

PULMONARY HYPERTENSION IN THE MODERN ERA: SCIENCE AND CLINICAL PRACTICE

EDITED BY: Elena Goncharova, Vinicio De Jesus Perez and Harm Jan Bogaard
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PULMONARY HYPERTENSION IN THE MODERN ERA: SCIENCE AND CLINICAL PRACTICE

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Editorial: Pulmonary Hypertension in the Modern Era: Science and Clinical Practice

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Editorial on the Research Topic

Pulmonary Hypertension in the Modern Era: Science and Clinical Practice

Pulmonary hypertension (PH) comprises five groups of progressive cardiopulmonary diseases defined as mean pulmonary arterial (PA) pressure (mPAP) ≥ 20 mmHg (1), leading to right heart failure and premature death. Despite past advances, the pathobiology of different groups of PH is incompletely understood, and diagnostic and treatment options are limited.

Pulmonary arterial hypertension (PAH, Group I PH) is a rare group of conditions that includes familiar and idiopathic forms. Multiple genetic and genomic risk factors have been identified (2), but much remains to discover. Using whole genome sequencing of two unrelated families with PAH, Pienkos et al. identified two novel rare variants in *TNIP2* and *TRAF2*, which could promote an altered immune response, pulmonary vascular remodeling and PH by permitting overactivation of NF- κ B. In parallel studies, Rathinasabapathy et al. created transgenic mice harboring the human *Cav1* mutation. These researchers found that the *Cav1* mutation results in metabolic deficiencies, development of mild PH, reduced spontaneous exercise, and an inflammatory phenotype. This study adds to emerging evidence suggesting a role for insulin deficiency and exercise intolerance, and further supports the role of inflammation in PAH.

Other critical players in PAH are PA endothelial cells (PAECs), and the mechanisms by which PAECs regulate immune function and inflammation in PAH continue to attract attention. In this issue of the Journal, Bhagwani et al. provide a comprehensive review of the role of toll-like receptors, central players in innate immunity, in the immune and non-immune functions of endothelial cells in PH. One important but under-used tool in PAH pre-clinical research is high-resolution imaging, which was successfully used to study pathobiology of other diseases (3). In this special issue, Klouda et al. combined super-resolution microscopy with precision-cut lung slices and clearing technology to develop a novel evolutionary method to study cell behavior *ex vivo*, trace and reconstruct pulmonary vasculature, and address fundamental questions relevant to a wide variety of vascular disorders including PAH.

In addition to idiopathic and familiar forms, group 1 PH includes portopulmonary hypertension (PoPH) that afflicts ~6–8% of patients with portal hypertension and is associated with high morbidity and mortality, particularly in the context of liver transplantation. In this issue, Thomas et al. review the most recent advances in our understanding of the pathogenesis and clinical course of PoPH. The last decade has seen exciting developments in the management of PoPH, with large clinical trials demonstrating the efficacy of several PH-specific therapies in improving

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hemodynamics and clinical outcomes. While there is more work to be done, the authors' state-of-the-art summary leaves us enthusiastic for future progress in the field of PoPH research.

As new oral prostacyclin therapies and prostacyclin agonists have become available for the treatment of PAH, there is an increasing need for methods for transitioning between oral, inhaled and parenteral formulations. Pan et al. describe their experience of successful transitioning between different prostanoid therapies. Particularly because a standardized approach in transitioning safely between different formulations without compromising treatment efficacy is lacking, their case series gives valuable insight into sensible clinical practice and the pharmacokinetics of various prostanoid formulations. In a parallel report, Kopp et al. share their experience with one particular method of prostanoid therapy: intravenous treprostenil administration *via* a Lenus Pro[®] Implantable Pump. While this method has clear advantages, the authors witnessed unexpected acceleration in treprostenil delivery in two patients. The flow rate delivered by the pumps in these 2 patients progressively exceeded the manufacturer admitted margin of error with increases of more than 50% of the expected flow rate at the end of follow up. Spontaneous flow increase from such an implantable pump is a potentially major pitfall, which needs to be identified and actively managed by the responsible clinicians. Prospective studies are needed to assess the long-term safety and efficacy of treprostenil administration by this device.

One of the most common and lethal forms of PH is PH due to left heart disease (PH-LHD, group 2 PH) (4). In their review article, Todd and Lai discuss the challenges and opportunities in developing biomarkers for diagnosing and managing PH-LHD. Presently, the only biomarkers available are B-type natriuretic peptide and N-terminal proBNP; however, data for these biomarkers comes mainly from studies in non-PH-LHD patients, which may not necessarily capture the complex phenotypes of PH-LHD patients. In their paper, the authors discuss exciting studies pointing at endothelin-1, vascular endothelial growth factor-D, and microRNA-206 as new potential circulating biomarkers for patients with PH-LHD and how they could serve as the basis of future risk stratification tools for making management decisions in PH-LHD patients.

A significant challenge for improving the quality of life and survival of PH-LHD patients is the lack of effective treatments. To date, there is no definite evidence that PH-specific therapies can be routinely used to treat patients with PH-LHD. Given this problem, there is an active interest in investigating experimental treatments that could target mechanisms inherent to the pathogenesis of PH-LHD. In this special issue, Mulkareddy and Simon provide a comprehensive review of the evidence

that supports metformin as a prime candidate for clinical trials in PH-LHD. Given the strong association between metabolic dysregulation, insulin resistance, and cardiovascular diseases, investigators have identified the anti-diabetic drug metformin as a potential treatment for PH-LHD. Besides counteracting insulin resistance in type 2 diabetes mellitus, metformin has a range of beneficial effects that could potentially halt the progression of cardiopulmonary remodeling in PH-LHD and improve cardiac function, a major source of morbidity and mortality in these patients. Repurposing metformin for PH-LHD also offers the advantage of improving access to a potentially life-saving drug since the availability of generic formulations could help make this drug affordable and more accessible to patients around the world.

Group 5 PH comprises heterogeneous group of PH secondary to multifactorial mechanisms.

Al-Qadi et al. provide a thorough review of the epidemiology, pathogenesis and clinical approach of group 5 PH, with a specific focus on hematologic disorders, sarcoidosis and chronic renal failure. These forms of PH are caused by a mix of pathophysiologic drivers and can present with precapillary as well as postcapillary disease. The pathogenesis of most forms of group 5 PH includes hypercoagulability, altered vasomotor tone and varied degrees of cardiomyopathy due to a chronic high flow state or intrinsic involvement of the myocardium. In addition, specific to sarcoidosis are advanced interstitial lung disease, granulomatous obliteration of arterioles or venules and extrinsic compression of pulmonary arteries (and venous stenosis) by enlarged lymph nodes or fibrosing mediastinitis. Cardiotoxic and pulmonary toxic effects of medication deserve mention too, particularly the use of tyrosine kinase and alkylating agents in myeloproliferative disorders. Management of group 5 PH primarily involves optimizing treatment of the underlying disease and general measures such as supplemental oxygen, cautiously used diuretics and anticoagulation. PH-targeted therapy is usually reserved for patients with moderate to severe (based on functional class) RHC-confirmed pre-capillary PH with relatively reduced cardiac output.

In conclusion, this special issue highlights continuous interest in pulmonary hypertension research, spanning genetics and genomics, molecular mechanisms of PH pathogenesis, novel imaging techniques, and clinical studies. Together, efforts of many researchers continue to improve our understanding of disease pathobiology and treatment options for patients with PH.

AUTHOR CONTRIBUTIONS

EG, HB, and VJ drafted and edited the manuscript. All authors contributed to the article and approved the submitted version.

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Transitioning Between Prostanoid Therapies in Pulmonary Arterial Hypertension

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Background: New oral prostacyclin therapies and prostacyclin agonists have become available for the treatment of pulmonary arterial hypertension (PAH). However, methods for transitioning between oral, inhaled, and parenteral formulations are not well-established, except in the form of case reports and case series. Collectively, these emphasize the lack of a standardized process and approach in transitioning patients from PAH prostanoid therapies. In this case series, we report our experience at an accredited Pulmonary Hypertension center in transitioning between various oral, inhaled, and parenteral prostanoids to offer additional guidance on safe transitions in therapy.

Methods: All cases of prostanoid transitions at an accredited Pulmonary Hypertension center from March 2018 to September 2019 were included in this report. The transition approach for each case was developed through a review of the literature, extrapolation of available pharmacokinetic data, and collaboration between pharmacists and clinicians.

Results: This case series describes the transition of 3 patients from selexipag to parenteral treprostinil; 1 patient transitioning from parenteral treprostinil to selexipag; 1 patient transitioning from oral treprostinil to parenteral treprostinil; and 1 patient transitioning from inhaled treprostinil to selexipag. Four of the 6 patients presented here were transitioned to an alternate prostanoid on account of clinical worsening, while the remaining 2 patients transitioned due to intolerance of parenteral therapy and poor medication adherence. This case series includes patients with various etiologies of PAH including idiopathic PAH, methamphetamine-associated PAH, and scleroderma-associated PAH. All patients successfully completed each transition without serious adverse events.

Conclusions: With the increasing utilization and availability of prostanoids, there is a critical need for a standardized approach in transitioning safely between different formulations without compromising treatment efficacy. In this case series, we present our clinical experiences, guided by available pharmacokinetic data, in transitioning between various prostanoid formulations.

Keywords: pulmonary hypertension, therapeutics, right heart failure, pharmacology, prostaglandin, prostacyclin

BACKGROUND

In recent years, new oral prostanoid therapies such as selexipag and oral treprostinil have become globally available for the treatment of pulmonary arterial hypertension (PAH). These agents are becoming increasingly utilized due to their efficacy, oral route of administration, and potentially favorable side effect profile based on data from the FREEDOM trials with oral treprostinil (1–4) and the GRIPHON trial with selexipag (5). Established guidance on transitions between various formulations within this drug class have not been standardized. A number of case reports and case series have been published describing such transitions but lack consistent processes (6–13).

The most common method of determining dose equivalency between parenteral treprostinil and oral treprostinil is derived from a small phase 2 trial consisting of 33 patients. This dose conversion strategy can be found in the package insert for oral treprostinil and is summarized by the following equation (14):

Oral treprostinil total dose/day (mg) = parenteral treprostinil dose (ng/kg/min) \times weight (kg) \times 0.0072

During the transition period, the package insert recommends reducing parenteral treprostinil by up to 30 ng/kg/min per day while also increasing the dose of oral treprostinil by up to 6 mg per day as tolerated (14). However, despite sharing similar dosing strategies based on these recommendations, there are still numerous variations between different institutional practices. Most notable are the durations over which the transitions are implemented (6–10).

There is no standardized dose conversion strategy for transitioning from oral treprostinil to inhaled treprostinil, and no recommendations are provided in the package insert (14). There is published data for patients on doses of parenteral treprostinil ranging from 22.5 to 111 ng/kg/min and inhaled treprostinil ranging from 9 to 54 mcg four times daily, who have subsequently been transitioned to oral treprostinil 1 mg twice daily to 14 mg three times daily (7, 9, 10). Given the linear pharmacokinetic relationship between treprostinil plasma concentrations and doses up to 125 ng/kg/min, relative dose equivalencies between different formulations can be estimated (15). Parenteral treprostinil 10 ng/kg/min results in plasma concentrations of \sim 1–2 ng/mL, which is roughly equivalent to inhaled treprostinil 54 mcg four times daily (15–17). Assuming inhaled treprostinil 54 mcg four times daily is approximately equivalent to parenteral treprostinil 5–10 ng/kg/min, the corresponding dose of oral treprostinil (for an average 70 kg patient) would be \sim 1–2 mg three times daily. Other studies have described successful transitions between inhaled and parenteral treprostinil using parenteral doses of up to 25 ng/kg/min (18, 19).

For transitions between parenteral or inhaled prostanoids and selexipag, there is considerably less guidance available. Selexipag is an oral selective IP receptor agonist, and the pharmacokinetic relationship between dose and equivalent plasma concentrations are unknown. In the TRANSIT-1 study, 34 PAH patients on inhaled treprostinil were transitioned to selexipag over 8 weeks with a 94.1% success rate. The mean dose of inhaled treprostinil at baseline was 59.3 mcg four times daily (13). In a case series of 4 patients on parenteral treprostinil, the highest dose

upon initiation of the transition was 46.5 ng/kg/min, which was successfully transitioned to selexipag 1,600 mcg twice daily (11). In another case series of 5 patients, parenteral treprostinil doses up to 96 ng/kg/min were successfully transitioned to selexipag 1,600 mcg twice daily (20). It is important to note the lack of information beyond 6 months following the transition, particularly in patients previously on high treprostinil doses.

In this case series, we report our experience in transitioning between oral, inhaled, and parenteral prostaglandin products at an accredited Pulmonary Hypertension (PH) center (Table 1).

