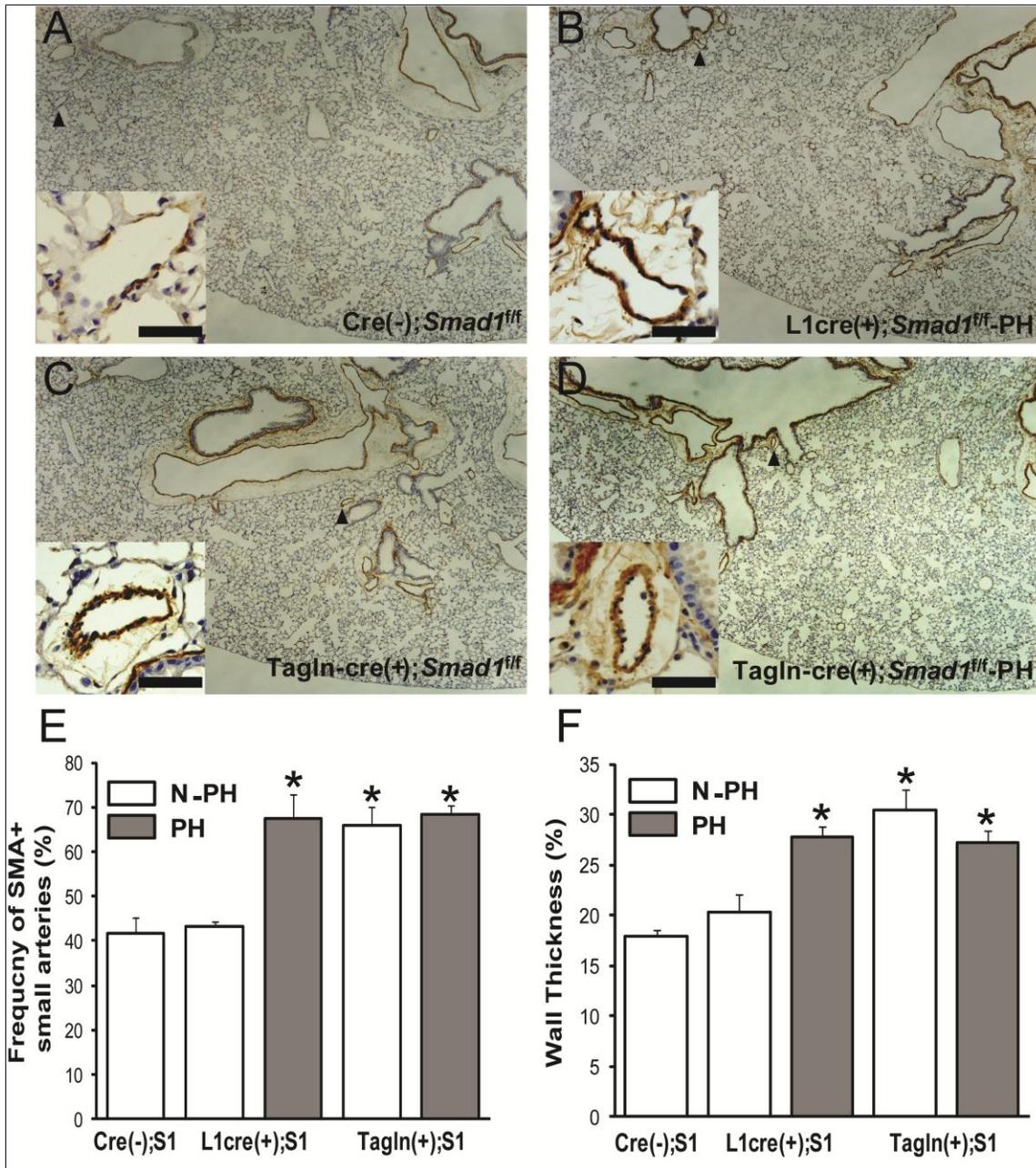


Figure 1-4. Elevated pulmonary pressure is associated with pulmonary vascular remodeling. (A-D), pulmonary arterioles were visualized by staining with anti- α SMA antibody. A representative vessel from each group (indicated by arrowheads) is shown at higher magnification at the left bottom corner of each frame. Scale bars in insets represent 50 μ m. PH mice of each *Smad1*-cKO mice showed muscularized vessels (E) and thickened vessel walls (F) compared with cre-negative controls. Interestingly, N-PH group of *Tagln*-cre(+);*Smad1*^{ff} also exhibited muscularized vessels with thickened vessels walls. * indicates significant statistical difference ($p < 0.05$) compared to cre-negative controls.

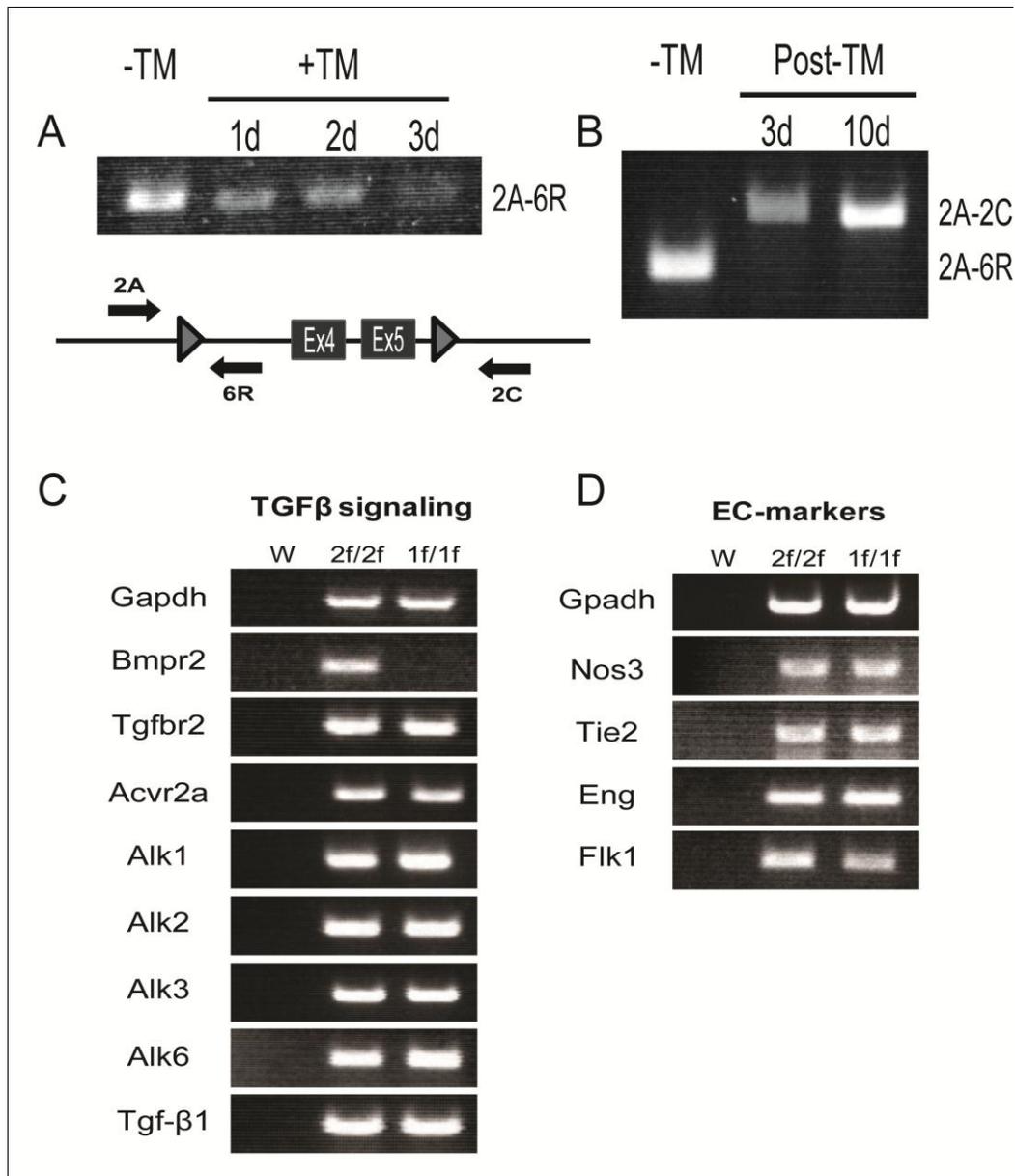


Figure 1-5. Deletion of *Bmpr2* gene in pulmonary endothelial cells (pECs). (A) pECs were treated with tamoxifen (TM) for 3 days. As shown in the diagram, TM enables cre recombinase to loop out the genomic segment containing 6R primer binding sequence and exon 4-5, which is flanked by two loxp sites (triangles). This deleted allele can produce a PCR product with the 2A-2C primer pair. Deletion of *Bmpr2* gene was examined using the 2A-6R primer pair. (B) Following growth in TM-free media, reappearance of an intact *Bmpr2* gene in TM-treated pECs was examined using primer pairs for the deleted allele [2A-2C] and the intact allele [2A-6R]. The mRNA levels of TGF- β signaling molecules (C) and endothelial markers (D) were examined by semi-quantitative RT-PCR to determine whether tamoxifen treatment for *Bmpr2* deletion affected expression of other molecules in TGF- β pathway and EC characteristic.

