#### 1 **Research Letter** 2 Decreasing ELK3 expression improves Bone Morphogenetic Protein Receptor 2 signaling 3 4 and pulmonary vascular cell function in PAH 5 Md Khadem Ali<sup>1,2</sup>, Lan Zhao<sup>1,2</sup>, Vinicio de Jesus Perez<sup>1,2</sup>, Mark R. Nicolls<sup>1,2</sup>, Edda F. 6 Spiekerkoetter<sup>1,2,\*</sup> 7 8 9 <sup>1</sup>Department of Medicine, Division of Pulmonary, Allergy and Critical Care Medicine, Stanford 10 University, Stanford, CA, USA 11 <sup>2</sup>Vera Moulton Wall Center for Pulmonary Vascular Disease, Stanford University, Stanford, CA, 12 USA 13 14 \*Corresponding author: 15 Edda Spiekerkoetter, MD 16 Associate Professor of Medicine 17 Department of Medicine, Pulmonary and Critical Care Medicine 18 Vera Moulton Wall Center for Pulmonary Vascular Disease 19 Stanford University 20 1701 Page Mill Road, Palo Alto, CA 94304, USA 21 Phone: +1 (650) 724-1493 22 Email: eddas@stanford.edu 23 24 **Current word count:** 1265 25 Number of Figures: 02 26 Number of Tables: 0 27 28 Support statement: This research was supported by funding from the National Institutes of 29 Health (R01 HL128734), Stanford Vera Moulton Wall Center for Pulmonary Vascular Diseases, 30 the U.S. Department of Defense (PR161256), and Stanford Translational Research and Applied

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Author contribution: MKA and ES conceptualised the study design. MKA performed the
 experiments and data analysis. All authors contributed to data collection, data interpretation,
 writing, and editing of the manuscript. ES: fund acquisition and supervision.

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#### 37 Abstract:

38 ELK3 is upregulated in blood and pulmonary vascular cells of PAH patients and may play a
39 significant role in PAH potentially through modulating BMPR2 signaling.

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### 41 Introduction:

42 Pulmonary arterial hypertension (PAH) is a rare but complex, severe, life-threatening condition 43 of the small pulmonary arteries. PAH is characterized by pulmonary arterial remodeling and 44 increased pulmonary vascular resistance; the consequent elevated pulmonary arterial pressures 45 causes right ventricular afterload and, ultimately, failure. The disease remains incurable despite 46 intense efforts to identify new therapies to treat PAH patients. In order to identify new therapies, 47 it is crucial to understand the cause and exact molecular mechanisms of the disease. Although the 48 exact cause of PAH is unclear, many genetic, epigenetic, and environmental factors have been 49 shown to contribute to the development and progression of the disease. For example, 50 heterozygous loss of function mutation in bone morphogenic protein receptor 2 (BMPR2) occurs 51 in 53-86% of familial PAH patients and 14-35% in sporadic idiopathic PAH patients(1). 52 However, the disease penetrance rate of the mutation carriers is low, indicating that other 53 unidentified factors may contribute to the disease development in addition to the gene mutation. 54 Significantly, BMPR2 signaling is thought to be impaired in PAH patients regardless of the 55 etiology of PAH, making the signaling a master switch in the disease. Previously, we and others 56 have shown that targeting BMPR2 signaling with repurposed drugs (FK506, Enzastaurin) or 57 rebalancing the pathway with BMP9 and Sotatercept improved PAH in animal models and pilot 58 studies in patients (1).

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To identify BMPR2 signaling modifier genes, our team previously performed an siRNAmediated high throughput screening (HTS) of ~22,000 genes in a BRE-ID1 incorporated mouse myoblastoma reporter cell line, which yielded two important novel BMPR2 modifier genes