METHODS

Approval of this study was obtained through the University of Utah Institutional Review Board. All cases that underwent transition between different prostanoid formulations at the University of Utah Pulmonary Hypertension Comprehensive Care Center from March 2018 to September 2019 were included in this report. No cases were omitted, and all available relevant data were included in this report. Written informed consent was obtained from the participants for the publication of this case report. The consecutive cases were collected prospectively. The different methodologies for transition were created after an extensive literature review and with collaboration between pharmacists and clinicians at University of Utah Health.

CASE PRESENTATION

Case #1: Selexipag to Parenteral Treprostinil

A 66-year-old female with World Health Organization (WHO) group 1, functional class (FC) III scleroderma-associated PAH underwent a planned admission to transition from selexipag to parenteral treprostinil due to declining clinical status. Prior to starting any PAH therapies, her right heart catheterization (RHC) showed a right atrial pressure (RAP) of 25 mmHg, pulmonary artery pressure (PAP) of 61/26 mmHg with a mean PAP of 43 mmHg, pulmonary capillary wedge pressure (PCWP) of 9 mmHg, pulmonary vascular resistance (PVR) of 6.7 Wood Units (WU), and cardiac index (CI) (Fick) of 2.3 L/min/m². PAH medications prior to the transition were: ambrisentan 10 mg daily, sildenafil 20 mg three times daily, and selexipag 600 mcg twice daily. The patient had received selexipag for \sim 7 months. Her other medications included apixaban 5 mg twice daily, bumetanide 2 mg twice daily, and metolazone 2.5 mg twice weekly. Prior to the transition, while on PAH therapies, the patient had a 6 minute walk distance (6MWD) of 213 m; brain natriuretic peptide (BNP) of 288 pg/mL; and a transthoracic echocardiogram (TTE) demonstrating an estimated right ventricular systolic pressure (RVSP) of 61 mmHg, a moderate pericardial effusion, and normal right ventricle (RV) size and function. On admission, her total body weight was 101.4 kg (BMI 38.4 kg/m²). She was hemodynamically stable with an oxygen saturation of 99% on her baseline oxygen requirement of 1L. Selexipag total daily dose was decreased by 400 mcg per day (split between the morning and evening

TABLE 1 | Baseline characteristics.

	Case #1	Case #2	Case #3	Case #4	Case #5	Case #6
Age (yr)	66	41	40	32	20	56
Type of PAH	Scleroderma-associated	Idiopathic	Methamphetamine-associated	Methamphetamine-associated	Idiopathic	Methamphetamine-associated
Transitioning agents	Selexipag to SQ Treprostinil	Selexipag to SQ Treprostinil	Selexipag to SQ Treprostinil	SQ Treprostinil to Selexipag	Oral Treprostinil to SQ Treprostinil	Inhaled Treprostinil to Selexipag
6MWD (m)	213	213	383	671	520	312
BNP (pg/mL)	288	579	NT-proBNP 599 pg/mL	21	43	189
Hemodynamics	RAP 25 mmHg PAP 61/26 mmHg mPAP 43 mmHg PCWP 9 mmHg PVR 6.7 WU CI 2.3 L/min/m ²	RAP 19 mmHg PAP 80/40 mmHg mPAP 54 mmHg PCWP 18 mmHg PVR 8.6 WU CI 2.56 L/min/m ²	RAP 5 mmHg PAP 87/32 mmHg mPAP 57 mmHg PCWP 12 mmHg PVR 9.3 WU CI 2.46 L/min/m ²	RAP 3 mmHg PAP 120/57 mmHg mPAP 78 mmHg PCWP 17 mmHg PVR 9.2 WU CI 3.7 L/min/m ²	RAP 10 mmHg PAP 100/55 mmHg mPAP 79 mmHg PCWP 21 mmHg PVR 13.68 WU CI 2.35 L/min/m ²	RAP 6 mmHg PAP 63/25 mmHg mPAP 44 mmHg PCWP 7 mmHg PVR 11.7 WU CI 1.6 L/min/m ²

TABLE 2 | Selexipag to parenteral treprostinil transitions (Cases #1-#3).

Drug Titration	Case 1	Case 2	Case 3
Selexipag dose prior to transition	600 mcg twice daily	1,600 mcg twice daily	1,400 mcg twice daily
Selexipag dose at time of treprostinil initiation	400 mcg	1,200 mcg	1,200 mcg
Treprostinil initiation dose	5 ng/kg/min	5 ng/kg/min	5 ng/kg/min
Selexipag dose down-titration	Total daily dose decreased by 400 mcg every 24 h	Total daily dose decreased by 800 mcg every 24 h until 400 mcg dose reached, then decreased by 400 mcg every 24 h	Decreased by 200 mcg per dose every 12 h
Treprostinil dose up-titration	Increased by 5 ng/kg/min every 24 h until selexipag off, then increased by 1 ng/kg/min every 24 h as tolerated	Increased by 3–5 ng/kg/min every 24 h until selexipag off, then increased by 1 ng/kg/min every 24 h as tolerated	Increased by 5 ng/kg/min every 24 h until selexipag off, then increased by 1 ng/kg/min every 24 h as tolerated
Treprostinil dosing weight used	54.2 kg (IBW)	70 kg (IBW)	50.1 kg (IBW)
Duration of transition	3 days	5 days	3 days
Treprostinil dose reached at hospital discharge	13 ng/kg/min	20 ng/kg/min	20 ng/kg/min

doses) starting on the day of hospital admission. Intravenous treprostinil was initiated at 5 ng/kg/min using an ideal body weight of 54.2 kg on day 1 of admission and increased by increments of 5 ng/kg/min every 24 hours to 10 ng/kg/min by the time of selexipag discontinuation. The transition was completed over 3 days (Table 2). The patient tolerated the transition well with no reports of headaches, flushing, flu-like symptoms, or symptomatic hypotension. Throughout the transition, she required 1 to 2 L/min of oxygen to maintain oxygen saturations >90% at rest. Her hospital stay was complicated by fluid overload and acute kidney injury likely secondary to cardiorenal syndrome, despite an adequate cardiac output shown on RHC. Following the transition, her RHC hemodynamics showed a RAP of 23 mmHg, PAP of 67/29 mmHg with a mean PAP of 44 mmHg, PCWP of 27 mmHg (increased PCWP secondary to right ventricular compression of the left ventricle), PVR of 2.2 WU, and CI (Fick) of 3.8 L/min/m². Prior to hospital discharge, the patient was transitioned from IV to SQ treprostinil and was instructed to continue up-titrating by 1 ng/kg/min every 2–4 days until

goal of 40 ng/kg/min. At a 7-month follow-up, she was on SQ treprostinil at a rate of 43 ng/kg/min. Her BNP was stable at 261 pg/mL, and her TTE was unchanged. However, her 6MWD had decreased to 168 m, and she required 3 L/min of oxygen to maintain an oxygen saturation of >88%. Repeat RHC showed a RAP of 19 mmHg, PAP of 61/25 mmHg with a mean PAP of 37 mmHg, PCWP of 22 mmHg, PVR of 1.7 WU, and CI (Fick) of 4.14 L/min/m².

Case #2: Selexipag to Parenteral Treprostinil

A 41-year-old female with WHO group 1, FC IV idiopathic PAH underwent a planned admission to transition from selexipag to parenteral treprostinil. She had worsening functional class on ambrisentan 10 mg daily, sildenafil 20 mg three times daily, and selexipag 1,600 mcg twice daily. Her concomitant medications included torsemide 40 mg daily, spironolactone 25 mg daily, digoxin 250 mcg daily, and apixaban 5 mg twice daily. Prior to the transition, while on PAH therapies, the patient had a

6MWD of 213 m, BNP of 579 pg/mL, and a TTE demonstrating an estimated RVSP of 69 mmHg and a severely enlarged RV with mildly reduced RV systolic function. Her RHC on admission showed a RAP of 19 mmHg, PAP of 80/40 mmHg with a mean PAP of 54 mmHg, PCWP of 18 mmHg, PVR of 8.6 WU, and CI (Fick) of 2.56 L/min/m². On admission, her total body weight was 107 kg (BMI 44.6 kg/m²). She was hemodynamically stable with an oxygen saturation of 92% on her baseline oxygen requirement of 4L. Selexipag was reduced to 1,200 mcg twice daily (75% of her home dose) on the day of admission and subsequently decreased by 800 mcg per day (split between the morning and evening doses). Once 400 mcg was reached, the dose was further decreased to 200 mcg after 24 h before stopping. Intravenous treprostinil was initiated on hospital day 1 at 5 ng/kg/min using an ideal body weight of 70 kg and was increased by 3–5 ng/kg/min every 24 h to 20 ng/kg/min by the time of selexipag discontinuation. The transition took place over the course of 5 days (Table 2). The patient tolerated the transition with no complaints. The patient did develop a rash during the admission, which was thought to be contact dermatitis secondary to an adhesive. Throughout the transition, her oxygen saturations remained >88% on 4 to 6 L/min NC, and her renal function remained unchanged. Prior to discharge, the patient was transitioned from IV to SQ treprostinil. She was continued on a gradual up-titration as an outpatient with a goal of 40 ng/kg/min and had reached 30 ng/kg/min by the time of her 1 month follow up.

Case #3: Selexipag to Parenteral Treprostinil

A 40-year-old female with WHO group 1, FC III methamphetamine-associated PAH underwent a planned admission for transition from oral selexipag to parenteral treprostinil. She initially presented to our PH clinic with progression of disease and worsening functional status despite treatment with tadalafil 40 mg daily, macitentan 10 mg daily, and selexipag 1,400 mcg twice daily for nearly 3 years. Her other medications included furosemide 40 mg in the morning and 20 mg in the afternoon and spironolactone 25 mg daily. Prior to her transition, the patient had a 6MWD of 382 m, NT-proBNP of 599 pg/mL, and a TTE revealing an estimated RVSP of 91 mmHg and a severely enlarged RV with mildly reduced RV systolic function. Her RHC showed a RAP of 5 mmHg, PAP of 87/32 mmHg with a mean PAP of 57 mmHg, PCWP of 12 mmHg, PVR of 9.3 WU, and CI (Fick) of 2.46 L/min/m². On admission, her total body weight was 94.5 kg (BMI 38.1 kg/m²). She was hemodynamically stable with an oxygen saturation of 90% on her baseline oxygen requirement of 4L. Selexipag was decreased by 200 mcg every 12 h from her home dose of 1,400 mcg twice daily. IV treprostinil was initiated at 5 ng/kg/min using an ideal body weight of 50.1 kg at the time of the first selexipag dose decrease and subsequently uptitrated by 5 ng/kg/min every 24 h to 15 ng/kg/min by the time of selexipag discontinuation. The transition took place over a period of 3 days (Table 2). After selexipag discontinuation,

treprostinil was increased to 20 ng/kg/min prior to hospital discharge. The patient tolerated the transition well with some mild nausea but had no other complaints. Her oxygen saturations remained >88% on 4–6 L/min of oxygen at night and on 1–3 L/min during the day throughout the transition. She was discharged on her home oxygen requirement of 4 L/min at night. On the day of discharge, the patient was transitioned from IV to SQ treprostinil with the plan to increase treprostinil by 3 ng/kg/min every 3–6 days to a goal of 35 ng/kg/min. At the time of publication, the patient is doing well but has not yet had follow up studies performed after her transition.

Case #4: Parenteral Treprostinil to Selexipag

A 32-year-old male with WHO group 1, FC II methamphetamine-associated PAH was admitted for transition from subcutaneous treprostinil to oral selexipag. The reason for transition was patient preference as his quality of life had significantly declined due to being limited by daily pump management. This transition was strongly discouraged by his PH care team, but after several conversations regarding his perspective and the risks associated, he chose to transition with close follow up. The patient was diagnosed with PAH 10 months prior to admission and was originally started on epoprostenol at the time of diagnosis but was quickly transitioned to subcutaneous treprostinil. His dose of treprostinil at the time of this transition was 32 ng/kg/min using a dosing weight of 70.6 kg. He was also on tadalafil 40 mg daily and macitentan 10 mg daily. Prior to the transition, while on PAH therapies, the patient had a 6MWD of 671 m, BNP of 21 pg/mL, and a TTE with an RVSP of 71.6 mmHg with normal RV systolic function and mild enlargement. His most recent RHC measurements immediately prior to the transition demonstrated a RAP of 3 mmHg, PAP of 120/57 mmHg with a mean PAP of 78 mmHg, PCWP of 17 mmHg, PVR of 9.2 WU, and CI (Fick) of 3.7 L/min/m². On admission, his total body weight was 65.5 kg (BMI 23.3 kg/m²). The patient was hemodynamically stable with an oxygen saturation of 96% on room air while on his home treprostinil, tadalafil, and macitentan. The patient underwent a simultaneous down-titration of treprostinil by 4 ng/kg/min and up-titration of selexipag by 200 mcg every 12 h. The transition took place over a period of 5 hospital days (Table 3). The patient had no complaints of side effects during the transition. The patient's renal function remained stable during the transition, and his oxygen saturation remained >88% on his baseline requirement of 3L nocturnal oxygen and room air throughout the day. The patient successfully reached 1,600 mcg BID at the end of the transition and was discharged home on tadalafil, macitentan, and selexipag. At a 3-month follow-up, the patient had a 6MWD of 649 m, a BNP of 29 pg/mL, and a TTE that demonstrated an RVSP of 94.4 mmHg and severe RV enlargement and normal RV systolic function. He had complaints of flushing and jaw pain on selexipag 1,600 mcg BID, and thus his dose was reduced to 1,400 mcg daily in the morning and 1,600 mcg daily at bedtime.