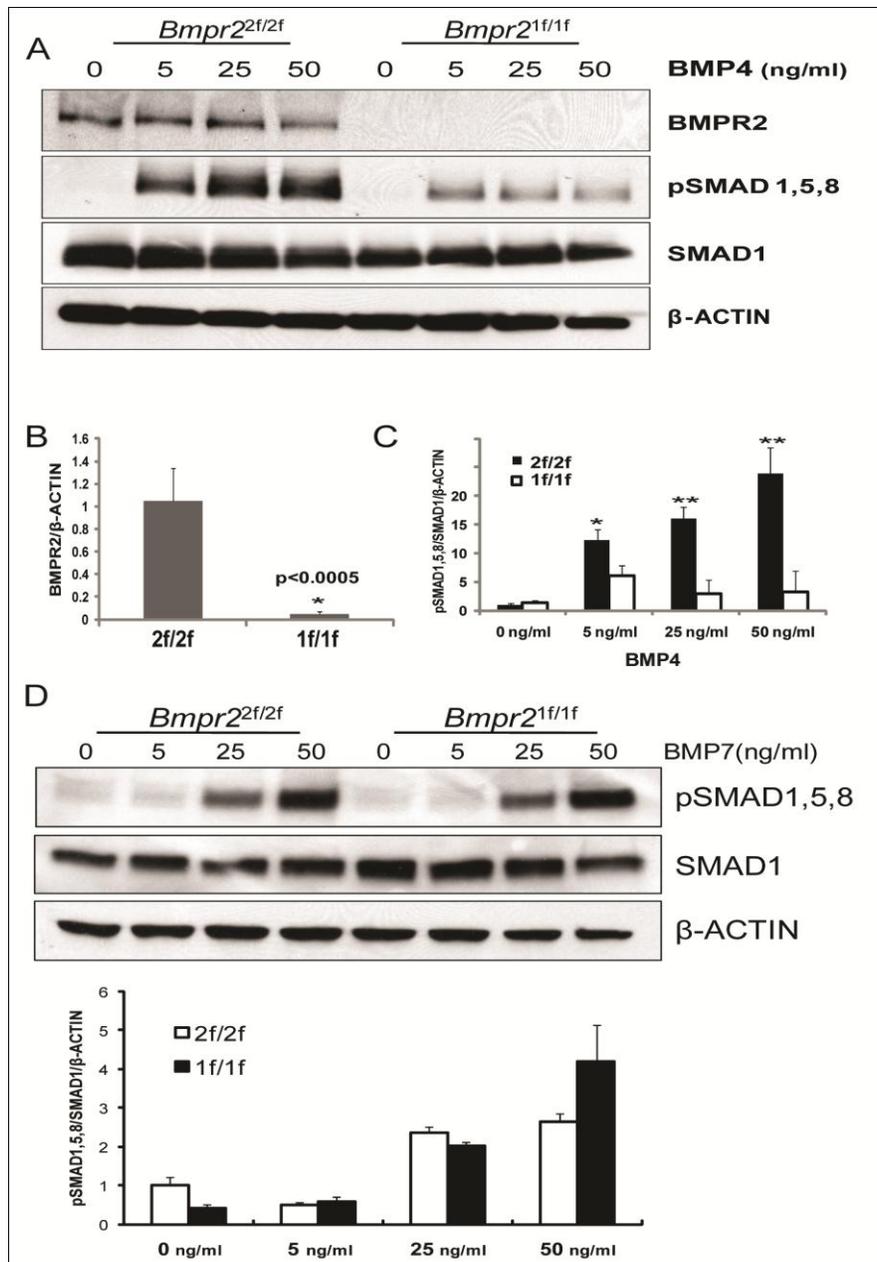


Figure 1-6. *Bmpr2*-deleted pECs displayed a reduced response to BMP4 but not BMP7. (A) BMP4 (0 – 50 ng/ml) was added to *Bmpr2*-intact (*Bmpr2*^{2f/2f}) and *Bmpr2*-deleted (*Bmpr2*^{1f/1f}) cells for 30 minutes after serum starvation. The levels of pSMAD1/5/8 were examined to evaluate the BMP4 response. (B) The protein levels of *Bmpr2* were determined for *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells. (C) BMP4 response was quantified by the ratio of pSMAD1,5,8/SMAD1/β-actin in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells. (D) BMP7 (0 – 50 ng/ml) was added to *Bmpr2*-intact (*Bmpr2*^{2f/2f}) and *Bmpr2*-deleted (*Bmpr2*^{1f/1f}) cells for 30 minutes after serum starvation. The levels of pSMAD1/5/8 were determined to evaluate differences in response to BMP7. The BMP7 response was quantified by the ratio of pSMAD1,5,8/SMAD1/β-actin in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells.

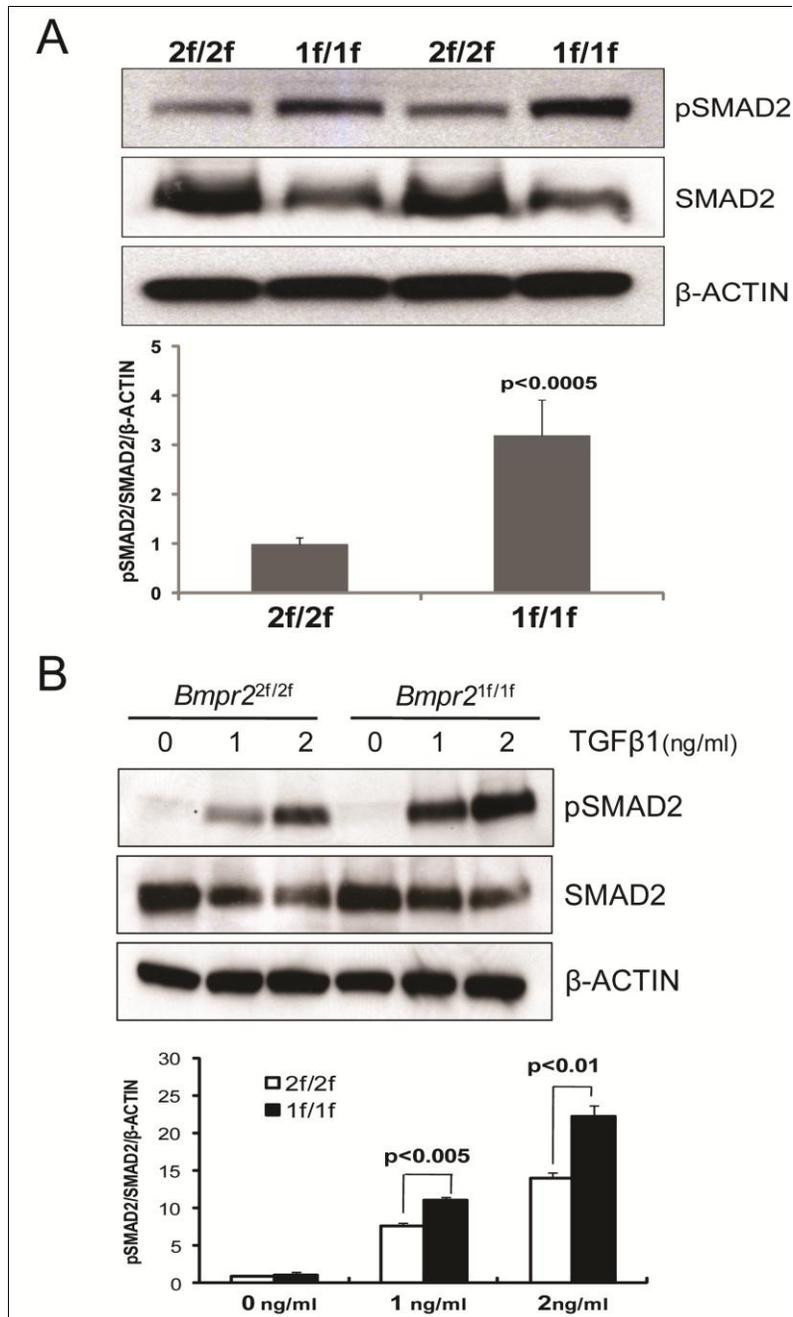


Figure 1-7. Reduced BMP response induced overactivation of TGF- β signaling. (A) The levels of pSMAD2 were examined to compare basal TGF- β activation in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells under normal media condition without additional cytokine treatment. TGF β activation was quantified by the ratio of pSMAD2/SMAD2/ β -actin in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells. (B) In order to compare TGF- β response in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells, TGF- β 1 (0-2 ng/ml) was added for 30 minutes after serum starvation. TGF- β response was quantified by the ratio of pSMAD2/SMAD2/ β -actin in *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells.

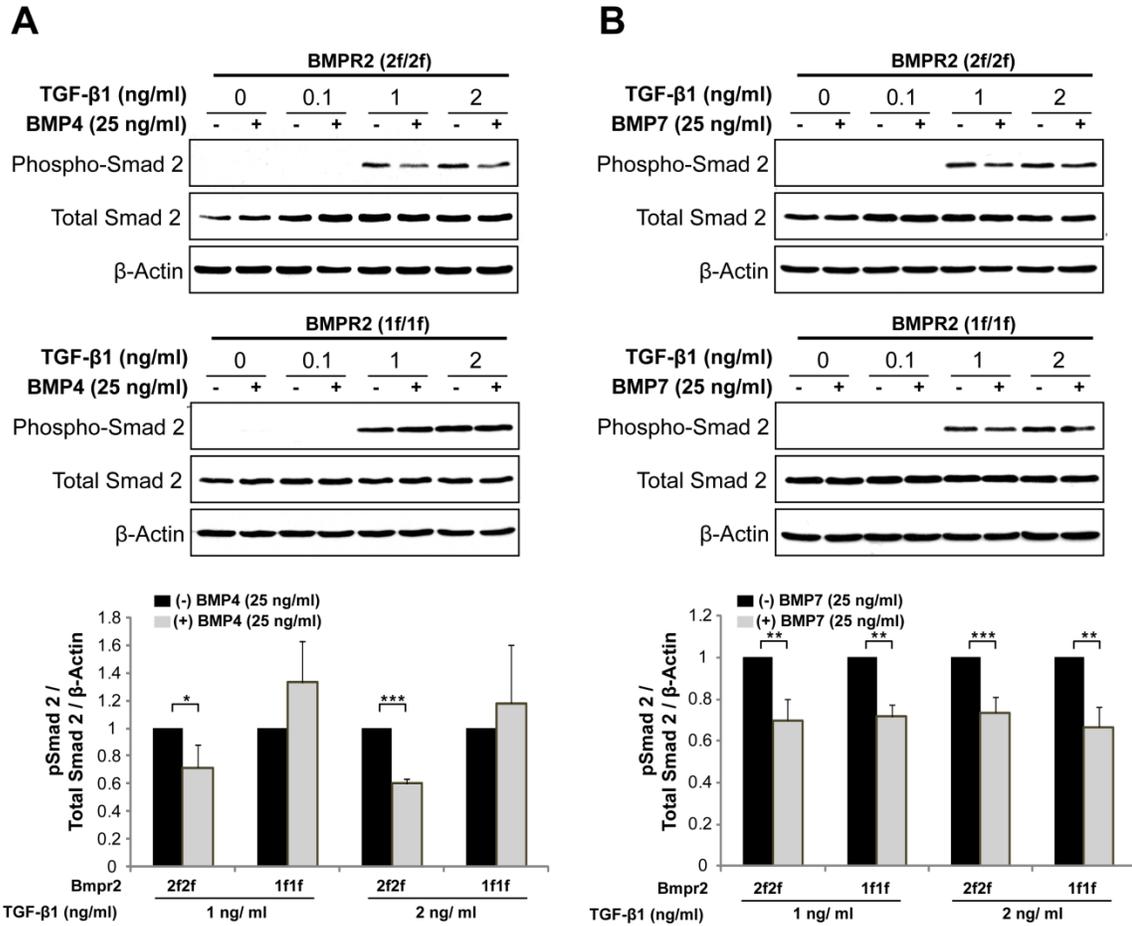


Figure 1-8. BMP and TGF-β signaling form an opposing balance in pECs (A) Activation of TGF-β signaling was suppressed by either BMP4 or BMP7 in *Bmpr2*-intact pECs. TGFβ-1 (0-2 ng/ml) was treated on *Bmpr2*^{2f/2f} and *Bmpr2*^{1f/1f} cells for 30 minutes after serum starvation. TGF-β response was quantified by the ratio of pSMAD2/SMAD2/β-actin. However, (B) this suppression was induced only by BMP7, not by BMP4 in *Bmpr2*-deficient pECs.