63 (FHIT, LCK) in PAH (2). In the HTS data set, we also found that E26 transformation-specific 64 transcription factor (ELK3) knockdown decreased Id1 levels. Surprisingly, our validation 65 experiments in human pulmonary arterial endothelial cells (PAECs) and pulmonary arterial smooth muscle cells (PASMCs) showed that ELK3 modulated BMPR2 signaling in the opposite 66 67 direction compared to the HTS experiments using the mouse myoblastoma cell line. ELK3 68 knockdown increased BMPR2 signaling. ELK3, also known as NET/SAP-2/ERP, is a 69 transcription factor which can form a ternary complex with serum response factor and DNA. 70 While ELK3 generally acts as a transcriptional repressor, it can also work as a transcriptional 71 activator when phosphorylated by the Ras/mitogen-activated protein kinase signaling pathway. 72 ELK3 regulates various biological processes in health and disease, such as proliferation, 73 apoptosis, migration, and angiogenesis. Elevated expression of ELK3 plays a significant role in 74 accelerating the progression and metastasis of different cancers, such as prostate, breast, bladder, 75 gastric, and liver cancer. A significant upregulation of ELK3 expression was also observed in rat 76 carotid arteries following balloon-injury and in human plaques (3). ELK3 was also found to 77 attenuate angiogenesis in VEGF-induced angiogenesis assays in vitro and in vivo (4). While the 78 role of ELK3 has been studied in different cancers and cardiovascular diseases, its role in PAH is 79 unknown.

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#### 81 **Results and Discussion:**

82 We first checked whether ELK3 expression is dysregulated in the whole blood of PAH patients 83 using a large RNA sequencing (RNAseq) data set comprising 359 patients with idiopathic, 84 heritable, and drug-induced PAH as well as 72 age- and sex-matched healthy (5). We found a 85 significant upregulation of ELK3 in the blood of PAH patients compared to healthy controls 86 (Figure 1A). We next tried to determine whether the upregulation of ELK3 in peripheral blood is 87 mirrored by an ELK3 upregulation in the PAH pulmonary vasculature. Thus, we re-analyzed 88 publicly available RNAseq data sets generated from two critical pulmonary vascular cell types in 89 PAH, PAECs and PASMCs of PAH patients and healthy controls. We found a significant 90 increase in ELK3 expression in the PASMCs of 4 idiopathic (I)PAH patients compared to 4 91 healthy control PASMCs (Figure 1B, GSE144274). We also measured the expression of ELK3 92 in PASMCs of a different cohort of 4 healthy controls and 3 PAH patients by quantitative reverse 93 transcription PCR (qRT-PCR) but did not observe a significant change in ELK3 expression

94 between the two groups (Figure 1C). Possible confounding factors are low ELK3 mRNA 95 expression, low sample size and different PAH etiologies, which warrants further validation in a 96 larger cohort. Previously, several studies showed increased expression of ELK3 in PAECs of 97 patients. In a single-cell RNAseq analysis study of lung tissues from 6 healthy controls and 4 98 IPAH patients, ELK3 expression was shown to be up-regulated in endothelial cells of IPAH 99 patients compared to healthy controls(6). As another example, Reyes-Palomares and colleagues 100 identified that the transcriptional activity of ELK3 was more active in PAH patients than healthy 101 controls (7). We therefore analyzed ELK3 expression in PAECs of PAH patients using a publicly 102 available RNAseq data set comprising 9 healthy controls and 8 PAH patients (GSE0126262). We 103 did not observe a significant change in ELK3 expression in PAECs of PAH patients in this 104 RNAseq dataset (Figure 1D). The increased detection of ELK3 in blood of PAH patients could 105 not be attributed to PASCMs and PAECs in our relatively small sample size, given that these 106 cells could be the source of ELK3 in blood but that was not demonstrated in this study.

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108 ELK3 is predicted to be connected with the inhibitor of DNA (ID) signaling pathway 109 (wikipathways). Furthermore, homozygous deficiency in ELK3 was shown to up-regulate 110 expression of SERPINE1 (Serpin family E member 1), also called plasminogen activator 111 inhibitor 1 (PAI-1) in prostate cancer cells(8). PAI-1 is a known downstream target of the BMP 112 signaling pathway and is strongly linked to PAH(9). A recent integrated bioinformatic analysis 113 revealed SERPINE1 (PAI-1) as one of the most significant markers in PAH(10). We therefore 114 hypothesized that the observed increase in ELK3 might downregulate BMPR2 signaling and 115 thereby be involved in PAH pathogenesis. We further hypothesized that decreasing ELK3 might 116 do the opposite – increase BMPR2 signaling – and thereby might be beneficial in PAH. We 117 conducted experiments in PAECs subjected to ELK3 or non-target siRNA with and without BMP9 stimulation. We demonstrated that ELK3 silencing further increased BMP9-induced 118 119 phospho-SMAD1/5/9 levels measured by western blot (Figures 2A-B). ELK3 protein 120 concentration is very low in PASMCs, as seen in **Figure 2C**, and therefore we used qRT-PCR to 121 assess the effect of ELK3 knockdown on BMPR2 signaling. We found a significant increase in 122 BMPR2 as well as ID1 expression, a downstream target of the BMPR2 signaling, following the 123 knockdown of ELK3 with siRNA in PASMCs, while BMPR2 knockdown did not change the 124 ELK3 expression (Figures 2D-G), suggesting ELK3 to be upstream of BMPR2. We also