TABLE 3 | Parenteral treprostinil to selexipag transition (Case #4).

	Treprostinil dose titration	Selexipag dose titration
Day 1 AM	32 ng/kg/min	0 mcg
Day 1 PM	28 ng/kg/min	200 mcg
Day 2 AM	24 ng/kg/min	400 mcg
Day 2 PM	20 ng/kg/min	600 mcg
Day 3 AM	16 ng/kg/min	800 mcg
Day 3 PM	12 ng/kg/min	1,000 mcg
Day 4 AM	8 ng/kg/min	1,200 mcg
Day 4 PM	4 ng/kg/min	1,400 mcg
Day 5 AM	0 ng/kg/min	1,600 mcg

Case #5: Oral Treprostinil to Parenteral Treprostinil

A 20-year-old male with WHO group 1, FC III idiopathic PAH was admitted for transition from oral treprostinil to subcutaneous treprostinil. The patient had a diagnosis of PAH since the age of 4 and had been on PAH-specific therapies prior to presentation to our center. The reason for transition was declining functional status on tadalafil 40 mg daily, macitentan 10 mg daily, and treprostinil diolamine 4 mg three times daily. Other medications included aspirin 81 mg daily and digoxin 125 mcg daily. His 6MWD at baseline on PAH therapies was 520 m, with a desaturation to 69%, requiring 25L via nasal cannula to bring his oxygen saturation to 82% at the end of 6 minutes. His initial BNP was 43 pg/mL. His TTE showed an RVSP of 97.9 mmHg, moderate RV enlargement, severe RV wall thickness, and normal RV systolic function. His RHC immediately prior to his transition to IV treprostinil showed a RAP of 10 mmHg, PAP of 100/55 mmHg with a mean PAP of 79 mmHg, PCWP of 21 mmHg, PVR of 13.68 WU, and CI (Fick) of 2.35 L/min/m². On admission, the patient's total body weight was 69.9 kg (BMI 23.4 kg/m²). The patient was hemodynamically stable on his home treprostinil and had an oxygen saturation of 90% on room air. The patient underwent a simultaneous down-titration of oral treprostinil by 1 mg three times daily every 24 h while initiating and increasing IV treprostinil by 6 ng/kg/min every 24 h. The transition took place over a period of 4 hospital days (Table 4). The patient had no complaints of side effects during the transition, and the patient's renal function remained stable during the transition. His oxygen saturation at rest remained >90% throughout the entire transition without supplemental oxygen and was 94% on room air at the time of discharge. The patient successfully reached 24 ng/kg/min at the end of the transition and was converted to subcutaneous treprostinil and discharged home without any problems. At 2-month follow-up, the patient was WHO-FC II and had reached a dose of 40 ng/kg/min. His 6MWD had improved to 613 m, but he still required 25L of oxygen to achieve an oxygen saturation of 83%. His treprostinil was gradually increased to 61 ng/kg/min. At his 6-month follow-up, there was no new RHC data, his BNP was stable at 52 pg/mL and a TTE demonstrated normal RV size, severe RV wall thickness, and normal RV function.

TABLE 4 | Oral Treprostinil to Parenteral Treprostinil (Case #5).

	Oral treprostinil down-titration	IV treprostinil dose titration
Day 1 AM	4 mg	
Day 1 Mid-day	4 mg	6 ng/kg/min
Day 1 PM	3 mg	6 ng/kg/min
Day 2 AM	3 mg	6 ng/kg/min
Day 2 Mid-day	3 mg	12 ng/kg/min
Day 2 PM	2 mg	12 ng/kg/min
Day 3 AM	2 mg	12 ng/kg/min
Day 3 Mid-day	2 mg	18 ng/kg/min
Day 3 PM	1 mg	18 ng/kg/min
Day 4 AM	1 mg	18 ng/kg/min
Day 4 Mid-day	1 mg	24 ng/kg/min
Day 4 PM	OFF	24 ng/kg/min
	Discharged	

Case #6: Inhaled Treprostinil to Selexipag

A 56-year-old male with WHO group 1 methamphetamine-associated PAH, diagnosed 3.5 years previously, presented to our PH clinic after being lost to follow-up for 15 months. At the time of reestablishing care, the patient was WHO-FC III and was directly admitted to the hospital for intravenous diuresis for right heart failure and re-initiation and optimization of PAH-specific therapies. The patient was experiencing intolerable cough and poor adherence to inhaled treprostinil, and the decision was made to transition him to selexipag. The patient was also non-adherent to a regimen of tadalafil 40 mg daily and macitentan 10 mg daily. His other medications included torsemide 80 mg daily, metolazone 5 mg as needed, potassium chloride 20 mEq three times daily, spironolactone 100 mg daily, and aspirin 81 mg daily. Prior to admission, the patient had a 6MWD of 312 m and resting oxygen saturation of 90% on room air. His initial BNP was 189 pg/mL. His TTE showed an RVSP of 65.9 mmHg, severe RV enlargement, and severely reduced RV systolic function. RHC showed a RAP of 6 mmHg, PAP of 63/25 mmHg with a mean PAP of 44 mmHg, PCWP of 7 mmHg, PVR of 11.7 WU, and CI (Fick) of 1.6 L/min/m². On admission to the hospital, the patient was restarted on macitentan 10 mg daily, and treprostinil was changed to 6 puffs four times daily for medication optimization in anticipation of transitioning him from inhaled treprostinil to selexipag. He was WHO-FC II upon discharge from the hospital, but due to a delay in access had an interruption in macitentan for 6 days after discharge. The decision was then made to begin the transition from inhaled treprostinil to selexipag 2 weeks later as an outpatient. At the time of transition, the patient was using inhaled treprostinil 6 puffs three times daily. The plan was to initiate selexipag 200 mcg twice daily for 1 week while maintaining inhaled treprostinil at 6 puffs three times daily, then to continue the transition as outlined in Table 5. However, during the second week of the transition, the patient complained of increased cough related to inhaled treprostinil and wanted to discontinue use. The patient had also not increased selexipag as planned. The transition plan was modified as shown in Table 5.

TABLE 5 | Inhaled treprostinil to selexipag (Case #6).

Planned transition			Actual transition		
	Inhaled treprostinil down-titration	Selexipag up-titration		Inhaled treprostinil down-titration	Selexipag up-titration
Week 1	6 puffs three times daily	200 mcg twice daily	Week 1	6 puffs three times daily	200 mcg twice daily
Week 2	3 puffs three times daily	400 mcg twice daily	Week 2	3 puffs three times daily	400 mcg in the morning and 200 mcg in the evening
Week 3	OFF	600 mcg twice daily Increase every 1–2 weeks as tolerated to max of 1,600 mcg twice daily	Week 3	OFF	400 mcg twice daily Increase every 1–2 weeks as tolerated to max of 1,600 mcg twice daily

The patient reached a maintenance dose of 1,600 mcg twice daily approximately 6 weeks after starting selexipag. At a 6-month follow-up, he remained on selexipag 1,600 mcg twice daily. His 6MWD improved to 431 m, and his BNP was 164 pg/mL. His TTE at 8 months showed a reduction in RVSP to 53.4 mmHg, severe RV enlargement, and moderately to severely reduce RV function.

DISCUSSION

These 6 cases provide real world experience of transitioning patients between oral, inhaled, and parenteral prostanoids at a specialized PAH center over an 18-month period. This series illustrates one of the many challenges in contemporary PAH management where there is a lack of consensus regarding transitions between different agents. As prescriptions for oral prostanoid products continue to rise and the number of specialized PAH centers and providers continue to expand, it is increasingly essential to share experiences and institutional practice patterns. The purpose of this case series was to provide additional insight into transition strategies and serve as a resource for other practitioners faced with the challenge of transitioning patients between alternative prostanoid formulations.

We created individualized approaches for transitioning between prostanoid formulations based on reviews of the available literature and extensive discussion amongst our PAH providers. The absence of standardized protocols to facilitate prostanoid transitions provides a challenge to those caring for PAH patients. The risks of providing a suboptimal dose with prostanoid transitions includes acute right ventricular failure and death. The risk of transitioning patients to significantly higher doses too quickly include hypotension resulting in organ failures, as well as significant side effects. This in turn may go on to limit the amount prescribed, thus leading to inadequate treatment of patients with this severe, progressive disease.

When faced with the clinical scenario of transitioning patients between various prostanoid agents, we found the limited number of published reports describing real-world experiences to be challenging. The goal of this paper is to serve as a practical guide to our approach and details the challenges associated

with these transitions. This paper is distinct from prior reports because it is comprised of different strategies and methods of delivering prostanoids, namely oral, inhaled, intravenous, and subcutaneous. Prior reports largely focused on one type of transition.

Within this case series, the predominant reason for transitioning between prostanoid formulations was clinical deterioration. Less commonly, patients were transitioned due to intolerance or poor adherence to the previous delivery system. In addition to studying how best to safely implement transitions, it would also be worthwhile to further examine reasons to switch between different prostanoid analogs from both the patients' and providers' perspectives. In the era of shared-decision making, it is important to ensure patients have been thoroughly educated on initial prostanoid therapy options, and in turn, that transition between agents is feasible, as demonstrated in this case series (21).

More prostanoid analogs are currently under clinical investigation, including a dry powder inhaled form of treprostinil (22) and ralinepag (23), a novel prostacyclin IP receptor agonist. Unless efforts are made to study the optimum manner in which PAH providers and their patients can transition between agents, this area will become increasingly complex. It will be important for drug developers and those working in the field of PAH to study and develop clinical standards in transitioning between agents.

LIMITATIONS

The cases presented here are small in number and come from a single center. Other reports have described more standardized protocols, with larger sample sizes and with some homogeneity between cases (7, 13).

In clinical practice there are well-known difficulties in categorizing patients with pulmonary hypertension. For example, in this series, several patients displayed a PCWP >15 mmHg at various time points during their clinical care. This reflects the difficulty with interpreting hemodynamics in PAH patients and emphasizes one of the challenges in PH management. All patients met WHO group I PAH criteria at the time of therapy initiation. However, over time, several patients developed

progressive heart failure with elevated PCWP likely due to left ventricular compression from an oversized right ventricle. As is routine in our and other PH programs, these cases are reviewed thoroughly and repeatedly in order to have a confirmatory diagnosis of PAH.

During the period under study, no patients were transitioned from or to epoprostenol. Therefore, such transitions are not presented in this paper. Although our transitions were well-tolerated clinically, it remains difficult to compare approaches between different institutions.

CONCLUSIONS

With the increased use of oral prostanoids, there is a critical need to provide evidence- and pharmacokinetic-based guidance on how to transition between oral and parenteral prostanoids. In this case series, we present our approach for this practice in the hope to generate discussion and to determine best practice in this area.

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ETHICS STATEMENT

Written informed consent was obtained from the participants for the publication of this case report.

AUTHOR CONTRIBUTIONS

All authors contributed to reviewing the source data and crafting the manuscript.

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From 2D to 3D: Promising Advances in Imaging Lung Structure

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The delicate structure of murine lungs poses many challenges for acquiring high-quality images that truly represent the living lung. Here, we describe several optimized procedures for obtaining and imaging murine lung tissue. Compared to traditional paraffin cross-section and optimal cutting temperature (OCT), agarose-inflated vibratome sections (aka precision-cut lung slices), combines comparable structural preservation with experimental flexibility. In particular, we discuss an optimized procedure to precision-cut lung slices that can be used to visualize three-dimensional cell-cell interactions beyond the limitations of two-dimensional imaging. Super-resolution microscopy can then be used to reveal the fine structure of lung tissue's cellular bodies and processes that regular confocal cannot. Lastly, we evaluate the entire lung vasculature with clearing technology that allows imaging of the entire volume of the lung without sectioning. In this manuscript, we combine the above procedures to create a novel and evolutionary method to study cell behavior *ex vivo*, trace and reconstruct pulmonary vasculature, address fundamental questions relevant to a wide variety of vascular disorders, and perceive implications to better imaging clinical tissue.

Keywords: lung structure, vibratome, precision cut lung slices, optical clearing, confocal, STED

INTRODUCTION

High resolution imaging of intact tissue is vital to understanding the anatomy and physiology underlying human disease. Especially in the lung, where cells and tissues interact with the different environmental and internal stimuli, robust visualization of tissue throughout the intact lung helps identify the patterns and characteristics of diseases from asthma to fibrosis to COPD (1).

Detailed imaging of murine lung tissue that preserves the structure and cellular components helps advance the knowledge and understanding of diseases so therapeutic approaches can be applied to human subjects. The ability to label protein, RNA, and other biological compounds with a high signal-to-noise ratio while maintaining physiological structure broadens the understanding of pathology and can demonstrate the spatial approximation of different biological compounds and cell types *in situ*.