CHAPTER 2
THERAPEUTIC EFFECT OF VEGF BLOCKADE ON ARTERIOVENOUS
MALFORMATIONS (AVMS) IN AN ANIMAL MODEL FOR HEREDITARY
HEMORRHAGIC TELANGIECTASIA

Introduction

Hereditary Hemorrhagic Telangiectasia (HHT)

Hereditary Hemorrhagic Telangiectasia (HHT) is an inherited autosomal dominant vascular disease affecting 1 in 5,000–8,000 individuals worldwide.¹³⁹⁻¹⁴¹ The disease is characterized by epistaxis (spontaneous and recurrent nosebleeds), mucocutaneous telangiectases as well as arteriovenous malformations (AVMs) in the brain, lungs, and visceral organs including the liver and GI tract.¹⁴²⁻¹⁴⁵ An AVM is a direct connection between arteries and veins without intervening capillaries¹⁴² and is an underlying defect associated with recurrent nose bleeding, telangiectases, and visceral hemorrhaging. In AVMs, high velocity-arterial blood invades venous vascular beds creating turbulent and swirling flows leading to tortuous and irregular vascular remodeling. When these blood vessels rupture, it results in chronic and severe anemia due to massive blood loss. Brain and lung hemorrhages are life-threatening and the major cause of death in young patients.^{146, 147} To compensate for the lost blood from GI and nose bleeding, patients receive regular blood transfusions and/or iron supplementation therapy. More than 90% of HHT patients experience recurrent nosebleeds by their 60s¹⁴⁸ and this affliction impacts profoundly on the patients quality of life.^{149, 150} All the patients receiving Young's procedure for completely blocking nostrils said that the quality of their lives was improved following the surgery.¹⁵¹

Limitation of Current Therapeutic Interventions in HHT

Therapeutic treatments are available to alleviate telangiectasia and visceral lesions. Embolization can prevent blood flow to AVM lesions by occluding blood vessels with metal coils and is effective for pulmonary and cerebral AVMs.^{152, 153} For acute nosebleeds, packing is a common method and in chronic and recurrent nose bleeding, humidification with topical lubricant, laser therapy, and surgeries including Young's procedure are applied depending on the unique condition of the patient.¹⁵⁴ Systemic hormonal treatments or prothrombotic strategies using antifibrinolytic agents have been useful in limiting excessive hemorrhaging.^{155, 156} However, the mechanism of hormonal efficacy remains unknown and the negative impacts of prothrombotic agents must be considered before treatment.¹⁵⁷ Thus, the development of therapies targeting the pathogenic mechanisms underlying AVM formation remains an urgent issue.

Genetic Studies in HHT

Heterozygous mutations in HHT-causing genes have been identified: *ENG* for HHT1,¹⁵⁸⁻¹⁶⁰ *ALK1* for HHT2,^{161, 162} and *SMAD4* for Juvenile Polyposis and HHT (JP-HHT).¹⁶³ Over 80% of HHT patients possess heterozygous mutations in one of these genes, while mutations at additional loci are responsible for the remaining 20%. These loci include the HHT3 locus (chromosome 5)¹⁶⁴ and the HHT4 locus (chromosome 7)¹⁶⁵ even though causative genes remain unidentified.

Genetic Mouse Models for HHT

The role of *ALK1* in HHT pathogenesis has been explored using genetically modified mouse models.^{166, 167} Heterozygous *Alk1* null mice displayed HHT-like phenotypes such as age-dependent subcutaneous and mucocutaneous vascular lesions as well as hemorrhages in the lungs, GI tract, liver, brain and spleen.

Penetrance, however, was not complete and symptoms when observed were not as severe as observed in humans. Homozygous *Alk1* knockout (KO) mice died *in utero* displaying vascular defects such as hyper-dilated blood vessels and AVMs.^{166, 168} Targeted *Alk1*-deletion in endothelial cells of transgenic mice resulted in AVMs and hemorrhages in the vascular beds of brain, lung and GI tract.¹⁰⁹ Recently, our lab demonstrated that the conditional deletion of *Alk1* in adult mice using the tamoxifen-inducible RosacreER driver recapitulated HHT phenotypes common in human patients, such as severe hemorrhaging and low hemoglobin levels.¹⁶⁹

Secondary Hits are Required for AVM Development.

AVM lesions in HHT patients and *Alk1* mutant mice appear in selective organs such as the lung, GI tract, liver and brain despite the global presence of *ALK1* or *ENG* mutations. Even in the same organ, some vascular beds form AVMs and hemorrhage while other vessels remain normal in appearance. This suggests that genetic predisposition is not sufficient to cause disease and other genetic or environmental factors play a role. When excisional wounds were performed on the dorsal skin and ear of *Alk1*-deficient adult mice, vascular dysplasia, or “*de novo* AVM” was induced around to the wounded area.¹⁶⁹ Vessels removed from the wounded area were unaffected. This was evidence to suggest that ALK1 deficiency and injury combined to induce the vascular lesion. Angiogenesis is variably regulated by processes associated with wound-healing. These include the actions of infiltrating inflammatory cells, the turnover of extracellular matrix via matrix metalloproteinase (MMP) secretion, and the local activity of growth factors and cytokines produced by endogenous and recruited cells.¹⁷⁰ Which event is most critical for AVM formation in *Alk1*-deficient blood vessels is a question we addressed in this study. We focused on LPS-induced inflammation and

growth factor-induced angiogenesis. Mucocutaneous telangiectasia occurs frequently on the lips, tongue, and nose which are constantly exposed to infection. Increased levels of inflammatory markers factors such as MMP9 and interleukin-6 (IL-6) have been demonstrated in brain AVM tissue.^{171, 172} Inflammatory cells including neutrophils and macrophages/microglia were frequently identified in the vascular walls of AVM tissue.¹⁷³ In addition, the tissues and plasma of HHT patients contain high levels of vascular endothelial growth factor (VEGF). Vessel density was also increased.^{174, 175} Brain AVMs usually develop congenitally during vascularization. Some case reports showed that anti-angiogenesis therapy using bevacizumab and thalidomide worked well to prevent epistaxis and liver AVMs.^{176, 177}

In this report, we demonstrate that inflammation or direct angiogenic stimulation can induce AVMs in *Alk1*-deficient subdermal vessels. Angiogenic stimulation is a necessary sequella to inflammation in the induction of AVM formation. Then, angiogenesis blockade using a VEGF-neutralizing antibody is effective in alleviating wound-induced AVMs and internal bleeding in GI tract and lungs in ALK1-deficient adult mice. Taken together, our findings suggest that angiogenesis blockades could be promising and fundamental medication in future clinical trials for HHT patients.

Results

Vascular Endothelial Growth Factor (VEGF) or Lipopolysaccharide (LPS) can Induce De Novo AVMs in *Alk1*-Deficient Subdermal Vessels.

In order to examine whether inflammation or angiogenesis can induce AVM formation in *Alk1*-deficient mice, VEGF or LPS was administered to mice in the form of PLGA particles implanted subdermally under the dorsal skin. LPS-PLGA and VEGF-PLGA particles were injected into cre-negative mice as a control to assess whether

either agent alone could induce vascular lesions in the absence of an *Alk1* deficiency. After 7 days, blue latex dye was infused through the left heart to visualize the blood vessels and to locate arterial-venous (AV) shunts. The dye should be prohibited from traversing capillary beds unless an AV shunt is present. The particles formed an orange round aggregation on the subdermal area because of rhodamine. After removing particle-aggregates, the skin was cleared to reveal the vascular network. In *Alk1* wild type mice, a higher vascular density was evident around LPS and VEGF particle-injected areas compared to neighboring regions, but vascular malformation such as AV shunts were not observed (Fig. 2-1A, B). PLGA particles themselves occasionally induced higher vascular densities around implantation sites in *Alk1*-deficient mice but did not induce AV shunts. This was demonstrated in tamoxifen-treated *R26^{creER/+};Alk1^{2f/2f}* mice in which PBS-PLGA was implanted into the dorsal skin. PBS-PLGA had no affect on subdermal vessels (Fig. 2-1C) or, on rare occasions, induced a moderate level of abnormal vasculature (Fig. 2-1D). When VEGF-PLGA was injected into *Alk1*-deficient skin, tortuous, irregular, and excessive vasculature developed around the particles and surrounding veins were dilated and contained latex dye, indicating the presence of AV shunts (Fig. 2-1E). LPS-PLGA particles also induced a comparable level of tortuous vessels and AV shunting (Fig. 2-1F).

VEGF and LPS Stimulated Angiogenesis.

VEGF is a strong angiogenic stimulator in vascularization^{178, 179} and tumors¹⁸⁰ and LPS has been to stimulate angiogenesis.^{181, 182} Therefore, we questioned whether VEGF or LPS released from PLGA particles elicited an angiogenic response. We observed the vascular morphology around VEGF- or LPS-PLGA-injected area 4 days following PLGA-particle injection. Sprouting blood vessels were seen to be drawn

towards the VEGF and LPS particles (Fig. 2-1G, H). This result was consistent with angiogenesis being a common, and necessary, element to both VEGF and LPS-induced AVM formation. It could also be a critical factor in wound-induced AVMs.

VEGF-Blockade Suppressed Wound-Induced AVM in *Alk1*-Deficient Subdermal Vessels.