observed that inhibition of ELK3 decreased PASMCs proliferation (Figure 2H), which is in line
with the proposed model that ELK3 inhibition improves BMPR2 signaling and function in
PAEC and PASMC (Figure 2I).

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Previous studies suggested that ELK3 is downregulated during hypoxia, releasing repression of several genes and leading to increased expression of Egr1 and VEGF, as well as PHD2, PHD3, and Siah2 destabilizing HIF1 $\alpha$  (6). Therefore, its enhanced expression in PAH is surprising and might be expected to inhibit angiogenesis. Further studies need to explore the exact role and molecular mechanisms of ELK3 in PAH.

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This study has several limitations. *First,* we were not able to validate the RNAseq expression of ELK3 by qRT-PCR. *Second,* while ELK3 inhibition increases BMPR2 signaling, it is critical to explore whether overexpression of ELK3 does the opposite, that is, inhibits BMPR2 expression and signaling and induces PAH. *Third,* the molecular mechanisms of how ELK3 regulates BMPR2 signaling, and PAH still need to be clarified.

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#### 141 **Conclusion:**

In summary, through preliminary experimental and clinical sample analysis, we uncovered ELK3 as a clinically meaningful BMPR2 signaling modulator that influences pulmonary vascular cell function. Further studies are required to fully elucidate the role and molecular mechanisms of ELK3 expression in the pathogenesis of PAH.

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## 147 **Conflict of Interest statement:**

148 The authors declared no conflict of interest exists.

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# 157 Author contribution:

158 MKA and ES conceptualised the study design. MKA performed the experiments and data

analysis. All authors contributed to data collection, data interpretation, writing, and editing the

- 160 manuscript. ES: fund acquisition and supervision.
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194 Figure 1. ELK3 is upregulated in the blood and PASMCs of PAH patients. A) RNAseq

analysis of the whole blood collected from the 72 healthy controls (HC) and 359 patients with

196 IPAH, APH or HPAH showed a significant increase in ELK3 expression in PAH compared to 197 HC. For detail subject characteristics and hemodynamic data, please see (5). B) ELK3 expression 198 was found to be upregulated in PASMCs of IPAH patients in a publicly available RNAseq data 199 seta comprising 4 HC and 4 IPAH patients (GSE144274). C) ELK3 expression was not altered in 200 PASMCs of a small cohort of PAH patients by qRT-PCR. D) ELK3 expression was not changed 201 in PAECs of a small cohort of PAH patients (GSE126262). Wilcoxon rank sum test with 202 continuity correction was performed to compare ELK3 expression in the whole blood RNAseq 203 data. Student t-test was used to compare ELK3 expression in the PASMCs of PAH RNAseq 204 data. \*\*P<0.01. TPM, transcripts per million.

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## 206 Figure 2. ELK3 is involved in regulating BMPR2 signaling and PASMCs proliferation. A-

207 B) Inhibition of ELK3 with siRNA increased BMP9-induced pSMAD1/5/9 in PAECs, as 208 measured by western blot, 48h knock down, 2h BMP9 stimulation. C) Western blot verification 209 of ELK3 knockdown with siRNA in PASMCs. D-G) siRNA-mediated knockdown of ELK3 210 increased ID1 levels in PASMCs. BMPR2, ID1 and ELK3 levels were measured by qRT-PCR 211 following 72 hours of ELK3 or BMPR2 knockdown with 60nM siRNA and 3ul RNAimax. Data 212 are represented as mean +/- standard error mean (n=6/group). Student t test, \*\*P<0.01. H) 213 Silencing of ELK3 decreases hPASMCs proliferation as measured by MTT assay. Student t test, 214 \*\*\*\*P<0.0001. I) Proposed model for how ELK3 regulates BMPR2 signaling and PAH.

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#### **Figure 1**



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- 235 **Figure 2**



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