Despite sophisticated advancements in imaging technology, acquiring high-quality, clinically relevant data remains challenging. The inherent difficulties in lung tissue preparation are the technical bottleneck. Sub-optimally prepared lung tissue can lead to inaccurate alveolar architecture, light scattering and difficulties with fluorescent staining, which may potentially produce an inaccurate model.

The natural respiratory cycle is driven by negative intrathoracic pressure generated by the respiratory muscles (2). Without this pressure gradient, as when a lung is harvested, the delicate pulmonary tissue collapses and no longer resembles its physiological structure. Preparing lung tissue for imaging requires inflating the collapsed airway and perfusing the vasculature, which changes the dynamic lung into a static fixture, maintaining structural integrity and molecular components. Given the different configuration and composition in alveolar and endothelial tissue, along with the rate at which the airways and vasculature taper in the lung, distinct techniques must be used to prepare tissue as realistically as possible.

Numerous different strategies and techniques exist for preparing animal and human tissue for visualization (1, 3, 4). The standard of practice for pathologists and histologists is preparing tissue in thin slices for microscopic visualization. However, advances in biomedical photonics has allowed advanced viewing of lung tissue, improving structure, resolution and clarity of samples. Precision-cut lung slices (PCLS) provides the advantage of maintaining the three-dimensional architecture of the lung compared to thin slices, allowing for dynamic views similar to physiological conditions. Advances in microscopy, including stimulated emission depletion microscopy (STED), bypass the diffraction limit of light microscopy providing super resolution images. Clearing optical techniques, such as solvent-based clearing protocols, has significantly improved the viewing of these biological specimens by decreasing the amount of light scattering (5). Current techniques focus on optimizing tissue clarity, retaining anatomical structure, and maintaining cellular molecules for fluorescence staining.

The primary purpose of this manuscript is to describe a novel, evolutionary method to prepare thin-sectioned, thick-sectioned, and whole-organ murine lung tissue that combines OCT tissue clearing, PCLS and tissue clearing into one protocol. The optimized sample provides high-quality images of preserved structures that can be fluorescence-stained. A secondary purpose of this manuscript is to review commonly used tissue preparation methods, discussing the pros and cons of each strategy.

MATERIALS AND METHODS

Please refer to **Table 1** for instruments and equipment, chemicals and reagents and solutions used during methods.

Animals

Animals used were 10–12 weeks, 22–25 g, male C57BL6 or NG2-Cre-ER/ROSA26RtdTomato (NG2-tdT) mice, as previously published (6). The NG2-tdT mouse line exhibits selective pericyte labeling by tdTomato when B6.Cg-Tg (Cspg4-Cre/Esr1*) B6Kik/J (NG2-Cre-ER) mice (<https://www.jax.org/strain/008538>) are crossed with with tdTomato Ai14 (<https://www.jax.org/strain/007908>). NG2-tdT mice were treated with 2 mg tamoxifen dissolved in corn oil (20 mg/ml) for two consecutive injections. After tamoxifen treatments, mice rested for 7 days before tissue harvest.

TABLE 1 | Equipment, Chemicals and reagents, Instruments and solutions used in the materials and methods.

Equipment

- Cryostat slicer (HM525NX, ThermoFisher)
- Vibration slicer (Leica VT1000s, Nussloch, Germany)
- Confocal microscope (Zeiss 880 with Airyscan)
- Light Sheet microscope (LaVision)
- Facility Line Confocal/STED microscopy (Abberior Instruments America LLC)

Chemicals and reagents

- UltraPure Low Melting Point Agarose (2%, Invitrogen, 16520050)
- Mouse-anti-mouse/human SMA-Cy3 (1:300, Sigma, C6198-.2ML)
- Rat-anti-mouse CD31 (1:100; BD-Pharmingen, 553370 or 550274)
- Rabbit-anti-mouse-RFP (1:100, Rockland, 600-401-379)
- Goat anti-rabbit STAR RED (1:100, Abberior Instruments GmbH, STRED)

Instruments and solutions

- Flushing buffer (1X PBS + 0.1% heparin)
- PLP perfusion buffer (0.075 M lysine, 0.37 M sodium phosphate pH 7.2, 2% formaldehyde, and 0.01 M NaIO₄, a detailed receipt can be found at Cold Spring Harbor Protocols)
- Inflation solution (½ volume of 100% OCT mixed with ½ volume of 30% sucrose)
- Expansion solution (2% agarose in 1XPBS)
- Blocking buffer (5% serum (goat, determined by species of secondary antibodies), 0.3% Triton X100 in 1XPBS)
- Antibody dilution buffer (1% BSA, 0.3% Triton X-100 in 1XPBS)
- MPBS (Percentage of Methanol in 1XPBS)
- BABB solution (1 volume of Benzyl Alcohol mix with 2 volume of Benzyl Benzoate)

Harvest, Cannulation, and Perfusion of Mouse Lung

Anesthetize the animal with 3% isoflurane to ensure the lack of pain reflex, then sacrifice by neck isolation. Open the chest and create a small nick through the suprarenal abdominal aorta. Insert a 25–30 g butterfly needle* (**Figures 1A,A'**) connected to a 30 cc syringe into the right ventricle. Flush with a 20 ml flushing buffer slowly but steadily until the fluid extravasating from the abdominal aorta is clear and colorless. *Note: use of the butterfly needle prevents fluid leakage due to multiple cannulations.

(1) OCT Tissue Preparation

Switch syringe at infusion port of butterfly needle and flush with Periodate-Lysine-Paraformaldehyde (PLP) buffer. Cannulate trachea with 22 g blunt-ended needle and inflate with 2.5–4 ml OCT inflation solution. Immediately following inflation, separate and put each lobe in the tissue mold completely covered with OCT. Place the mold on a metal block and submerge in liquid nitrogen. After the OCT is completely frozen, store at -80° . Proceed with cryosectioning, immunolabeling, counterstaining, and imaging (each step is summarized in **Figure 2**).

(2) Vibratome

Cannulate trachea with 22 g blunt-ended needle and inflate with 2.5–4 ml 2% agarose solution. Incubate the dissected lungs in 4% paraformaldehyde solution overnight at 4° on an agitator.

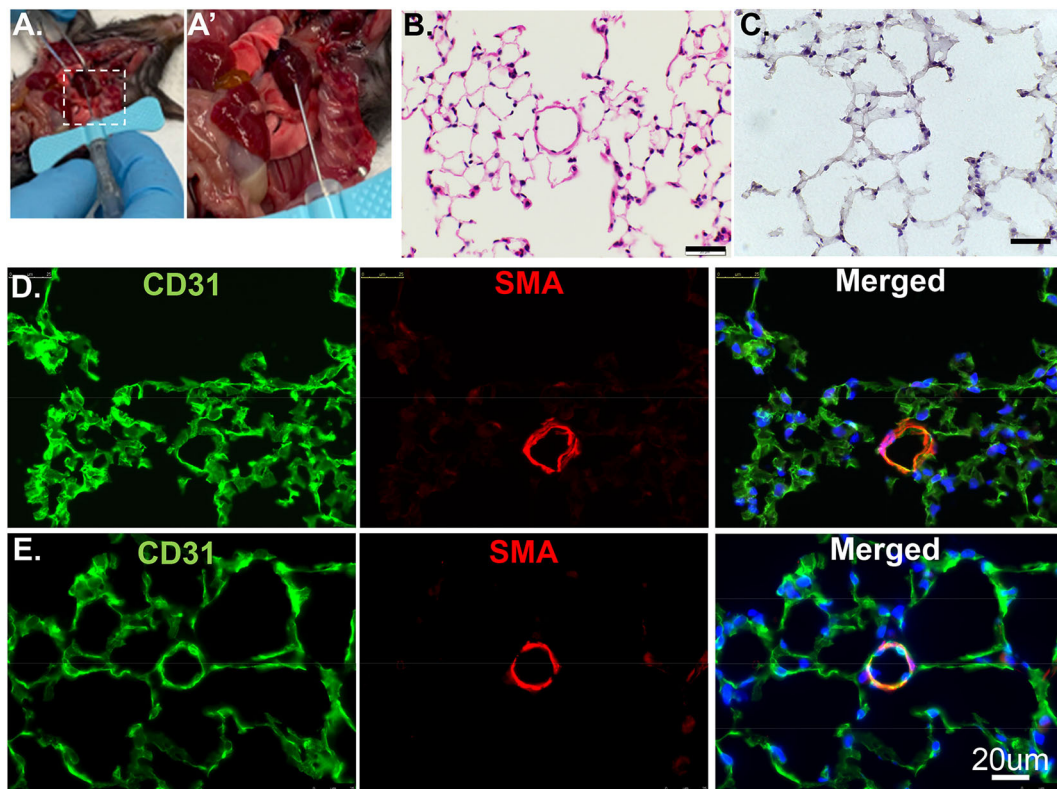


FIGURE 1 | Cross section of lung tissues using H&E or IF. **(A)** The RV is cannulated with a 25G butterfly needle to facilitate switching solutions from flushing buffer to PLP for pulmonary vascular perfusion. **(A')** Enlarged image of dashed rectangle in **A**. **(B)** Wildtype murine lung structure using H&E. Scale bar: 20 μ m. **(C)** Wildtype murine lung structure using H&E after inflation with 100% OCT. Scale bar: 20 μ m. **(D)** The same procedure as in **C** but applied IF staining. CD31: green, stained for endothelial; SMA: red, stained for smooth muscle layer; DAPI: blue, stained for nuclei. **(E)** PLP buffered murine lung described in manuscript providing improved structural integrity.

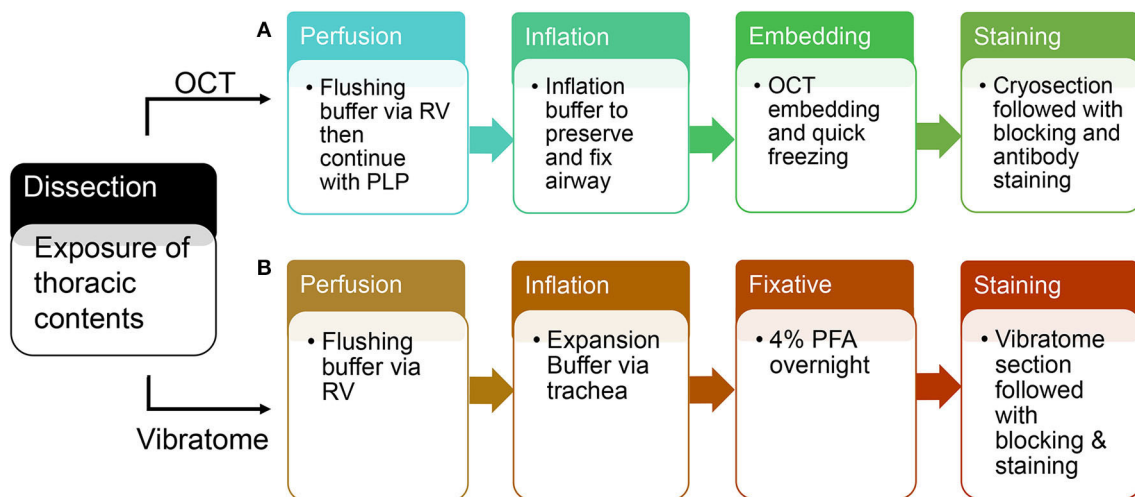


FIGURE 2 | A schematic flow of OCT tissue preparation and vibratome tissue preparation. **(A)** An overview of steps in tissue preparation using OCT. **(B)** An overview of steps in tissue preparation using low melting agarose.

The next day, wash with 1XPBS at 4° and store for future use. Separate and section each lobe with Vibration Slicer at 300 μ m thickness. Block slices in serum overnight and incubate with

primary/secondary antibodies in antibody dilution buffer, and lastly image with Zeiss confocal 880 (each step is summarized in **Figure 2**). Vibratome-sliced sections prepared for STED imaging

will use the same protocol as Confocal, but incubate with STED designated secondary antibody (7), such as STAR Red.

(3) Tissue Clearing

Tissue clearing is effective with any thickness of tissue slice, including the whole lobe. Wash with PBS, then the following serial dilutions of MPBS: 25, 50, 75, 100% (dehydration) to 75, 50, 25%, PBS (rehydration), with an overnight incubation at each step. Incubation time will vary based on the thickness of the sample being prepared, with a minimum of 30 min recommended for thin samples (300 μ m) or overnight for whole lung lobes. The rest of the staining will be performed at 4°. Incubate in serum blocking buffer (the selected serum depends on the species of the secondary antibody, e.g., 5% Goat serum) overnight. Incubate primary antibodies in blocking buffer overnight. Wash slices in the blocking buffer at least three times. Incubate in secondary antibodies at least overnight. Wash slices in the blocking buffer at least three times. After a brief wash in PBS, dehydrate with serial dilutions of MPBS and stop at 100% methanol. Submerge tissue in BABB* and gently agitate until the tissue is colorless. The organ must be as dehydrated as possible, as BABB will fail to clear tissue in the presence of water. Proceed with imaging with Confocal or Light-Sheet Microscopy. *Note: Once combined, BABB is extremely volatile and toxic on skin contact.