To test the participation of VEGF in the development of inflammation-induced or wound-induced AVM formation, experiments were repeated in the presence of VEGF-neutralizing antibody to block angiogenesis. *Alk1*-deficient and wounded mice were simultaneously treated with VEGF antibody or saline alone. The local vasculature was evaluated 9 days following the application of the wound. Saline-treated control mice exhibited a typical AV shunts with excessive, tortuous, and dilated vessels surrounding the wounded area (Fig. 2-2A, top panels). In contrast, AV shunts were only occasionally observed in VEGF antibody-treated animals and there were fewer tortuous and irregular vessels near the wound (Fig. 2-2A, bottom panels). Blood vessels positive for latex dye were quantified to measure the severity of AVMs in saline-treated and VEGF antibody-treated groups. The VEGF antibody-treated group (mean: 17.38 % vessel area) displayed a reduction in vessel content of approximately 40% in mean vessel area relative to saline-treated group (17.38 % versus 29.04 %, respectively). Thus VEGF antibody was capable of suppressing AVM formation in *Alk1*-deficient subdermal vessels responding to a wound (Fig. 2-2B).

VEGF-Blockade Alleviated Internal Bleeding and Low Hemoglobin Occurring in *Alk1*-Deficient Mice.

Alk1-deleted mice (tamoxifen-treated $R26^{creER/+}; Alk1^{2f/2f}$ mice) die within 2-3 weeks of tamoxifen treatment displaying severe GI, lung, and uterine hemorrhage.¹⁶⁹ Since AVMs are a major cause of bleeding in HHT, we questioned whether

angiogenesis blockade would suppress visceral bleeding in *Alk1*-deficient mice. VEGF antibody (or saline, as a control) was injected into *R26^{creER/+};Alk1^{2f/2f}* mice following tamoxifen treatment. After 9 days, the concentration of blood hemoglobin was determined as an indirect measure of internal hemorrhaging. Normal mice had hemoglobin levels at 15-17 g/dl (data not shown). *Alk1*-deficient mice treated with saline exhibited significantly lower hemoglobin levels (mean: 8.6 g/dl) indicative of severe hemorrhaging. On the other hand, VEGF antibody-treated mice displayed near normal levels of hemoglobin (13.8 g/dl), suggesting that internal bleeding was greatly reduced by VEGF antibody treatment (Fig. 2-3A). Thoracic and abdominal cavities were explored for direct evidence of hemorrhaging. In addition, heart size was compared as well because poorly oxygenated blood due to severe hemorrhage would force the heart to work harder to maintain oxygen levels, leading to enlargement of the heart. The relative severity of GI and lung bleeding and heart enlargement was categorized as weak, moderate, or severe (Fig. 2-7). The saline-treated group showed moderate to severe bleeding in the lungs and GI tract (11/12 animals; Table 2-1, Fig. 2-3B). The majority of the VEGF antibody-treated group displayed weak GI and lung hemorrhaging (9/12 mice; Table 2-1, Fig. 2-3B). The hearts of saline-treated group were more enlarged compared to the VEGF antibody-treated group (Fig. 2-3B) suggestive that severe hemorrhaging in the mock-protected group adversely affected the size of the heart.

VEGF-Blockade Suppressed LPS-Induced AVMs in *Alk1*-Deficient Subdermal Vessels.

To determine whether inflammation alone, in the absence of angiogenesis, could induce AVMs in *Alk1*-deficient mice, LPS particles were implanted in subdermal sites and animals were simultaneously treated with VEGF-antibody to block angiogenesis.

VEGF-antibody suppressed AVM formation associated with LPS implantation and mock treatment with vehicle (saline; Fig. 2-4A through 2-4D). This result indicates that LPS-induced AVM formation is linked to angiogenic stimulation.

TNF- α did not induce severe AVM formation compared to LPS.

Inflammatory stimulation largely independent of an angiogenic response was evaluated with the implantation of TNF- α -PLGA particles into *Alk1*-deficient mice. TNF- α is less actively involved in the monocyte-induced angiogenesis compared to other inflammatory cytokines such as IL-8.¹⁸³ A higher vascular density was observed around TNF- α particles, indicating that TNF- α induced neovascularization. However, it did not provoke severe form of AVMs shown around LPS-particles (Fig. 2-5D through 2-5F)

Discussion

Although HHT is a genetic disease, vascular lesions appear only in limited vascular beds, indicating that localized secondary genetic or environmental insult is involved in development of AVMs. Recently, we have demonstrated that wound could induce AVMs in the subdermal vessels of a mouse model of HHT2.¹⁶⁹ In the present study, we showed that angiogenic stimulation by VEGF or LPS could mimic the wound effect for *de novo* AVM formation in subdermal vessels of ALK1-deficient mice (Fig. 2-6). We further demonstrated that treatment with VEGF-neutralizing antibody could inhibit both wound- and LPS-induced AVMs (Fig. 2-6), and could alleviate the internal bleeding in the *Alk1* conditional KO model. These data attest to the use of angiogenesis blockade for treating epistaxis and GI bleeding in HHT patients.

Therapeutic Potential of Anti-Angiogenic Drugs in HHT

Recent anecdotal clinical evidence suggests that anti-angiogenic drugs can be an effective therapy for HHT. The first ground-breaking report came from an Australian

group showing VEGF-neutralizing antibody, Bevacizumab (Avastin) treatment effectively regressed hepatic AVMs.¹⁷⁷ More recently, Bevacizumab treatment was shown to be effective for pancreatic AVMs,¹⁸⁴ GI bleeding,^{185, 186} and epistaxis¹⁸⁷⁻¹⁹⁰ in HHT patients. Lebrin et al showed that thalidomide, known to have anti-angiogenic and anti-inflammatory functions, enhanced mural cell recruitment in *Eng*^{+/-} mouse model, and alleviated the frequency and severity of nosebleeds in HHT patients.¹⁷⁶ Various side effects of Bevacizumab including cardiac failure, GI bleeding, wound-healing complication, and arterial thrombo embolic events have been reported.¹⁹¹ In order to reduce side effects through systemic administration of Bevacizumab, topical application to nasal area using spray has been tried and shown to be effective.^{192, 193}

Therapeutic Potential of Anti-Inflammatory Intervention

More than 90% of patients experience nosebleeds that stem from nasal Telangiectasia,¹⁹⁴ indicating that the nasal mucosa is very susceptible to AVM formation. We speculate that chronic inflammation, infection and immune activation in the nose might be related to the high susceptibility. Inflammatory cytokines gene expression was implicated in AVM formation as the promoter polymorphisms in IL-1 β , IL-6, TNF- α and APOE ϵ 2 are associated with AVM susceptibility and clinical presentation of intracranial hemorrhage.^{171, 195-197} More recently, Torsney et al showed that inflammation in the ears and eyelids almost invariably caused bleeding from thin-walled dilated vessels in the *Eng*^{+/-} mouse.¹⁹⁸ Our skin wound model demonstrated that AVMs form only in the wound areas in *Alk1*-deficient mice,¹⁶⁹ and here we showed that LPS could mimic the wound effect.

Inflammatory Stimulation Associated with Angiogenesis is Important for Inducing AVM Formation.

LPS is an endotoxin that provokes an acute inflammatory response by releasing proinflammatory cytokines including tumor necrosis factor- α (TNF- α), IL-1 β , IL-6, IL-8.¹⁹⁹ TNF- α is a cytokine that promotes a strong inflammatory response by recruiting immune cells, induces additional proinflammatory cytokines such as IL-1 and IL-6 and inhibits apoptosis of inflammatory cells.^{200, 201} We show that LPS can induce angiogenic stimulation in the context of an inflammatory response. Neovascularization and recruitment of sprouting blood vessels occurred around LPS-injected sites. Mattsby-Baltzer et al have suggested that endotoxin-mediated neovascularization is a component of inflammation and wound healing.¹⁸¹ LPS is associated with increased angiogenesis, vascular permeability, tumor cell invasion, and VEGF expression in macrophages.^{182, 202} TNF- α also induced angiogenesis via macrophage stimulation.²⁰³ Activation of the Eck RPTK by its inducible ligand B61 is involved in TNF- α -induced angiogenesis.²⁰⁴ We showed that the AVMs formed by LPS were more severe than those by TNF- α . Perhaps it happened because LPS might have provoked stronger inflammatory stimulation by releasing various proinflammatory cytokines (including TNF- α) compared to TNF- α -induced inflammation, leading to more intensive angiogenic stimulation. For instance, it was shown that production of IL-8 which may play a role in a more lasting myeloid cell-driven angiogenesis could be induced by LPS, but not TNF- α .¹⁸³

Taken together, our data suggest that blocking angiogenesis or inflammation could be an effective therapy for epistaxis and GI bleeding in HHT patients. Number of FDA approved drugs inhibiting angiogenesis or inflammation is rapidly increasing. In addition

to angiogenesis blockade and inflammatory inhibitors, drugs targeting some other aspect of AVM formation such as antifibrinolytic agent (Tranexamic acid),^{156, 205} antioxidants (N-Acetyl Cystein),²⁰⁶ estrogen analogs (Raloxifene),²⁰⁷ have been suggested for treating HHT. The preclinical model presented here would be an invaluable resource with which these potential drugs can be screened for effectiveness in preventing skin de novo AVMs and GI bleeding. Thus, inflammation prevention could be therapeutic for HHT patients suffering nosebleeds.

Table 2-1. The number of mice with hemorrhage in saline- and VEGF-antibody-treated groups

| | | Saline (n=12) | VEGF-Ab (n=12) |
|---------------|----------|---------------|----------------|
| GI bleeding | Weak | 1 | 7 |
| | Moderate | 6 | 3 |
| | Severe | 5 | 2 |
| Lung bleeding | Weak | 2 | 9 |
| | Moderate | 5 | 2 |
| | Severe | 5 | 1 |
| Liver AVM | Weak | 11 | 10 |
| | Moderate | 0 | 1 |
| | Severe | 1 | 1 |

Lung, GI bleeding and liver AVM of each mouse were categorized into three groups (weak, moderate, severe) and the number of groups was counted in saline-treated (n=12) and VEGF-treated (n=12) Alk1-deficient mice.

