- 11. Spagnolo P, Balestro E, Aliberti S, Cocconcelli E, Biondini D, Casa GD, et al. Pulmonary fibrosis secondary to COVID-19: a call to arms? Lancet Respir Med 2020;8:750–752.
- 12. Chang YC, Yu CJ, Chang SC, Galvin JR, Liu HM, Hsiao CH, et al. Pulmonary sequelae in convalescent patients after severe acute respiratory syndrome: evaluation with thin-section CT. Radiology 2005;236:1067–1075.

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In Search of "Hepatic Factor": Lack of Evidence for ALK1 Ligands BMP9 and BMP10

To the Editor:

Considerable evidence suggests that the liver produces or modifies a circulating factor critical for preventing pulmonary arteriovenous malformations (PAVMs). In hepatopulmonary syndrome, liver dysfunction is associated with hypoxemia secondary to PAVMs, and PAVMs are reversed by liver transplantation (1). Additionally, portosystemic shunts that allow gut venous effluent to bypass the liver lead to PAVMs, which resolve when the shunt is closed (2). Further evidence comes from single-ventricle patients who undergo a threestaged surgery to relieve hemodynamic burden on the heart and correct oxygen desaturation. The second-stage surgery, the bidirectional Glenn, directs passively draining venous return from the superior vena cava (SVC) to the pulmonary circulation, with venous return from the inferior vena cava (IVC) pumped to the systemic circulation. Although the Glenn effectively decreases ventricular hemodynamic stress, intrapulmonary arteriovenous shunting is pervasive, and up to 25% of Glenn patients develop clinically significant hypoxemia secondary to diffuse PAVMs (3, 4). Although early theories of PAVM development focused on the absence of pulsatile flow or increased lower lobe perfusion (3), later evidence implicated the exclusion of a liver-derived substance from the pulmonary vasculature. This "hepatic factor" was postulated based on correlation between laterality of PAVMs and laterality of exclusion of hepatic venous effluent (5), and its existence is strongly supported by evidence that the third-stage Fontan procedure (completion of the total cavopulmonary anastomosis), which reroutes IVC flow to the lungs without restoring pulsatility, is strongly associated with PAVM regression (6). Despite the strong evidence for hepatic factor, its identity remains unknown.

Approximately 80% of PAVMs are associated with hereditary hemorrhagic telangiectasia, a genetic disorder caused primarily by mutations in BMP (bone morphogenetic protein) receptors ENG (endoglin) and ACVRL1 (activin A receptor like type 1, which encodes ALK1) (7). This pathway is active in lung endothelium, and ligands include BMP9 and BMP10 homodimers and BMP9/10 heterodimer $(8, 9)$. Both BMP9 and BMP10 are transcribed in hepatic stellate cells (9). Given the strong relationship between PAVMs and

hereditary hemorrhagic telangiectasia, the hepatic origins of BMP9 and BMP10, and evidence of decreased plasma BMP9 in hepatopulmonary syndrome (10), we hypothesized that ALK1 ligands may be the hepatic factor required for PAVM prevention. We expect that hepatic factor is either labile or actively removed from circulation on first pass through the systemic circulation, making it unavailable to the lung vasculature in Glenn circulation. Accordingly, in normal circulation, we hypothesized that concentrations of ALK1 ligands would be higher in the right atrium and pulmonary artery compared with the SVC and infrahepatic IVC. Some of the results of these studies have been previously reported in the form of an abstract (11), and some have been previously reported in the form of a preprint [\(https://](https://doi.org/10.1101/2020.07.09.20148320) doi.org/10.1101/2020.07.09.20148320).

Methods

This study was approved by the University of Pittsburgh Institutional Review Board. Participants undergoing clinically indicated cardiac catheterization were recruited between September 2015 and February 2017 and provided informed child assent and/or parental consent. Patients with bidirectional Glenn, prior to Fontan, were compared with two-ventricle control subjects. Excluded diagnoses among control subjects included single ventricle physiology, unrepaired complex congenital heart disease, and large shunt lesions. Patients with liver disease, anemia (Hb \leq 8 g/dl), cardiac surgery within 30 days, or transfusion within 48 hours were excluded from both cohorts.