RESULTS

Paraffin embedding is the traditional technique to preserve and evaluate lung structure. Paraffin sections are a valuable resource to reveal structural and matrix components details, such as procedures Movat, Trichrome, H&E staining (Figure 1B). However, paraffin embedding is suboptimal for immunofluorescence imaging. In this method, the lung is infused through airways with 3% formalin to preserve the structure and prevent artifacts that can preclude accurate histological evaluation. The fixative also cross-links certain proteins in and on the cells. Thus, antigen retrieval to “unmask” cross-linked antigens is often essential and required. Unfortunately, some antigens can be destroyed by the high-temperature processing used in “unmasking” and are no longer recognized by their designated antibodies. Sometimes, high antibody concentration throughout the tissue may cause non-specific binding as false-negative results.

Embedding in OCT (aka frozen sections) compound, a method with minimal antigen damage is much better for two-dimensional immunofluorescence (IF) imaging. OCT compound was developed specifically to facilitate tissue cryo-sectioning and permeability. Tissue preparation with OCT allows for faster start-to-finish tissue processing, thinner sections (10 μ m), with more consistent antigen penetration for immunolabeling when compared to paraffin preservation and sectioning.

Previously, we used PBS to flush blood from the harvested tissue, 3% formalin as a fixative, then inflated through airways with 100% OCT. After H&E and antibody staining, we found discontinuous layers of alveolar pneumocytes and transparent/matrix basement membrane, which failed to reflect

the real structure (Figures 1C,D). To optimize this procedure, we used Periodate-Lysine-Paraformaldehyde (PLP) and inflation solution to preserve structural integrity. After a flushing buffer, PLP solution fixes the vasculature from the inside out while further clearing the tissue of blood. The inflation buffer optimizes the structural preservation of the alveolar/capillary interface even at the periphery and reduces autofluorescence (Figure 1E). We recommend performing the inflation step as quickly as possible to facilitate rapid coating with OCT and freezing. This step will exclude air bubbles and optimize the overall structure appearance.

To avoid regional sampling bias when analyzing lung structure in a whole-mount context, we applied the expansion solution via trachea and sectioned into 300–600 μ m vibratome slices (the slice thickness can be determined by the working distance of confocal objective lens), or as known as Precision Cut Lung Slices (PCLS). These slices can be easily permeabilized and stained with antibodies CD31 for endothelium and SMA for smooth muscle. We then used confocal Z-stack/reconstitution to assess CD31/SMA-positive cell coverage on individual vessels (Figure 3). Quantification of our results can be achieved by counting the number of overlapped nuclei with or without positive antibody staining (8).

As proof of principle, we used a previously created a NG2-selective reporter mouse and showed that intraperitoneal tamoxifen selectively marks NG2 promoter-driven cells with red fluorescence in capillaries (6). The vibratome slices collected from this transgenic line was stained with RFP and imaged using Stimulated emission depletion (STED) microscopy (Figure 4A). Compared to the out-of-focus blur and an artifact seen by confocal (Figure 4B), STED allowed high resolution and distinction of single positive RFP on the cytoplasm at high spatial density (Figure 4C).

Molecular and optical interrogation of large-sized biological tissue (>1 mm) can be achieved by combining optical tissue clearing and light-sheet microscopy (Figure 5). It enables a true representation of cell location in a whole organ level that allows the immunolabeling of proteins that are essential under normal and pathologic conditions. BABB clearing after systematic tissue dehydration, rehydration, labeling, and then second dehydration provides for a relatively fast protocol for this visualization. Positive SMA stain area quantification can be calculated by processing with the “Surface and Filament Tracer” tool of Imaris (Bitplane) (9).

DISCUSSION

In this manuscript, we outline new, enhanced methodologies to produce high-resolution images that preserve the structure and optimize experimental flexibility on thin-section, thick-sections, and whole organ images. The combination of tissue clearing preparation with lung inflation enables an unparalleled system-based visualization of biological pathways within the whole organ. Inflating the lung gives us a better perspective of distal vs. proximal structures and provides superior spatial detail compared to methods not used in combination.

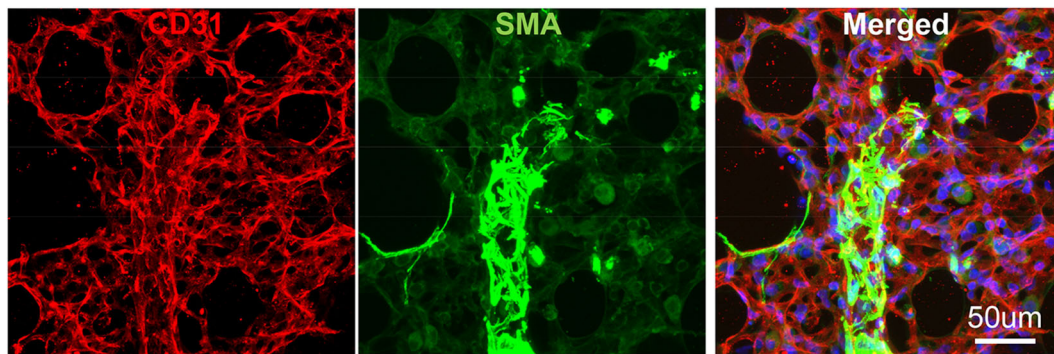


FIGURE 3 | Vibratome section of lung tissues using IF. The wildtype murine lung after flushing was inflated with agarose and proceed with fixative. CD31: green, stained for endothelial; SMA: Red, stained for smooth muscle layer; DAPI: blue, stained for nuclei.

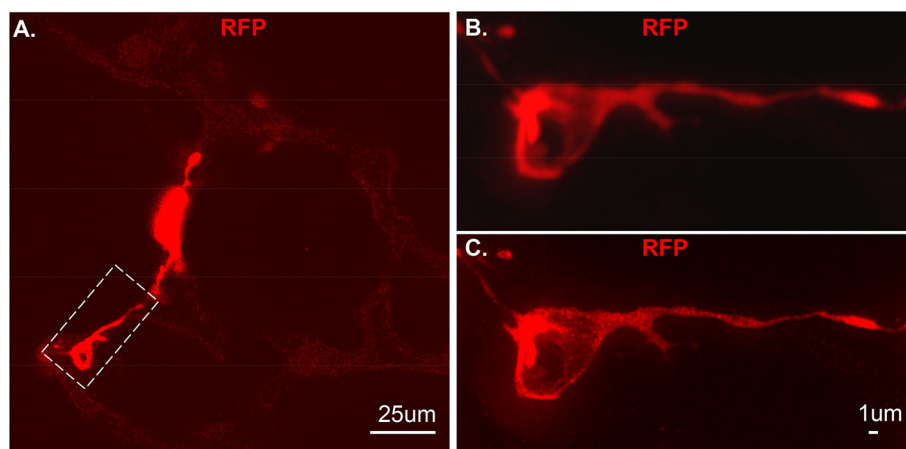


FIGURE 4 | An individual tdT positive cell from vibratome sections. The section was stained using RFP and the secondary antibody used is anti-rabbit-STAR RED. **(A)** An overall view of NG2-tdT lung using STED. A RFP positive cell in dash rectangle was captured using Confocal **(B)** or captured using STED **(C)**. RFP: red fluorescence protein.

The preparation of tissue microslices is a technique that uses a vibratome, or vibrating blade, to produce precision-cut organotypic slices. This allows for increased accuracy and reproducibility when generating precision-cut lung slices (PCLS) compared to alternate methods (10). Tissue slices are time- and cost-effective compared to other *in vivo* work (11, 12). PCLS has been used in human and animal studies focusing on lung anatomy, toxic exposures, infectious disease and immunological studies (13, 14). They can maintain the structural framework at both the tissue and cellular level, making them useful for studying anatomical abnormalities seen in diseases, such as asthma or COPD (15). This makes PCLS the ideal platform to study tissue-specificity and cancer cell selectivity of gene therapy vectors prior to *in vivo* trials (4).

However, PCLS does not come without limitations. PCLS displays a snapshot of the cells and molecules residing in lung tissue at the time of removal and does not capture the heterogeneity seen in some pathological processes. The PCLS model is also static, which means that any testing involving

mechanical stress, such as barotrauma, is limited. Samples have a viability of 3–6 days, limiting the extend that PCLS can model *in vivo* situations (12). While a PCLS is an excellent model for physiologic and toxicologic studies, future investigation is needed to describe methods for improved viability incorporation of immune components to enhance PCLS.

Confocal imaging was patented in 1957 and aimed to overcome the limitations of traditional fluorescence microscopes when viewing thin-cut samples (16). It is superior to conventional microscopy by focusing light inside tissue and limiting the emitted light returned, allowing the specimen to be imaged at a single “point” at a time. This allows the reconstruction of high-resolution, high contrast three-dimensional images without the artifacts that limit conventical microscopy. However, confocal imaging is restricted by its prolonged scanning time, photobleaching secondary to long scan times, causing reduced signal-to-noise ratios, and the inability to simultaneously visualize multiple specimen layers (17).

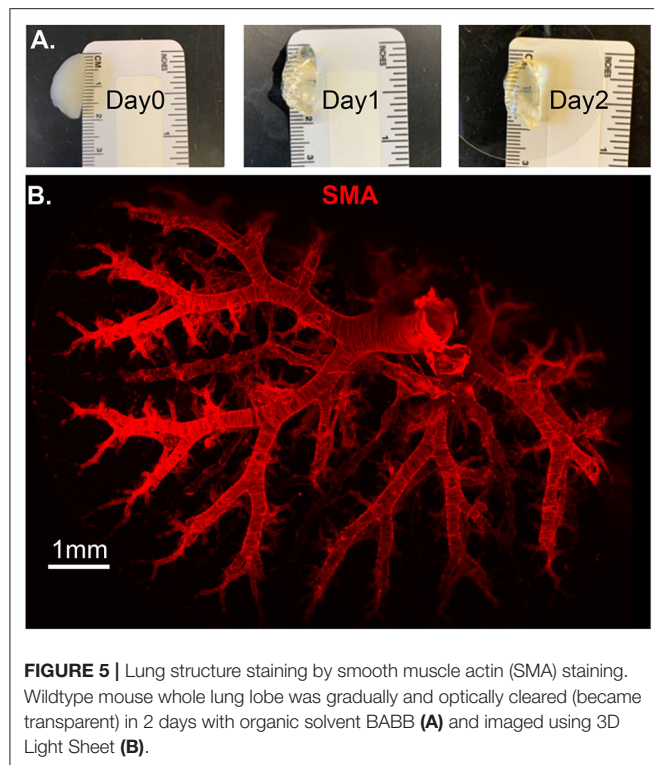


FIGURE 5 | Lung structure staining by smooth muscle actin (SMA) staining. Wildtype mouse whole lung lobe was gradually and optically cleared (became transparent) in 2 days with organic solvent BABB (A) and imaged using 3D Light Sheet (B).

Due to these limitations, techniques were developed which split fluorescence light into multiple planes. This method, or light sheet microscopy, significantly reduced scanning time and some of the limitations seen with confocal imaging. Light sheet microscopy has been employed to investigate inflammation, cancer, hemopoiesis and infection in lung and other tissues (18–21). It has also been used to investigate the distribution of *Aspergillus fumigatus* and identify the local inflammatory response secondary to pulmonary infections (22). However, this technique is limited by optical aberrations and a point-by-point scanning technique (17).

Stimulated emission depletion (STED) microscopy creates super-resolution images at a molecular level. This overcomes the diffraction limitations seen in confocal microscopy to improve resolution and provide nanoscale visualization of individually labeled molecules (23). This technique has been recently used in neuroscience to better visualize dendritic cells and understand their structure and function (24). STED has been applied to lung tissue by detecting filamentous human respiratory syncytial virus particles, providing a better understanding and mechanism of how the virus spreads cell to cell (e.g., microbiota activity) (25). STED microscopy can potentially aid in understanding the molecular mechanisms of receptors and their specific ligands in disease, such as pulmonary hypertension, fibrosis and malignancies (26, 27). An understanding of these molecules at their response can lead to the development of possible pharmaceutical therapies.

Tissue optical clearing is a chemical process aiming to improve light penetration throughout intact tissue, rendering them transparent and allowing fluorescent microscope imaging

(28). Organic-solvent based clearing protocols, such as 3DISCO can achieve high levels of tissue transparency, allowing imaging of large samples, such as brain, tumors and embryos (5, 29). However, the inability to preserve fluorescent protein emission with solvent-based clearing techniques has led to a pursue with aqueous-based solutions. Current techniques include passive immersion, hyperhydration, or hydrogel embedding. While these techniques traditionally do not clear as well as solvent-based methods, they are easy to implement and useful in samples where a wide range of fluorescent dyes and proteins are required. CLARITY is a method for chemically transforming intact biological tissue into a hydrogel-tissue hybrid (3). This hybrid model is amendable to light and macromolecular labels while also retaining structure and biological molecules, such as proteins and nucleic acids. CLARITY has the additional benefit of being plastic-safe, making materials easier to select for processing and imaging safer on most institutional equipment due to lack of necessity for imaging while submerged within an organic solvent. The method has also been successfully applied to numerous animal and human brain models, with the potential to be applied to different, large organ systems. For lung tissue mainly, we found the solvent-based method is more stable and less time-consuming than CLARITY due to variable lipid and protein contents.