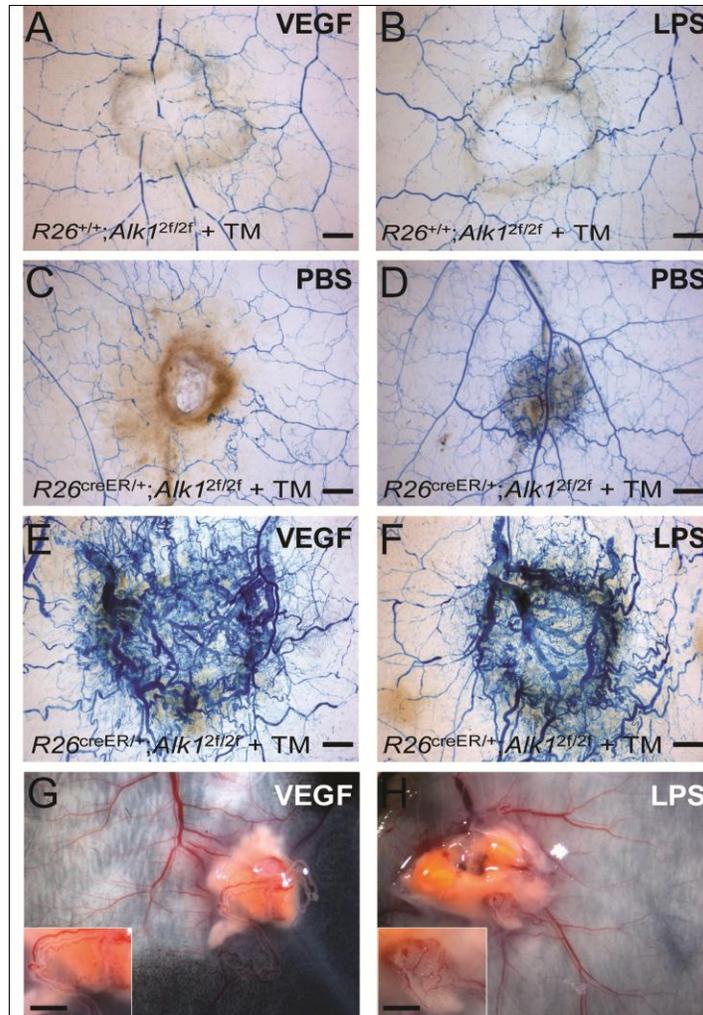


Figure 2-1. Angiogenic or inflammatory stimulation induces AVMs in *Alk1*-deficient adult subdermal vessels but not *Alk1*-wild type vessels. VEGF-PLGA and LPS-PLGA were implanted under dorsal skin as an angiogenic or inflammatory stimulus, respectively. Tamoxifen (TM) was intraperitoneally injected to effect *Alk1*-deletion. After 7 days, blue latex dye was infused through the heart to visualize blood vessels in the dorsal skin and particle-injected skins were cleared (see methods). (A-B) VEGF (A) and LPS (B) particles did not induce AVMs in Cre-negative controls ($R26^{+/+}; Alk1^{2f/2f} + TM$) bearing wild type *Alk1* alleles. (C-D) PBS-PLGA was implanted to investigate whether PLGA itself can induce AVM formation in *Alk1*-deficient mice ($R26^{creER/+}; Alk1^{2f/2f} + TM$). PBS-PLGA did not induce AVMs (C) or rarely formed abnormal vasculature (D). (E-F) VEGF (E) and LPS (F) particles induced severe AVMs and tortuous vessels in ALK1-depleted vessels ($R26^{creER/+}; Alk1^{2f/2f} + TM$). VEGF (G) and LPS (H) particles induced angiogenesis by recruiting sprouting vessels toward particles as seen 4 days post-implantation of loaded particles. At left bottom are higher magnifications of recruited sprouting vessels. Scale bars in each panel indicate 1 mm.

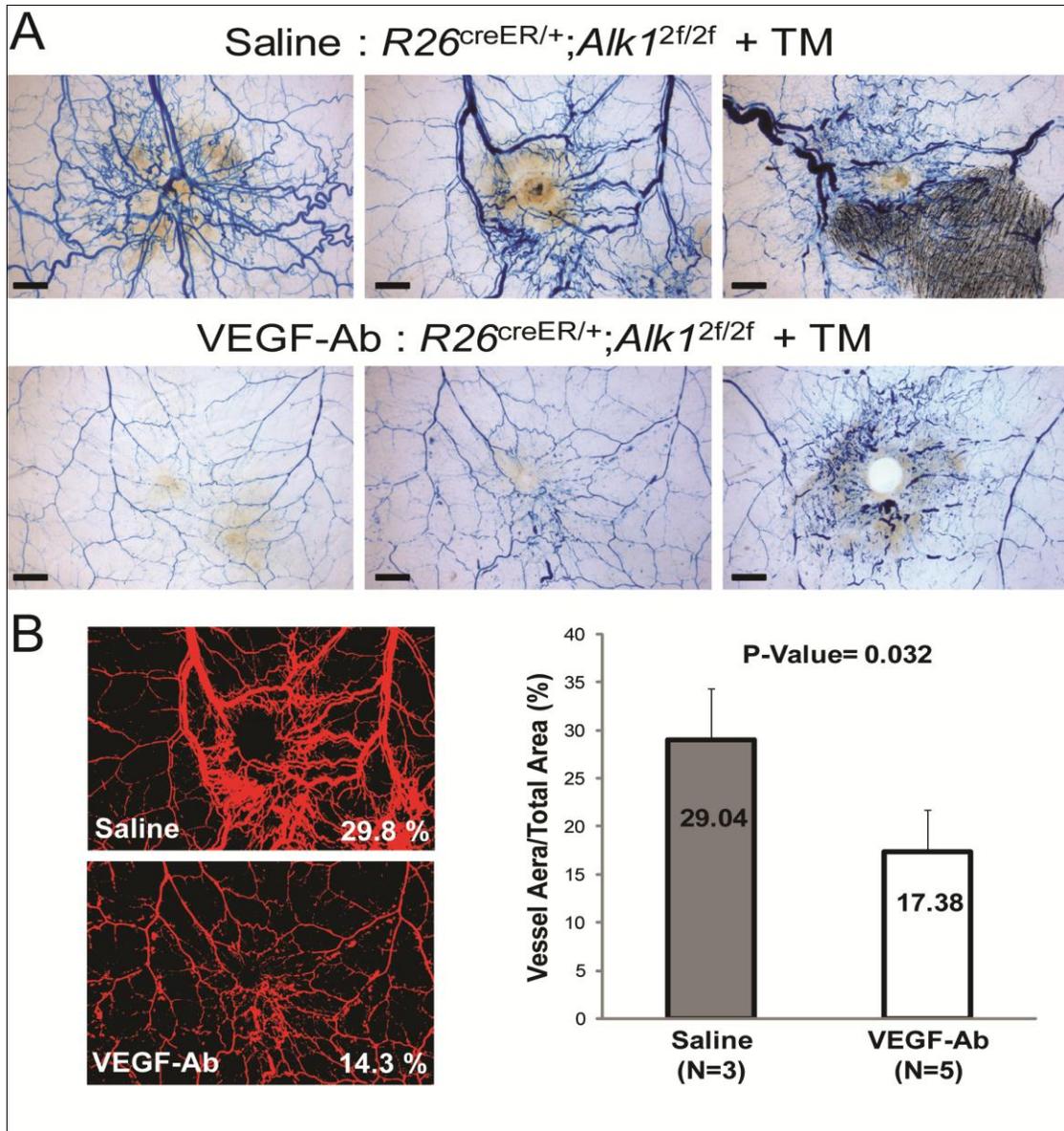


Figure 2-2. Angiogenesis blockade (VEGF-neutralizing antibody) suppressed wound-induced AVM formation in *Alk1*-deficient adult subdermal vessels. (A) Saline or VEGF-antibody (Ab) were injected into *Alk1*-deficient adult mice ($R26^{creER/+};Alk1^{2f/2f}$) to block angiogenesis. AVMs were induced by incisional wound of dorsal skin. VEGF-Ab suppressed wound-induced AVMs compared to saline-treated group. Tamoxifen (TM) was treated to effect *Alk1* deletion. VEGF-Ab (bottom panels) suppressed wound-induced AVMs compared to saline-treated animals (top panels). (B) The severity of AVMs formed surrounding wound sites was calculated as the percentage of vessel area containing latex dye (MATLAB program). Panels on the left are representative of one experimental comparison. The graph on the right is a compilation of all data showing a significant reduction in the percentage of vessel area in VEGF-treated group compared to the saline-treated group. Scale bars in each panel indicate 1 mm.

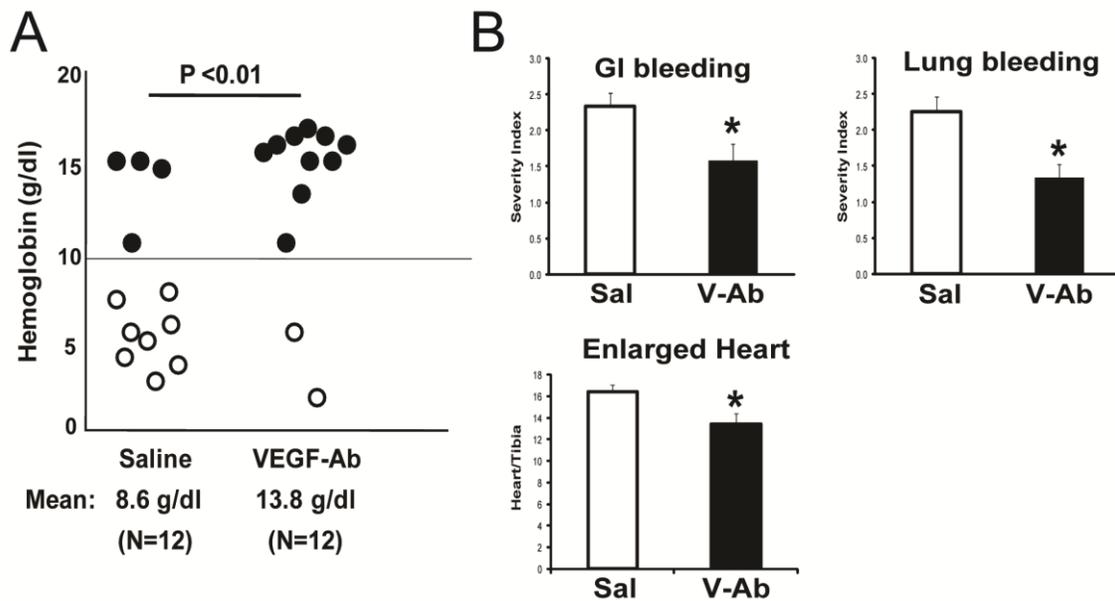


Figure 2-3. Angiogenesis blockade improved hemoglobin levels and visceral hemorrhage in *Alk1*-deficient adult mice. (A) Hemoglobin concentration was measured in saline-treated (n=12) and VEGF-treated (n=12) *Alk1*-deficient mice 9 days after tamoxifen treatment to effect *Alk1* deletion. VEGF-blockade improved hemoglobin levels in *Alk1*-deficient mice relative to the saline-treated group. (B) The severity of GI and lung bleeding (top) was expressed as a severity index based on direct observation and categorization. Hypertrophy of the heart (bottom) was expressed as a ratio of heart volume and tibia length ($\mu\text{l}/\text{cm}$). VEGF-blockade significantly suppressed related hemorrhaging and heart enlargement in *Alk1*-deficient mice.