We collected 1 ml blood in K_2EDTA tubes from five sites: the right atrium, pulmonary artery, aorta, SVC, and infrahepatic IVC. We measured ligands in duplicate in 30 μ l of plasma via sandwich ELISAs (R&D Systems) using DY3209 (BMP9), MAB2926 and BAF3956 (BMP10), and MAB2926 and BAF3209 (BMP9/10), with in-house generated recombinant proteins for the latter two standard curves. We fit data to a four-parameter logistic curve and performed statistical analysis using GraphPad Prism. We ran all samples from an individual on a single plate, and the operator was blinded to sample identity. Sample volume limitations prevented us from assaying all ligands in every individual.

Results

Diagnoses in 38 control subjects (mean age, 5.8 yr [4 mo to 12.6 yr]; 21 males, 17 females) included small shunt lesions (21), repaired forms of congenital heart disease with two-ventricle physiology (11), vascular stenosis (5), valvar obstructive lesions (2), and hypertrophic cardiomyopathy (1). Primary cardiac diagnoses in nine Glenn cases (mean age, 2.9 yr [range, 22 mo to 5.1 yr]; 7 males, 2 females) included variants of hypoplastic left heart syndrome (5), pulmonary atresia with intact ventricular septum (2), double outlet right ventricle with pulmonary atresia (1), and heterotaxy with right atrial isomerism (1).

BMPs are generated as proprotein dimers that are cleaved between the N-terminal prodomains and C-terminal growth factor domains, releasing the disulfide-bonded GFD (growth factor dimer). In control plasma, we detected BMP9 GFD, BMP10 proprotein, and BMP9/10 GFD (Figure 1) but not BMP10 GFD (DY2926; R&D Systems; data not shown). However, we found no differences in plasma concentrations of any ALK1 ligand when comparing withinsubject values across the right atrium, pulmonary artery, aorta, SVC, and IVC (Figure 1). This result suggests that these ligands are neither particularly labile nor actively removed on first pass through the systemic or pulmonary circulation, failing to support the hypothesis that they represent the hepatic factor required to prevent PAVMs.

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Figure 1. Plasma BMP9, BMP10, and BMP9/10 concentrations do not differ between pulmonary inflow and systemic venous circulation. Plasma from control subjects was sampled from the right atrium, pulmonary artery, aorta, superior vena cava, and infrahepatic inferior vena cava and ligands quantified by sandwich ELISA. All values are growth factor dimer equivalents. Colored lines connect samples from a single patient. The following were not significant by repeated-measures one-way ANOVA: BMP9, $n = 31$, $P = 0.13$; BMP10, $n = 31$, $P = 0.17$; and BMP9/10, $n = 21$, $P = 0.21$. IVC = inferior vena cava; PA = pulmonary artery; RA = right atrium; SVC = superior vena cava.

Evaluation of plasma from Glenn cases similarly revealed no significant differences in within-subject ligand concentrations across sampling sites (data not shown). However, after collapsing data across sites, comparison of grand means revealed significant decreases in plasma concentrations of all ligands in Glenn cases compared with control subjects (Figure 2), and significance persisted after age adjustment (multiple linear regression; BMP9,

 $P = 0.04$; BMP10, $P = 0.0002$, BMP9/10, $P = 0.002$). Although our sample set is underpowered, we saw no correlation between ligand concentration and PAVMs (Figure 2).

Discussion

We found no within-subject differences in plasma concentrations of BMP9 GFD, BMP10 proprotein, or BMP9/10 GFD across different

Figure 2. Plasma BMP9, BMP10, and BMP9/10 concentrations are lower in Glenn cases compared with control subjects. Within-patient data were averaged across all sampling sites (right atrium, pulmonary artery, aorta, superior vena cava, and inferior vena cava) for control subjects and Glenn cases and evaluated by Welch's t test. Error bars represent SD. BMP9: control, $n = 31$; Glenn, $n = 5$; $P = 0.02$. BMP10: control, $n = 31$; Glenn, $n = 7$; $P = 0.0006$. BMP9/10: control, $n = 21$; Glenn, $n = 6$; $P = 0.01$. * $P < 0.05$ and ** $P < 0.001$. Open circles indicate cases with pulmonary arteriovenous malformations.

sampling sites, in agreement with a recent report regarding BMP9 GFD (12) and failing to support the idea that ALK1 ligands are the "hepatic factor" required to prevent PAVMs. However, it remains possible that liver-derived BMP9 or BMP9/10 proproteins (not assayed) may exhibit site-dependent concentration differences or that enzymes required to cleave proproteins are unavailable in the Glenn circulation. Surprisingly, we found that Glenn cases had significantly lower concentrations of all three ligands compared with control subjects. Measurement of these ligands in additional Glenn cases and in Fontan cases will be required to determine the biological significance of this finding with respect to Glenn-associated PAVMs.