In summary, our study provides novel and insightful methods to improve and enhance experimental procedures to preserve pulmonary vasculature with minimal change from regional to a whole-organ large scale. Additionally, all presented methods can have a broader and more comprehensive application, such as assessment of lung pathology in clinical samples.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The animal study was reviewed and approved by IACUC at Boston Children's Hospital and Stanford University.

AUTHOR CONTRIBUTIONS

KY: conception and design, acquisition of data, analysis and interpretation of data, drafted and revised the manuscript. TK and DC: analysis and interpretation of data, drafted and revised the manuscript. YH, WT, ML, AC, MN, XZ, and BR made substantial contributions to the acquisition of data, analysis and interpretation of data. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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When Innate Immunity Meets Angiogenesis—The Role of Toll-Like Receptors in Endothelial Cells and Pulmonary Hypertension

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Toll-like receptors serve a central role in innate immunity, but they can also modulate cell function in various non-immune cell types including endothelial cells. Endothelial cells are necessary for the organized function of the vascular system, and part of their fundamental role is also the regulation of immune function and inflammation. In this review, we summarize the current knowledge of how Toll-like receptors contribute to the immune and non-immune functions of the endothelial cells.

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TOLL-LIKE RECEPTORS

The human body is constantly exposed to exogenous immunological triggers and reacts to these triggers after recognizing their associated molecules. Janeway proposed over 20 years ago that invading microorganisms, such as viruses and bacteria, have specific molecular patterns, so-called pathogen-associated molecular patterns (PAMPs), that trigger recognition by the immune system and named these pattern recognition receptors (PRRs) (1). The PAMPs are sensed by germline-encoded evolutionary conserved host sensors called pathogen recognition receptors or PRRs and form a key element of the innate immune system (2, 3). PRRs are not only present in typical immune cells, such as monocytes, macrophages, and T lymphocytes, but also are expressed in non-immune cells, including endothelial cells (4). Four classes of PRRs are known today: (1) the Toll-like receptors (TLRs) which we will focus on in this review; (2) C-type lectin receptors (CLRs): this large family of cell surface transmembrane receptors bind carbohydrates via specific recognition domains and are important for the immune response to fungal pathogens (5); (3) retinoic acid-inducible gene-I (RIG-I)-like receptors (RLR) which are cytoplasmic sensors of viral RNA or self-processed RNA (6, 7); and (4) nucleotide-binding oligomerization domain (NOD)-like receptors (NLRs) which are cytoplasmic receptors that recognize PAMPs. In contrast to TLRs, NLRs mainly signal via the formation of a multimeric protein complex called the inflammasome (8). Of these four receptor families, TLRs are the most extensively studied class. Although TLRs are an integral part of the innate immune system, their expression is not limited to immune cells but their presence can also be detected in non-immune cells such as endothelial cells, which are further discussed in this review article. The name “Toll-like receptor” originates from the structural homology of TLRs with the Toll transmembrane protein which is important for embryonic development in *Drosophila melanogaster* (9). Toll has an important role for antibacterial defense in *Drosophila*

(10), because mutations in the Toll gene decrease the antimicrobial and antifungal response, leading to increased susceptibility to infection and death (11). The cytoplasmic domain of the Toll protein shares homology with human interleukin-1 (IL-1) receptors (IL-1Rs), leading to similar biochemical signal transduction by IL-1Rs and Toll protein (12). Human transmembrane receptors with structural homology to the *Drosophila* Toll protein were categorized as Toll-like receptors (TLRs) (13). At present, 10 TLRs have been identified in humans (TLR1–10) and 12 (TLR1–9, 11, and 13) in mice (14). TLRs 1–9 are highly conserved in mammals, TLR10 is non-functional in mice, and the human TLR11 gene contains a stop codon that results in lack of production of functional TLR11 (15, 16). The TLRs are localized to either the cell surface membrane or the membranes of intracellular compartments. TLRs 1, 2, 4, 5, 6, and 11 are found at the cell surface and detect extracellular PAMPs, whereas TLR 3, 7, 8, and 9 bind intracellular PAMPs localized to intracellular vesicles such as endosomes and lysosomes, or vesicles derived from the endoplasmic reticulum (17).

In addition to recognizing extracellular and intracellular PAMPs, the immune system plays an important role in the response to non-pathogenic conditions, such as trauma, ischemia, and autoimmune disorders. Polly Matzinger proposed a danger signal model paralleling the concept of PAMPs and suggested that any molecule that is normally not secreted from the cell could activate an immune response if released from the cell in response to injury (18). This damage model further evolved to the concept of damage-associated molecular patterns (DAMPs) (19). Similar to the recognition of PAMPs by PRRs, certain PRRs, including TLR2 and TLR4, bind DAMPs (20). A list of the most common PAMPs (pathogen-associated molecules) and DAMPs (self-molecules) that activate the different TLRs is shown in **Table 1**.

TOLL-LIKE RECEPTOR SIGNALING

TLRs are evolutionary conserved type 1 transmembrane glycoprotein receptors with an ectodomain and a cytosolic domain. The ectodomain contains varying numbers of leucine-rich repeats (LRRs) that are required for ligand binding. Vertebrate TLRs contain 16–28 LRRs and human TLRs 19–25 LRRs. These LRRs form a continuous structure and adopt a horseshoe shape, which facilitates ligand binding (49). After ligand binding, the receptor forms an m-shaped dimer that sandwiches the ligand, bringing the cytoplasmic and transmembrane domains together to initiate signaling (50). Ligand binding causes either homodimerization or heterodimerization of TLRs and the formation of heterodimers promotes ligand diversity, whereas homodimerization increases ligand specificity (51). For example, TLR2 and TLR4 homodimerize but only the TLR4 homodimer can initiate TNF α signaling, whereas TLR1, TLR2, and TLR6 need to heterodimerize to initiate TNF α signaling (52). Differences in LRRs in the extracellular domains along with combinations of different TLRs not only promote ligand diversity to recognize

TABLE 1 | Ligands for various toll like receptors.

TLR	Ligand	
	PAMP	DAMP
TLR1	Bacterial lipoproteins (21, 22)	
TLR2	Soluble peptidoglycan (SPGN) (23)	Biglycan (24)
	Lipoteichoic acid (LTA) (23, 25)	High Mobility Group Box 1 HMGB1(26)
	Pam ₃ CSK ₄ (27)	Monosodium urate crystals (28, 29)
		Calcium Pyrophosphate Dihydrate (28)
		Human cardiac myosin and COC1f fragment of cardiac myosin binding protein-C (30, 31)
TLR3	Viral dsRNA and Polyinosinic:polycytidylic acid (poly(I:C)) (32)	dsRNA from necrotic cells (33)
	siRNA (34)	mRNA (35),
TLR4	Lipopolysaccharide LPS (36)	Biglycan (24)
		High-mobility group box 1 (HMGB1) (26)
		Fibrinogen (37)
		Heparan sulfate (38, 39)
		COC1f fragment of cardiac myosin binding protein-C (30)
TLR5	Flagellin (40)	
TLR6	Diacylated lipoproteins (41)	
TLR7	Guanosine and uridine-containing ssRNA (42)	Human cardiac myosin (31)
TLR8	Single-stranded RNA (ssRNA), bacterial RNA (43–45)	
TLR9	Unmethylated CpG oligodinucleotides (ODNs) from bacterial DNA (46)	Mitochondrial CpG-ODN (47)
TLR10	Human immunization virus-1 (HIV-1) proteins (48)	

Ligands for various Toll-like receptors differentiated into PAMPs and DAMPs. PAMP, pathogen-associated molecular patterns; DAMP, damage-associated molecular patterns; dsRNA, double-stranded RNA; ssRNA, single-stranded RNA; ODN, oligodinucleotides; mRNA, messenger RNA.

a large number of PAMPs but also determine differences in the downstream signaling (53, 54).

Ligand binding then initiates a cascade of downstream signaling via the cytosolic Toll/interleukin-1 receptor (TIR) domain, which is a conserved domain shared by TLRs and the interleukin 1 receptor (IL-1R) superfamily (55). The TIR domains of TLRs dimerize upon ligand binding. This self-association leads to the recruitment of intracellular TIR-containing adaptor proteins via TIR–TIR interactions, forming a trimer. The recruitment of additional intermediates then elongates this trimer. The sequential and cooperative

binding of the TIR domains amplifies the signal and results in a highly sensitive response (56). The main TIR-containing adaptor proteins are myeloid differentiation factor 88 (MyD88), Toll-interleukin 1 receptor (TIR) domain-containing adaptor protein (MAL or TIRAP), and TIR domain-containing adapter molecules (TICAM), such as TIR-domain-containing adapter-inducing interferon- β (TRIF or TICAM-1) and translocating chain-associated membrane protein (TRAM or TICAM-2). These adaptor proteins bind to TLRs and initiate major downstream signaling pathways, including nuclear factor κ -light-chain enhancer of activated B cells (NF- κ B), activator protein 1 (AP-1), and interferon-regulatory factor (IRF) (56, 57).

All TLRs except TLR3 follow the MyD88-dependent pathway-activating mitogen-activated protein kinase kinase kinase 7 (MAPKKK7 or TAK1) which is downstream of TRAF6. TAK1

then leads to activation of both enzyme I κ B kinase (IKK)-activating NF- κ B and MAPK family c-Jun N-terminal kinases (JNK), extracellular signal-regulated kinases (ERK), and p38 leading to AP-1 activation and proinflammatory cytokine production (58). However, endosomal TLR3 and TLR4 signal via a MyD88-independent pathway involving TRIF. TLR3 associates with TRIF directly (59), whereas TLR4 associates with TRIF through TRAM (60). TRIF then activates TRAF3 activating the IRF3/7 signaling pathway leading to type 1 IFN production (61) and TRAF6 activating the NF- κ B and AP-1 signaling response (62, 63).

Activation of these pathways results in the release of cytokines such as interleukin-4 (IL-4), IL-13, tumor necrosis factor- α (TNF- α), IL-1 β , chemokines such as IL-8, monocyte chemoattractant protein 1 (MCP-1), macrophage inflammatory protein-1 β (MIP-1 β), and type I interferons (IFNs), such as IFN- α