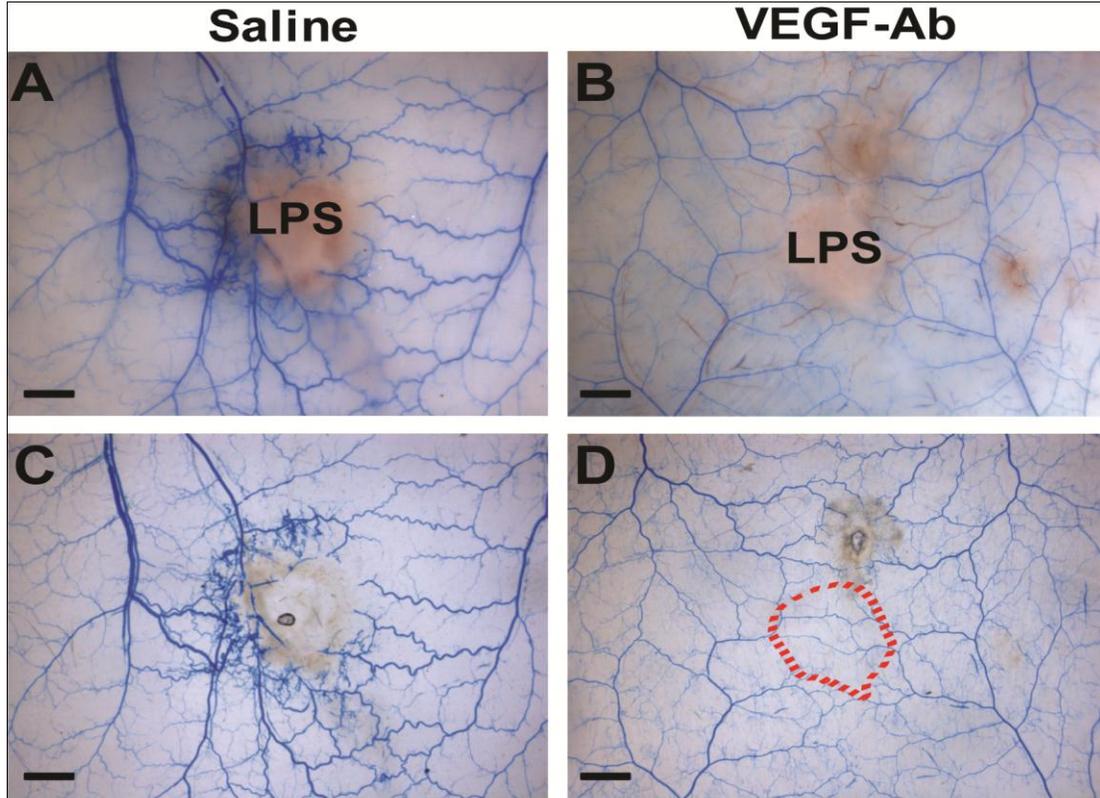


Figure 2-4. Angiogenesis blockade suppressed LPS-induced AVMs in *Alk1*-deficient mice. LPS particles formed aggregates on subdermal surfaces in both saline-treated (A) and VEGF-antibody-treated (B) *Alk1*-deficient mice. After skins were cleared to enhance the visualization of blood vessels, AVMs were apparent in saline-treated mice (C) but not mice treated with VEGF antibody (D). Scale bars in each panel indicate 1 mm.

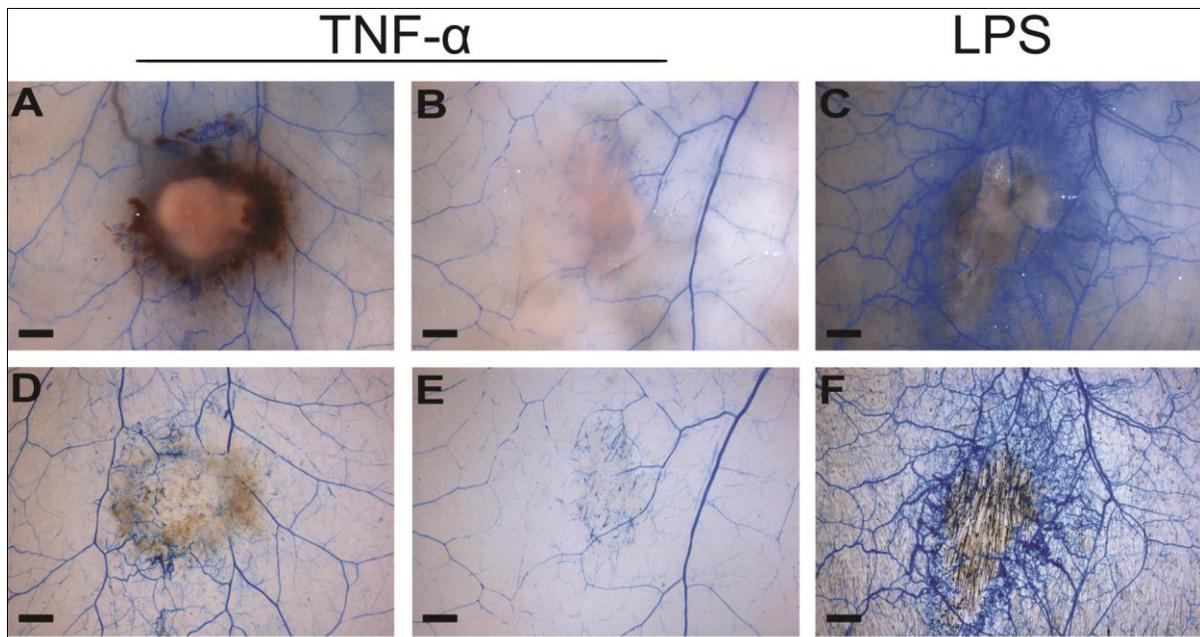
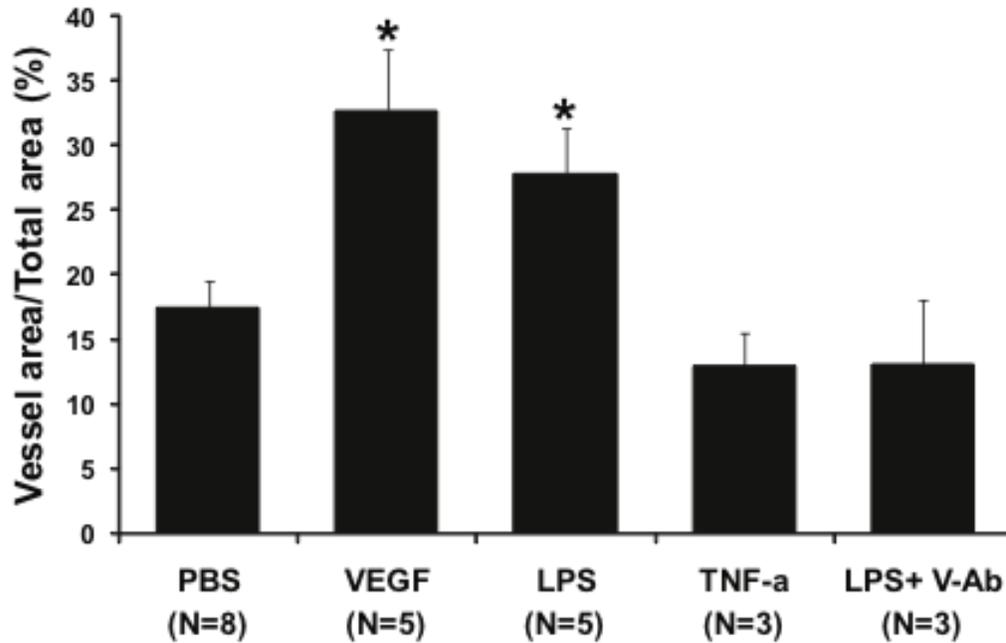


Figure 2-5. TNF- α induced mild AVM formation in *Alk1*-deficient mice. TNF- α PLGA particles were implanted under dorsal skin in tamoxifen-treated *Alk1*-deficient mice ($R26^{creER/+}; Alk1^{2f/2f}$). Vessels were observed after 7 days with the latex dye. TNF- α -treated mice (A, B and D, E) are compared with an LPS-treated mouse (C and F) before (A-C) and after (D-F) the clearing of aggregated particles. TNF- α induced neovascularization around particle aggregate. However, AVMs formed by TNF- α (D, E) were less numerous than those induced by LPS (F). Scale bars in each panel indicate 1 mm.



* : P <0.05 vs PBS

Figure 2-6. Quantification of AVM formation by various stimuli in Alk1-deficient subdermal vessels. VEGF and LPS-treated groups significantly promoted AVM formation compared to PBS-treated group. TNF- α did not significantly induce AVM formation in Alk1-deficient mice. LPS-induced AVM formation was suppressed by VEGF-neutralizing antibody (V-Ab).