[Author disclosures](http://www.atsjournals.org/doi/suppl/10.1164/rccm.202005-1937LE/suppl_file/disclosures.pdf) are available with the text of this letter at [www.atsjournals.org.](http://www.atsjournals.org)

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References

- 1. Krowka MJ, Porayko MK, Plevak DJ, Pappas SC, Steers JL, Krom RA, et al. Hepatopulmonary syndrome with progressive hypoxemia as an indication for liver transplantation: case reports and literature review. Mayo Clin Proc 1997;72:44–53.
- 2. Newman B, Feinstein JA, Cohen RA, Feingold B, Kreutzer J, Patel H, et al. Congenital extrahepatic portosystemic shunt associated with heterotaxy and polysplenia. Pediatr Radiol 2010;40:1222–1230.
- 3. Cloutier A, Ash JM, Smallhorn JF, Williams WG, Trusler GA, Rowe RD, et al. Abnormal distribution of pulmonary blood flow after the Glenn shunt or Fontan procedure: risk of development of arteriovenous fistulae. Circulation 1985;72:471–479.
- 4. Vettukattil JJ, Slavik Z, Lamb RK, Monro JL, Keeton BR, Tsang VT, et al. Intrapulmonary arteriovenous shunting may be a universal phenomenon in patients with the superior cavopulmonary anastomosis: a radionuclide study. Heart 2000;83:425–428.
- 5. Srivastava D, Preminger T, Lock JE, Mandell V, Keane JF, Mayer JE Jr, et al. Hepatic venous blood and the development of pulmonary arteriovenous malformations in congenital heart disease. Circulation 1995;92:1217–1222.
- 6. Shah MJ, Rychik J, Fogel MA, Murphy JD, Jacobs ML. Pulmonary AV malformations after superior cavopulmonary connection: resolution after inclusion of hepatic veins in the pulmonary circulation. Ann Thorac Surg 1997;63:960–963.
- 7. Roman BL, Hinck AP. ALK1 signaling in development and disease: new paradigms. Cell Mol Life Sci 2017;74:4539–4560.
- 8. Mahmoud M, Borthwick GM, Hislop AA, Arthur HM. Endoglin and activin receptor-like-kinase 1 are co-expressed in the distal vessels of the lung: implications for two familial vascular dysplasias, HHT and PAH. Lab Invest 2009;89:15–25.
- 9. Tillet E, Ouarné M, Desroches-Castan A, Mallet C, Subileau M, Didier R, et al. A heterodimer formed by bone morphogenetic protein 9 (BMP9) and BMP10 provides most BMP biological activity in plasma. J Biol Chem 2018;293:10963–10974.
- 10. Rochon ER, Krowka MJ, Bartolome S, Heresi GA, Bull T, Roberts K, et al. BMP9/10 in pulmonary vascular complications of liver disease. Am J Respir Crit Care Med 2020;201:1575–1578.
- 11. Treggiari D, Capasso T, Hindes M, Bloch J, Cook S, Trucco S, et al. Investigating the role of BMP9 in development of superior cavopulmonary anastomosis-associated pulmonary arteriovenous malformations. Presented at Vascular Biology and Genetics Workshop VII. October 14, 2017. Pacific Grove, CA. Abstract 14, p. 31.
- 12. Shirali AS, Lluri G, Guihard PJ, Conrad MB, Kim H, Pawlikowska L, et al. Angiopoietin-2 predicts morbidity in adults with Fontan physiology. Sci Rep 2019;9:18328.

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Pulmonary Vascular Pruning on Computed Tomography and Risk of Death in the Framingham Heart Study

To the Editor:

Pulmonary vascular disease, including pulmonary hypertension (PH), is a heterogeneous group of disorders that are associated with a high risk of death. Regardless of etiology, this condition is characterized histologically by the narrowing and loss of the small

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