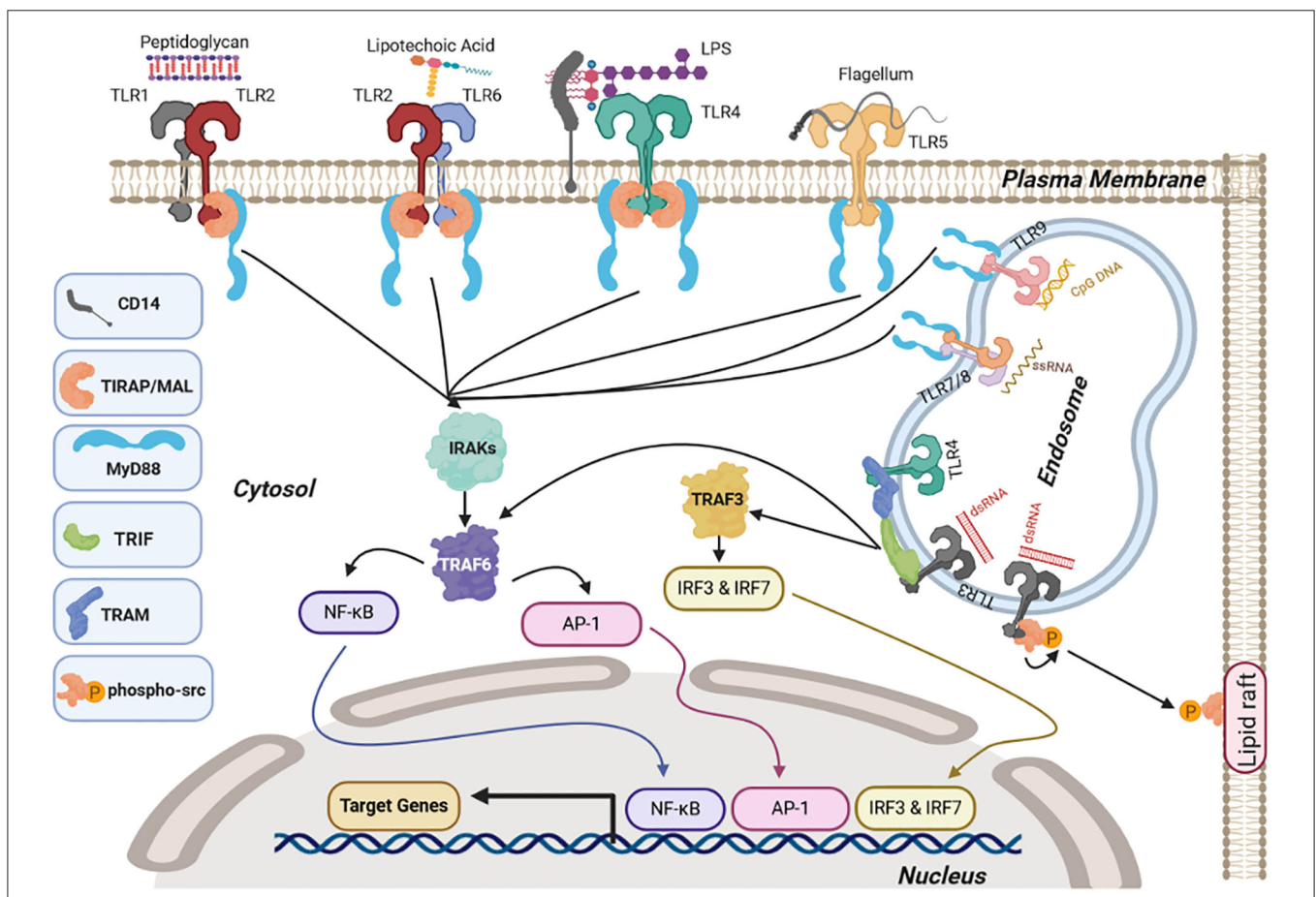


FIGURE 1 | Toll-like receptor (TLR) signaling. Upon ligand binding at the cell surface, TLR1/2, TLR2/6, and TLR4 interact with myeloid differentiation factor 88 (MyD88) via the adaptor molecule Toll-interleukin 1 receptor (TIR) domain-containing adaptor protein (MAL or TIRAP) whereas TLR5 and endosomal TLRs TLR7/8 and TLR9 interact directly through MyD88. This causes activation of nuclear factor κ -light-chain enhancer of activated B cells (NF- κ B) and activator protein-1 (AP-1), signaling via interleukin-1 receptor-associated kinase 4 (IRAK1 and 2) and tumor necrosis factor receptor (TNFR)-associated factor 6 TRAF6. Endosomal TLR3 and TLR4 induce NF- κ B and AP-1 via TRIF associating with TRAF6 along with interferon regulatory factor IRF3/7 signaling via TRIF and TRAF3. Once NF- κ B, AP-1, and IRFs are activated they translocate to the nucleus and activate transcription of target genes, including pro-inflammatory cytokines (NF- κ B and AP-1) and type I interferons (IRFs). TLR3 also induces auto-phosphorylation of the proto-oncogene Src, causing sequestration of Src in lipid rafts. Figures were drawn via biorender.com.

and IFN- β (64). The release of the cytokines varies and depends on receptor, cell type, and species (65, 66).

Not all TLRs signal via the same adaptor proteins; double-stranded RNA (ds-RNA) activation of TLR3 also initiates non-canonical signaling via recruitment and autophosphorylation of the proto-oncogene tyrosine-protein kinase Src. This leads to inhibition of cell migration, proliferation, and cell adhesion via functional sequestration of Src to lipid rafts (67). A summary of the signaling mechanisms downstream of TLRs is shown in **Figure 1**.

EXPRESSION AND ROLE OF TLRs IN ENDOTHELIAL CELLS

Endothelial cells (ECs) form the inner layer of the blood vessel, the tunica intima, and have contact to circulating cells in the lumen and vascular smooth muscle cells in the adjacent tunica media of the blood vessel. Therefore, ECs are uniquely positioned to maintain hemostasis and regulate the vascular tone via release of vasoactive mediators (68). However, ECs also respond to injury via secretion of cytokines, chemokines, and growth factors, as well as via expression of adhesion molecules (69). They hereby regulate the recruitment and activation of immune cells at the site of injury (70). ECs further recognize PAMPs and DAMPs through PRRs, particularly the TLRs (71). ECs from different species and tissues vary in the expression of TLRs. For example, human umbilical cord ECs (HUVECs) express high levels of TLR1-4, but low levels of TLR5-10, whereas human aortic ECs have high levels of all TLRs except for TLR3 and TLR9 compared to human peripheral blood mononuclear cells (PBMCs) as controls (72).

Similarly, the expression of TLRs varies between the different sections of the vascular tree, as major blood vessels have low endothelial expression of TLR2 and TLR4 (73). TLR expression is different among macrovascular and microvascular ECs. For example, baseline TLR4 levels are higher in dermal microvascular ECs than in aortic macrovascular ECs and elevated TLR4 levels correlated with increased chemokine and cytokine expression (74). In addition to mature ECs, TLRs are also found in precursor ECs, such as endothelial colony forming cells (ECFCs). ECFCs are a promising source for postnatal vascularization strategies and tissue repair (75) and are isolated from different sources (cord blood, peripheral blood, and lung) by outgrowth of EC colonies via limiting dilution from blood or tissues (76–79). ECFCs from cord blood and peripheral blood expressed mRNA for all TLRs, and umbilical cord ECFCs showed higher TLR4 expression than peripheral blood ECFCs, HUVECs, and PBMCs (80).

TLRs are also important for the differentiation and reprogramming of cells. Forced expression of a set of transcription factors is a common method to induce pluripotent stem cells. These transcription factors are octamer-binding transcription factor 4 (Oct4), sex determining region Y-box 2 (Sox2), Kruppel-like factor 4 (Klf4), and c-master regulator of cell cycle entry and proliferative metabolism (c-Myc). Lentiviral expression of these transcription factors increased the efficiency of generating induced pluripotent stem cells via

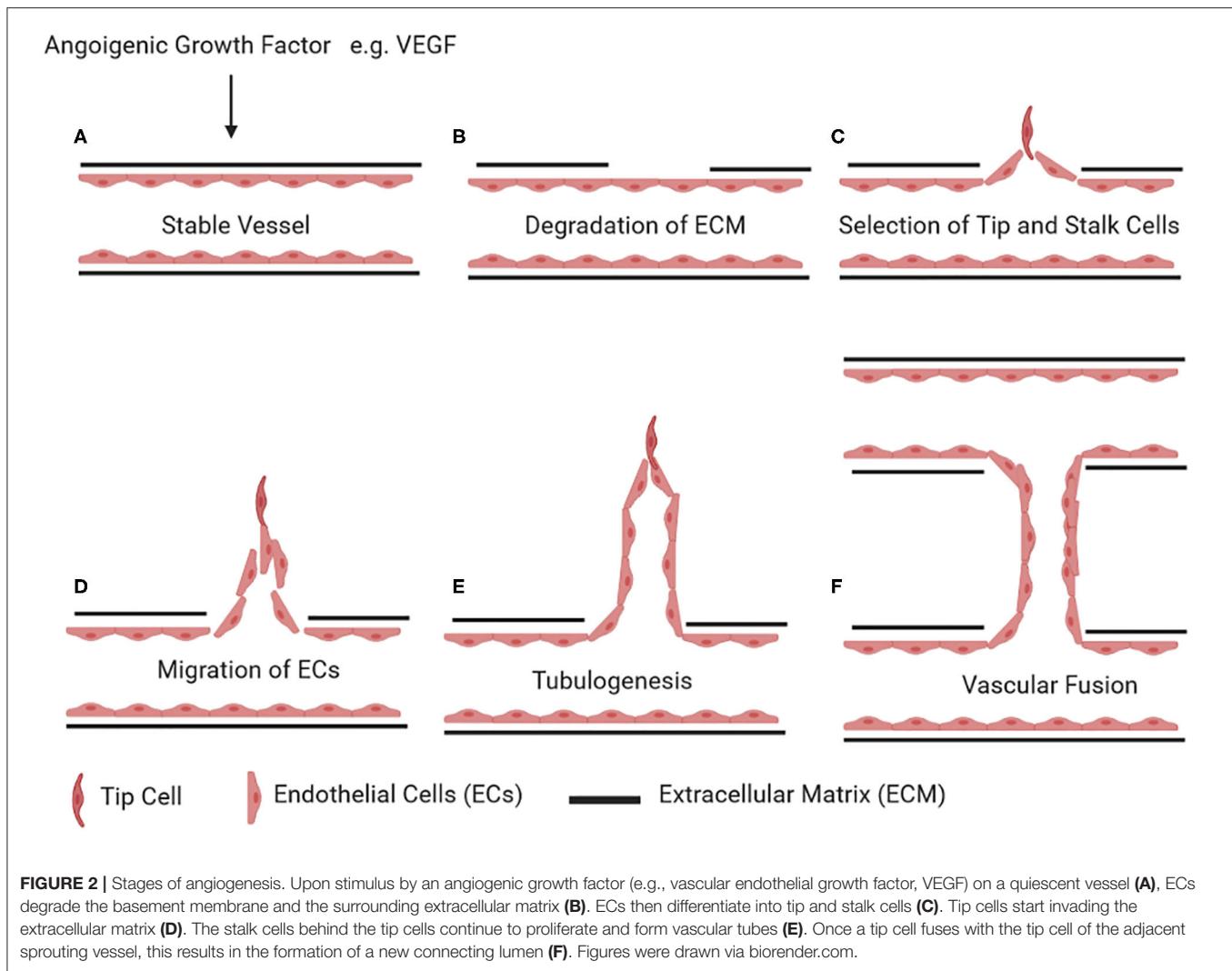
TLR3/TRIF compared to non-viral expression methods (81). In addition, activation of TLR3 by its ligand double-stranded RNA (dsRNA) combined with endothelial growth factors promoted differentiation of human fibroblasts to ECs (82).

ROLE OF TLRs IN BLOOD VESSEL FORMATION AND EXTENSION

The network of blood vessels expands via two main mechanisms: vasculogenesis is *de novo* formation of blood vessels, and angiogenesis is sprouting or splitting of existing blood vessels (83). Sprouting angiogenesis occurs in five steps: (A) After action of angiogenic growth factor (e.g., vascular endothelial growth factor, VEGF) on a quiescent blood vessel, there is (B) degradation of the capillary basement membrane followed by (C) EC proliferation and (D) selection of tip and stalk cells. (E) ECs then migrate from the existing blood vessel to form new vascular tubes (tubulogenesis) and finally (F) connect to another blood vessel by fusion (**Figure 2**) (84).

TLR signaling regulates the complex process of sprouting angiogenesis at different stages of the process: The initial step of angiogenesis requires the degradation of the ECM via matrix metalloproteinases (MMPs) 2 and 9 to facilitate EC migration and tube formation (85) along with degrading endothelial tight junctions (86). Tight junctions (TJ) seal the intercellular gap to maintain endothelial barrier function, and several proteins including occludins, claudins, and zonula occludens form these tight junctions (87). The endothelial tight junction protein claudin-5 increases EC proliferation (88) but decreases EC migration and permeability, hence reducing vascular sprouting (89) similar to ECs lacking zonula occludens 1 (ZO-1), which show less migration and vascular sprouting (90). MMP9 reduced ZO-1, occludin, and claudin-5 levels via proteolysis, resulting in enhanced endothelial permeability, migration, and tube formation, and this effect was reversed with MMP9 gene silencing (86, 91). Cytokines increase MMP9 expression in an NF- κ B-dependent manner via a NF- κ B-binding site on the MMP9 promoter (92). TLRs promote MMP9 expression via NF- κ B and extracellular signal-regulated kinase (ERK1/2) signaling pathways. TLR2 signaling influences endothelial MMP9 expression via ERK1/2 and c-Jun N-terminal kinase (JNK) signaling pathways along with MMP9-mediated reduction in tight junction proteins (93). Chlamydia pneumoniae infection promoted expression of VEGF and MMP9 in HUVECs via TLR2 and TLR4 (94). TLR4 results in EC hyperpermeability via NF- κ B and AP-1-induced stromal interaction molecule 1 (STIM1) expression in human lung microvascular ECs (HLMVECs) (95). TLR2/6 activation augments EC permeability via reduced claudin-5 expression and disappearance of tight junctions, which was partly mediated by ERK1/2 (96).

In addition to tight junctions, TLRs also influence the adherens junctions between ECs. Adherens junctions are cell-cell junctions with cadherins, such as vascular endothelial (VE-) cadherin, connecting neighboring plasma membranes via homophilic interactions (97). TLR2 stimulation reduced the expression of VE-cadherin resulting in increased EC detachment



and reduced EC migration while increasing MMP2/9 activity and EC apoptosis in human saphenous vein ECs (98). Similarly, in HLMVECs, TLR3 activation reduced the expression of claudin-5, ZO-1, and VE-cadherin (97).