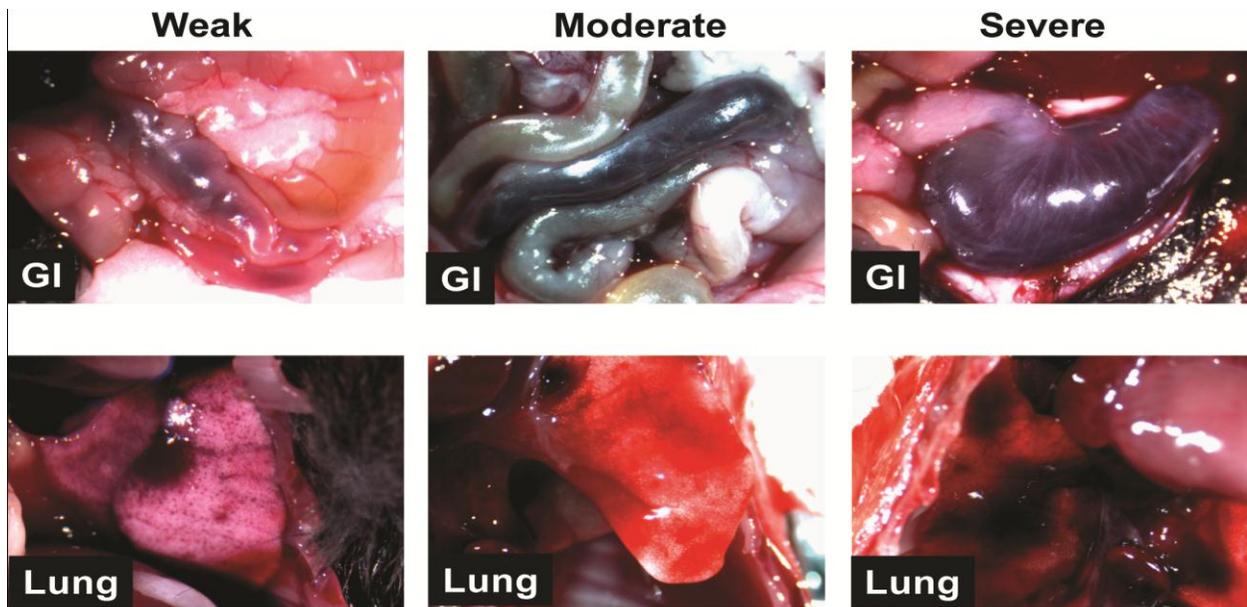


Figure 2-7. Lung (top) and GI bleeding (bottom) was categorized into three groups (weak, moderate, and severe) depending on the severity. Partial or focal hemorrhaging belonged to the weak level (left). Vast areas of hemorrhaging were categorized as severe (right). Intermediate levels of hemorrhaging were grouped as moderate (middle).

CHAPTER 3 MATERIALS AND METHODS

***Smad1*-Conditional Knockout Mice**

Generation of a conditional *Smad1* allele (*Smad1^f*) was previously described.²⁰⁸ Generation of the Tg(*Alk1-cre*)-L1 (L1cre) line has also been described.¹⁰⁹ Tg(*Tagln-cre*) and R26R mice were purchased from the Jackson Laboratory. *Smad1^{ff}* mice were intercrossed with L1cre or *Tagln-cre* lines. The L1cre(+);*Smad1^{f/+}* or L1cre(+);*Smad1^{+/2f}*;R26R males were further intercrossed with *Smad1^{ff}*;R26R females to produce L1cre(-);*Smad1^{ff}* and L1cre(+);*Smad1^{ff}*. *Tagln-cre*(-);*Smad1^{ff}* and *Tagln-cre*(+);*Smad1^{ff}* were produced by the same breeding scheme. More than half of the control and experimental mice contained the R26R allele to monitor the Cre activity. PCR primer sets and conditions for detecting the conditional as well as null alleles of *Smad1* were previously described.²⁰⁸ Primer sets for the genotype of L1cre, *Tagln-cre*, or R26R are shown in Table 1-1.

Hemodynamic Analysis

To evaluate pulmonary artery pressure, right ventricular systolic pressure (RVSP) was measured by right heart catheterization through the right jugular vein. Each mouse was anesthetized by isoflurane (1-2 %) and placed in the supine position. A 1-2 cm incision was made in the neck to expose the right jugular vein. The Mikro-Tip pressure transducer (SPR-835, Millar Instrument) was inserted into right external jugular vein and advanced into the right ventricle. Systemic blood pressure was recorded non-invasively using the tail-cuff method. A pneumatic pulse sensor was placed on the tail distal to an occlusion cuff controlled by a Programmed Electro-Sphygmomanometer (PE-300, Narco Bio-Systems), which is connected to the Powerlab system (ADInstrument). All

electrical outputs from the tail cuff, the pulse sensor and transducer were recorded and analyzed by the Powerlab 8/30 data acquisition system and associated Chart software (ADInstrument).

Right Ventricular Hypertrophy (RVH)

After hemodynamic analysis, mice were euthanized and using syringe-generated flow, the pulmonary circulation was perfused with PBS containing heparin (3 units/ml). The hearts were isolated and outflow tracts and atria were removed. The right ventricle was cut out from the heart by the spring scissor and the right ventricle and the remaining left ventricle (LV) plus septum (S) were weighed. Right ventricular hypertrophy was determined by the ratio of RV/LV+S.

Pulmonary Vessel Morphometry

After the hemodynamic analysis, the left lung was inflated with PBS for 20 minutes followed by formalin at a constant inflation pressure of 23cmH₂O, fixed with 4% paraformaldehyde overnight, and paraffin-embedded. Each lung sample was transversely sliced into 5 μ m thick sections and subjected to immunostaining with anti-smooth muscle α -actin (α SMA) antibody (Sigma-aldrich, mouse monoclonal, 1:800) using M.O.M kit (Vector laboratories) to visualize the vascular smooth muscle layer. To determine muscularization of pulmonary vessels, peripheral blood vessels ranging from 30-70 μ m in diameter were counted in at least four fields at 20X magnification with a Zeiss Axioplan-2 optical microscope. The counted vessels were categorized as fully muscularized (75-100% of medial layer covered by anti- α SMA staining), partially muscularized (1-74% of medial layer is covered by anti- α SMA staining), or nonmuscularized vessels at the level of alveolar ducts. The percentage of pulmonary vessels in each category was calculated by dividing the number of vessels by the total

number of counted vessels in the same field. To calculate the percentage of wall thickness (WT), circular and fully muscularized vessels were selected. WT1 (the thickness between the outer boundary and the inner boundary of α SMA positive medial layer) was measured at one point of the vessel wall and WT2 at the point which was diametrically opposite, guided by Openlab 5.03 Beta software (Improvision Inc.). External diameter (ED) was also measured at the same vessel. The percentage of medial wall thickness was calculated as $(WT1 + WT2) \times 100/ED$.

Establishment of Immortalized Pulmonary Endothelial Cells

Whole lungs were removed from an eight-week-old $R26^{CreER/+}; Bmpr2^{2f/2f}$ mouse and washed in HEPES followed by additional washing in DMEM. Lung tissues were finely minced using a sterile scalpel. The chopped tissues were subjected to serial digestion using 2 ml of a 1X trypsin solution [0.25% trypsin, 0.5 M EDTA (pH 8.0) in DMEM] at 37°C, with frequent shaking, for three times at 8 minutes each. Trypsin digestion was inactivated by adding 6 ml of normal endothelial cell media (ECM)[10% fetal bovine serum, 1 mg/ml heparine , 0.1 mg/ml endothelial mitogen (Biomedical technologies), 1 mM non-essential aminoacids (Cellgro), 1 mM sodium pyruvate, 50 units/ml penicillin/streptomycin]. After the large debris sinking down, the supernatant was carefully collected and plated into 2 wells of a 6-well culture plate. For immortalization, when the culture reached 50% confluency post isolation, they were transfected with 4.0 μ g of SV40 DNA: ATCC (VRMC-3), pUCSV40-B2E^{209, 210} using Lipofectamine (Invitrogen) following the manufacturer's protocol. After a couple of passages, the endothelial cells were sorted out by Fluorescence-Activated Cell Sorting (FACS) using Dio-Ac-LDL (Biomedical technologies) and lectin (Sigma-aldrich). For deleting the *Bmpr2* gene, the immortalized pulmonary ECs were plated on 6-well plates

(2×10^5 cells/well) and cultured with medium containing 1 μ M of 4-hydroxy tamoxifen (Sigma-aldrich) for 3 days. Thereafter the cells were cultured with 4-hydroxy tamoxifen-depleted growth medium. Primers (table 1) detecting null allele of *Bmpr2* (*Bmpr2*^{1f/1f}) were used to confirm the deletion of *Bmpr2* by tamoxifen.

Semi-Quantitative RT-PCR

To determine the levels of transcripts of endothelial cell-specific markers and genes involved in TGF- β superfamily signaling, *Bmpr2*-intact (*Bmpr2*^{2f/2f}) and *Bmpr2*-deleted (*Bmpr2*^{1f/1f}) pECs were harvested at 100% confluency from a 25 cm² culture flask. Total RNAs from the cells were extracted using the NucleoSpin RNA purification kit (Clontech). One μ g of RNA was used for reverse transcription (RT) reaction. The cDNAs were synthesized using SuperScript III First-Strand synthesis kit (Invitrogen). 2 μ l of cDNA was used as a template for PCR amplification for 25 cycles: denaturation at 94°C for 45 seconds, annealing at 60°C for 45 seconds and extension at 72°C for 1 minute. The transcription level of each gene was normalized to Gapdh expression. The primers used for RT-PCR analysis are shown in Table 1-1.

Western Blotting

Pulmonary ECs were harvested with a Chemicon lysis buffer containing 50 mM Tris (pH 6.8), 1 mM EDTA and 2% SDS and sonicated. Lysates were spun down at 13000 rpm for 15 minutes. The concentration of protein was determined using the Bio-rad DC Assay kit. 50 μ g of total protein was fractionated by 10% of SDS-PAGE and transferred to nitrocellulose membranes (Bio-rad). Membranes were incubated with the primary antibody followed by the horseradish peroxidase-linked secondary antibody. A chemiluminescent detection reagent (ECL PlusTM, Amersham Pharmacia Biotech Inc.) was used to visualize proteins. The antibodies used for Western blotting analysis are

the following: SMAD1 (rabbit polyclonal, 1:1000), SMAD2 (mouse monoclonal, 1:1000), pSMAD2 (rabbit polyclonal, 1:1000), pSMAD1/5/8 (rabbit polyclonal, 1:1000) from Cell Signaling, β -actin (mouse monoclonal, 1:10,000) from Sigma-aldrich, BMPR2 (mouse monoclonal, 1:500) from BD Transduction Laboratories, Secondary antibodies include: mouse (1:5000) and rabbit (1:5000) from Sigma-aldrich.

***Alk1*-Conditional Knockout Mice**

Establishment of the *Alk1*^{2f} allele in laboratory mice was described previously.¹⁰⁹ *R26*^{creER/+} allele-carrying mice were purchased from The Jackson Laboratory. By crossing *R26*^{creER/+} line with *Alk1*^{2f/2f} mice, *R26*^{creER/+}; *Alk1*^{2f/2f} mice were produced and 2-4 month-old mice were used in all experiments. The inactivation of the *Alk1* gene in adult mice was accomplished by treating animals as previously described.¹⁶⁹ All *in vivo* procedures were conducted in accordance with animal use guidelines established by the University of Florida Institutional Animal Care and Use Committee.

Preparation of PLGA Microparticles

Poly (d,l lactide-co-glycolide) (PLGA), 50:50 composition with inherent viscosity 0.55-0.75 dl/g in hexafluoroisopropanol, HFIP (Lactel) was used to generate the particles. The emulsion stabilizer, poly-vinyl alcohol (PVA) (MW ~ 100,000 g/mol) was purchased from Fisher Science. Phosphate buffered saline (PBS) solution (Hyclone) was used as the aqueous phase to form the emulsions while methylene chloride (Fisher Scientific) was used as an organic solvent to dissolve PLGA polymer. Microparticles were formed using a standard water-oil-water solvent evaporation technique.^{211, 212} Briefly, a 3% solution of PLGA polymer in methylene chloride was generated. 1 ml of 3% PLGA solution was emulsified with 10 μ l of 10 mg/ml Rhodamine 6G (Sigma-Aldrich) in PBS, 100 μ l of 1% bovine serum albumin (Fisher Scientific) in PBS, and either 100 μ l of 100

µg/ml VEGF (R&D systems) in PBS for VEGF-PLGA, 100 µl of 100 µg/ml TNF-α (R&D systems) in PBS for TNF-α-PLGA or 100 µl of 1 mg/ml LPS (Sigma-Aldrich) in PBS for LPS-PLGA at 26,500 rpm for 60 seconds using a tissue-miser homogenizer (Fisher Scientific) to form the primary emulsion. The primary emulsion was added to 10 ml of 5% PVA solution in PBS and the homogenizing was continued at 19,500 rpm for 120 seconds to form the secondary emulsion. Then, the secondary emulsion was added to 100 ml of 0.5% PVA made in dH₂O. The particles thus formed were agitated using a magnetic stirrer for 16 hours to evaporate residual methylene chloride. The remaining solution was centrifuged at 10,000 rpm for 10 minutes to collect microparticles which were subsequently washed three times with PBS. The particle suspension in 5 ml of PBS was then flash-frozen in liquid nitrogen and stored at -20⁰C until used.

Injection of PLGA Particles into Subdermal Area

The recipient mice (*R26^{creER/+};Alk1^{2f/2f}* and *R26^{+/+};Alk1^{2f/2f}*) were anesthetized by placing them within an induction chamber and introducing 4.0% isoflurane, and out of the chamber, anesthesia was maintained by 2.0%–3.0% isoflurane using a nose cone. The hairs on the back of the mice were shaved and 100 µl of PLGA particles containing 200 ng of VEGF, 2 µg of LPS, or 200 ng of TNF-α were implanted into the subdermal area of the mid-dorsal skin using 29-gauge 0.3-ml syringe (Monoject). Tamoxifen (TM, Sigma-aldrich) was injected intraperitoneally at a dose of 2.5 mg/25 g of body weight for *Alk1* gene deletion. Then, 0.6 ml of the blue latex dye (Connecticut Valley Biological Supply Co.) was injected via the left heart 7 days after TM injection to visualize the blood vessels.

Skin Wound Generation and VEGF-Antibody Treatment

The recipient mice ($R26^{creER/+};Alk1^{2f/2f}$) were anesthetized as described above. The hairs on the back of the mice were shaved and one 2 mm-diameter full-thickness excisional wounds were inflicted on the mid-dorsum using a 2 mm-dermal biopsy punch (Miltex). The wounds were left unsutured, and betadine was applied to the wounds. No analgesics or antibiotics were given in order not to affect the wound healing process. TM was injected intraperitoneally at a dose of 2.5 mg/25 g of body weight and VEGF-neutralizing antibody (100 μ l, 5 mg/kg of body weight, Genentech) or the same volume of saline was serially injected on the first, 4th, and 7th day of post-wounding. On the 9th day of post-wounding, 0.6 ml of the blue latex dye was injected via the left heart.

Latex Dye Injection and Image Processing

Mice were anesthetized with intraperitoneal injection of ketamine/xylazine (100 mg/15 mg/kg of body weight) and abdominal and thoracic cavities were opened. To perfuse all the blood out, heparin (200 unit/ml, Sigma-aldrich) in 10 ml PBS was infused through the left ventricle of the heart at 120 ml/hour using a syringe pump (KD scientific) after a small cut was made on the left atria. Then, 10 ml of PBS-dilator mixture containing heparin (10 unit/ml, Sigma-Aldrich), papaverine (0.04 mg/ml, Sigma-Aldrich), and sodium nitroprusside (100 μ M, Sigma-aldrich) was infused through the same hole made on the left ventricle to maximize dilation of blood vessels. To fix the dilated vessels, 10 ml of 10% formalin was infused through the same hole on the left ventricle. Then, 0.6 ml of the blue latex dye was slowly and steadily injected into the hole of left ventricle with a 26-gauge 1-ml syringe. Then, the hair removal cream (Veet) was put on the shaved back skin of dye-injected mice and the hairs were completely removed after 10 minutes incubation. The mice were washed briefly in water and fixed with 10%

formalin overnight. After fixation, the dorsal skin was peeled off, stretched, and flattened on the Styrofoam. The flattened skin was dehydrated by methanol series (20%, 50%, 75%, and 100%, each for 20 minutes) and cleared with organic solvent (benzyl alcohol/benzyl benzoate, 1:1; Sigma-aldrich). Blood vessels containing the latex dye in the cleared skin were imaged via a CCD camera (Leica) and processed by MATLAB (MathWorks) as described previously.¹⁶⁹

Hemoglobin Concentration and Hemorrhage Index

2-3 drops of blood were collected from the tails of VEGF-antibody-treated (N=12) and saline-treated mice (N=12) and soaked up by microcuvettes (STANBIO Laboratory). The concentration of hemoglobin was measured by the hemoglobin photometer (Hemopoint H2, STANBIO Laboratory) using the blood-soaked microcuvettes. Then, after anesthesia with ketamine/xylazine, abdominal and thoracic cavities were opened and hemorrhages of lungs and gastrointestinal tract and the size of the heart were observed and categorized into three levels: weak, moderate, and severe conditions according to the degree of severity (Figure 2-6). Hemorrhage levels were graded: 1.0 for weak, 2.0 for moderate, and 3.0 for severe level. The mean values were expressed as severity index.

Hypertrophy of the heart

After latex-dye injection through the heart, hearts were taken and all the blood vessels connecting to the heart were removed. Hearts were submerged in 1.5 ml tube containing 500 μ l PBS and increased volume of PBS was measured as a volume of heart. Right legs of mice were cut out in the middle of femur and were skinned off and were put in 1M NaOH overnight for digestion of tissues surrounding bones. After digestion, tissues were easily removed and the foot bone and broken femur were

separated from the tibia by slowly pulling them. The length of whole tibia was measured by the caliper.

Statistical Analysis

T-test was used to determine a statistical significance between two groups, and one-way Anova was used for more than 3 groups, and multiple pairwise comparisons were made by post-hoc tests (Tukey) using the Sigmaplot.

CHAPTER 4 CONCLUDING REMARKS AND FUTURE DIRECTIONS

Current therapeutic interventions for both PAH and HHT diseases are restricted to alleviate severe symptoms because they are not based on mechanistic pathology. There is a pressing need for more effective and pathology-specific therapies.

In PAH studies, I showed that loss of SMAD1 function in ECs or SMCs predisposes mice to pulmonary hypertension and Impaired BMP signaling makes TGF- β signaling overactivated and leads to an imbalance of TGF- β and BMP signaling in pECs, which might be a critical factor for PAH development. From these results, I can suggest several therapeutic targets. First, reduced SMAD1 activity may be associated with PH development. Therefore, inhibiting phosphatase activity could be effective to maintain SMAD1 phosphorylation and its activity although it should be specific to SMAD1/5/8, not SMAD2/3. Second, recovering from imbalance state of TGF- β /BMP signaling would be therapeutic for PAH patients by suppressing overactivated TGF- β . TGF- β antagonists such as the angiotensin II type 1 receptor (AT1) blocker, Losartan¹³³ or the activin receptor-like kinase 5 inhibitor, SB525334²¹³ may have therapeutic benefit. In addition to this, I found that BMP7 response was intact in *Bmpr2*-deleted pECs. Thus, treatment of BMP7 ligand could compensate for reduced level of pSMAD1/5/8 from BMPR2 deficiency.

To determine whether overactivation of TGF- β derived from reduced BMP response is a critical factor for PAH development *in vivo*, we are planning to test whether inhibition of TGF- β signaling prevent PAH manifestation in our PH mouse models-*Smad1* or *Bmpr2* cKO mice by additionally deleting genes involved in TGF- β

signaling such as *Tgfb2* and *Smad4* or treatment with inhibitors blocking TGF- β signaling, such as Activin Receptor Like Kinase 5 (ALK5) inhibitor.¹⁰³

Endothelial *Smad1* deletion did not result in 100% penetrance in PH, indicating that environmental or genetic second hits are required for PAH development. As shown in Figure 1-1, I observed the marked reduction of SMAD1/5/8 phosphorylation in the lungs of PH mice, suggesting that the complete loss of SMAD1 activity in the whole lung may be essential to manifest PAH phenotype. Thus, by mating *L1cre(+);Smad1^{fl/fl}* with *Tagln-cre(+);Smad1^{fl/fl}*, we can delete *Smad1* gene both in endothelial and smooth muscle cells. Although these double cre KO mice did not block SMAD1 activity in the whole lung, we can expect synergistic effect of smooth muscle thickening and endothelial dysfunction and may expect increase of penetrance.

In HHT studies, I demonstrated that inflammation or angiogenesis are sufficient to induce AVMs as second hits for ALK1 deficiency. In terms of therapeutic aspect, antagonizing inflammatory or angiogenic pathway may be effective for HHT patients. I showed that VEGF-neutralizing antibody can prevent AVM formation and internal bleeding in *Alk1*-deficient adult mice. Even though effective anti-inflammatory reagents are not available, IL-9-neutralizing antibody could be a useful reagent to test because IL-9 is essential factor for inflammation-induced angiogenesis.

Up to now, many studies have focused on assessing increased response to angiogenic stimulation in *Alk1*-deleted endothelial cells. However, what makes this increased sensitivity as a result of *Alk1* deficiency remains unclear. To address this question, we have many useful resources: pulmonary endothelial cells (pECs), mice carrying *Alk1^{2f/2f}* and *Alk1^{1f/1f}* allele respectively, and microarray data from pECs

(*Alk1*^{2f/2f} and *Alk1*^{1f/1f}). I showed VEGF signaling is required for AVM formation with *Alk1*-deficiency. With above materials, examining whether *Alk1* can suppress downstream of VEGF signaling or modulate interaction of VEGF and VEGF receptors and other possible hypotheses, would be interesting and invaluable research to find novel therapeutic targets.

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BIOGRAPHICAL SKETCH

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