TLR signaling further influences the expression of growth factors with a central role in regulating vascular growth. In human intestinal microvascular cells, bacterial ligands via TLRs increased angiogenic factors such as VEGF, VEGF receptor 2 (VEGFR-2), IL-8, and phosphorylation of focal adhesion kinase (p-FAK) (99). Of note, FAK promotes EC survival, angiogenesis, and vascular network stability (100). Among these growth factors, VEGF-A is a central angiogenic molecule that directs migration of endothelial tip cells during angiogenesis (101). Biglycan-stimulated TLR2 and TLR4 signaling increases VEGF-A levels, resulting in endothelial proliferation, migration, and tube formation. TLR2/4 stimulation activates NF- κ B, which interacts with the hypoxia-inducible factor-1 α (HIF-1 α) promoter to augment HIF-1 α levels. HIF-1 α in turn binds to the VEGF-A promoter increasing VEGF-A levels (102).

Activation of TLR5 by the bacterial ligand Flagellin promotes endothelial tube formation in human microvascular ECs and HUVECs through activation of the phosphoinositide 3-kinase (PI3K)/AKT1 signaling pathway (103). In rat, aortic cell culture TLR5 activation increased microvessel formation along with vessel survival but had no effect on *de novo* blood vessel formation (104).

However, TLRs also promote angiogenesis in a VEGF-independent manner. Mycoplasma lipopeptide MALP-2 in HUVECs and human monocytes/macrophages via TLR2/TLR6 promoted activation of granulocyte-macrophage colony-stimulating factor (GM-CSF). GM-CSF promoted angiogenesis in these endothelial cells while no enhanced VEGF levels in study were seen, suggesting that VEGF was not responsible for this angiogenesis in this regard (105). The end product of lipid oxidation, ω -(2-carboxyethyl)pyrrole (CEP), is generated during inflammation, wound healing, and aging. CEP interacted with TLR1/TLR2 heterodimer, promoting angiogenesis via MyD88-dependent NF- κ B signaling in multiple EC lineages from human umbilical vein, mouse lung, or aorta independent

of VEGF as CEP-mediated effects were unaffected by VEGFR kinase inhibition (106). In lung ECs, TLR4 signaling increases ERK-mediated activation of the transcription factor Forkhead box protein C2 (FoxC2), a transcription factor associated with lymph angiogenesis and endothelial specification by promoting delta-like 4 (DLL4) expression (107). DLL4 is a NOTCH ligand, which in angiogenic vasculature is associated with filipodia-rich endothelial tip cell formation responsible for guiding new sprouts (108).

Anti-TLR2 antibodies promoted angiogenesis in HUVECs in a manner similar to stromal cell-derived factor 1 (SDF1 also known as C-X-C motif chemokine 12 or CXCL12). This effect was mediated by Gi-protein-coupled receptor C-X-C motif chemokine receptor 4 (CXCR4) via ERK 1/2 and AKT, which was completely abolished by blocking G-protein and CXCR4, indicating a potential cross talk between TLR2 and CXCR4 pathways (109).

In contrast, TLR3 activation inhibited EC migration and tube formation in HUVECs via phosphorylation of Src independent of TRIF, TRAF3, MyD88, and IRF3 signaling (67). In primary choroidal ECs, TLR3 stimulation was anti-angiogenic and a TLR3 agonist reduced corneal neo-vascularization (34). Similarly, dsRNA of viral origin and polyinosinic:polycytidylic acid (poly(I:C)) promoted apoptosis and inhibited angiogenesis and migration in HUVECs (110). Our group has shown that TLR3 deficiency impaired migration of human pulmonary-artery ECs via caspase-3-dependent apoptosis (111). Similar to TLR3, TLR9 reduces angiogenesis, because the TLR9 agonist ODN1826 suppresses aortic angiogenesis, inhibits migration of tip cells in aortic ring assay, and suppresses corneal neovascularization *in vivo* (112). Based on these findings, activation of some TLRs appears to promote angiogenesis by downregulation of EC tight junction and adherens junction proteins, induction of angiogenic growth factors, increased proliferation, and salvation from apoptosis, but other TLRs, such as TLR3, seem to have anti-angiogenic effects following stimulation. However, further in-depth studies are required to fully understand how signaling through the different TLRs contributes to the various stages of angiogenesis and to identify the context-dependent effects of TLR signaling on angiogenesis.

ENDOTHELIAL TLRs AND INFLAMMATION

Blood vessels are the main highway for inflammatory cells to travel to the site of injury. Inflammation and angiogenesis are therefore intricately linked processes, and signaling pathways that promote inflammation frequently also facilitate angiogenesis by inducing EC proliferation and migration. It is therefore not surprising that many proinflammatory cytokines stimulate angiogenesis, including IL-6. In mouse aortic ring assays and lung endothelial cells and HUVECs, IL-6 stimulation increased angiogenesis independent of VEGF (113). One important pathway driving IL-6 expression is the NF- κ B pathway, and every member of the TLR family can signal via NF- κ B (114).

TLR expression varies between different EC populations. TLR4 levels were found to be higher in microvascular ECs

compared to macrovascular ECs, resulting in higher NF- κ B and IL-6 levels in microvascular ECs (74). Similarly, TLR2 activation promotes in human lung microvascular EC expression of cytokines and adhesion molecules, leading to adhesion of neutrophil granulocytes (115, 116). Similar to TLR2 and TLR4, TLR9 activation causes NF- κ B-mediated increase in tissue factor levels favoring a procoagulant phenotype in human coronary-artery ECs (117). In rat PAECs, TLR9 activation leads to NF- κ B-mediated IL-6 production (118). In human intestinal microvascular ECs, flagellin-mediated TLR5 activation promotes expression of leukocyte adhesion molecules such as intercellular adhesion molecule 1 (ICAM-1), leading to increased transendothelial migration of leukocytes (119). TLR9 activation further enhances neutrophil adhesion on HUVECs (120). These data show that TLR signaling in ECs also promotes cross talk between ECs and inflammatory cells.

As with the foreign molecules eliciting TLR responses, the internal molecules of the body also act as DAMPs to promote immune responses via TLRs (Table 1). One of the DAMPs is high-mobility group box 1 (HMGB1). HMGB1 is an endogenous inflammatory mediator released from dying and activated cells during many pathogenic conditions including pancreatitis, cancers, atherosclerosis, and myocardial infarction (121, 122). An interesting mechanism of inflammatory signaling in ECs is the TLR-high-mobility group box 1 (HMGB1) axis. HMGB1 is an alarmin that is released in response to injury from necrotic cells, triggering an inflammatory response (123). HMGB1 signals via TLRs 2, 3, 4, 7, and 9 (26, 123, 124) to induce expression of pro-inflammatory cytokines through MyD88/NF- κ B signaling (124), which in turn increases secretion of HMGB1, creating a positive feedback loop to amplify the HMGB1-mediated inflammatory effect (125). Likewise, extracellular histone-mediated activation of TLR2/TLR4 increases tissue factor expression via NF- κ B and AP-1, promoting thrombus formation in human coronary artery ECs (126). TLR-mediated inflammatory signaling can also utilize G proteins. The intercellular domains of TLR2, TLR3, and TLR4 contain a consensus motif for binding of pertussis toxin-sensitive heterotrimeric G proteins Gai/o. Gai/o activates MAPK and Akt and interferons downstream of TLR2, TLR3, and TLR4 in ECs while having no effect on NF- κ B signaling (127).

TLR 4 is the prototypical (128) and most studied TLR in literature. The functions of TLR4 as previously discussed not only are limited to its immunological role but also extend to various aspects of the vascular biology. One of the major cardiovascular disorders and the top most cause of death in the developing world is the coronary artery disease (CAD) or the ischemic heart disease (IHD), accounting for >9 million deaths globally in 2016 (129). In the heart, all the TLRs are expressed with the expression of TLR4 being the highest (130) and hence is a very important for the molecular pathogenesis of IHD. TLR4 along with TLR1 and TLR2 is highly expressed in human atherosclerotic plaques (131), with the highest expression in the shoulder regions of the plaques in the coronary arteries where the incidence of plaque rupture is highest (132). TLR4 significantly raised the levels of tissue factor (TF), a critical initiator of blood clotting from endothelial cells actively contributing to arterial thrombus formation (133). LPS-mediated TLR4 activation of human coronary artery endothelial

cells resulted in increased IL-1 β and TNF- α which are elevated in congestive heart failure (CHF) and CAD, hence contributing directly to their pathogenesis. Furthermore, TLR4 activation resulted in reduced cardiac function in mice whereas TLR4 deficiency promoted survival and reduction in septic shock and myocardial ischemia-induced cardiac dysfunction (134).

In addition to TLR4, endosomal TLR7 and TLR9 also play an important role in atherosclerotic lesions. Under normal conditions, major arteries of the body have a negligible expression of TLR7 and TLR9 (73). However, TLR7 levels were increased in endothelial cells, smooth muscle cells, and macrophages of mouse atherosclerotic lesions of the aortic arch (135). In apoE*3-Leiden mouse restenosis model, TLR7/9 activation significantly led to femoral artery cuff intimal hyperplasia and accelerated atherosclerosis and blockade of TLR7/9 significantly reduced neointima formation, atherosclerosis, and macrophage cytokine production (136). TLR7 activation may differentially respond according to disease condition. In patients with carotid endarterectomy, TLR7 was higher in atherosclerotic plaques, yet elevated TLR7 expression in the plaques was associated with better outcomes by production of anti-inflammatory cytokines (137).

RNA is another important DAMP that activates TLR3 (35, 138). We and others have shown that TLR3 expression and signaling are important in vascular biology and pulmonary hypertension (PH) (111, 139).

TLRs IN PULMONARY HYPERTENSION

Pulmonary hypertension (PH) is a chronic, progressive disorder of the lung vasculature characterized by abnormal pulmonary-artery vasoconstriction and remodeling, leading to right heart failure and death (140). Idiopathic pulmonary arterial hypertension (iPAH) shows dysregulated EC differentiation and growth (141), and EC dysfunction is now recognized as a central process in the initiation and progression of PH (142). There are multiple aspects of EC dysfunction, which include the emergence of apoptosis-resistant, hyperproliferative ECs, dysregulated release of mediators from ECs, and endothelial-to-mesenchymal transition. One concept suggests that during development of PAH, EC apoptosis results in the selection of these apoptosis-resistant, hyperproliferative ECs. In our recent publication, we showed TLR3 deficiency in pulmonary-artery ECs from PAH patients and in lungs from a rat model of severe PH with occlusive arteriopathy (111). Treatment with the TLR3 agonist and poly(I:C) reduced severe PH and occlusive arteriopathy in rats and increased endothelial TLR3 expression in an interleukin-10 (IL-10)-dependent manner (111). Our results mirror the protective role of TLR3 signaling after balloon injury in large systemic arteries (139). However, other conflicting results seemingly contradict a protective role of TLR3 in the vasculature, but these findings may be due to a more severe degree of endothelial injury in the model system (143). Because knockout of type I IFN receptor and type I IFN treatment have frequently been associated with mostly reversible PH in highly preselected patient groups

with significant comorbidities (144, 145), further evaluation of TLR3-targeted therapy is required in PAH. However, type I IFN therapy seems to reduce severe angioocclusive PH in rats (146).

Recently, the HMGB1–TLR4 axis has emerged as a potential driver of pulmonary vascular remodeling in PAH. HMGB1 was elevated in concentric and plexiform lesions from patients with iPAH and in the lungs of mice exposed to chronic hypoxia (147). In addition, monocrotaline (MCT), an EC toxin causing severe PH, enhances the release of HMGB1 from injured ECs (148). Pro-inflammatory cytokines promote HMGB1 secretion, and therefore HMGB1 secretion and TLR4 may be part of a positive feedback loop enhancing inflammation and lung vascular remodeling in PAH. In addition, activation of the HMGB1/TLR4 axis in rats exposed to chronic hypoxia caused a significant decline in bone morphogenic protein receptor 2 (BMPR2), connecting HMGB1 with a well-known pathway hypomorphism in PAH pathobiology (149, 150). An inhibitor of HMGB1–TLR4 interaction has been characterized as a novel potential therapeutic, translating the findings with the TLR4–HMGB1 pathway to a potential clinical treatment in PAH, although clinical evaluation is necessary as a next step after the initial preclinical study (151, 152). Therapeutic strategies targeting TLRs in PH could be a potential avenue where both agonists and antagonists can improve pathogenesis of PH, depending on the TLR in question. In contrast to reduced PH following treatment of PH rats with the TLR3 agonist poly(I:C) (111), P5779, a blocker of the HMGB1–TLR4 interaction, reduces PH in rats (151, 152). Although other different TLR agonists and antagonists are available, there is a need for further study of TLR signaling in PH to identify additional targets depending on the role of the particular TLR in PH pathobiology.

In addition, TLR signaling can also contribute to another pathogenic mechanism in PAH, endothelial-to-mesenchymal transition (EndMT). EndMT is a process by which ECs acquire a mesenchymal cell phenotype. There is growing evidence that EndMT contributes to development and progression of pulmonary vascular remodeling and severe PAH (153–156). TLR4 activation promoted EndMT and expression of the progenitor cell marker c-kit (CD117) in mouse pulmonary ECs, indicating a potential connection between TLR4 and EndMT (157). Our group has recently shown that clonally expanded CD117⁺ ECs promote the formation of occlusive arteriopathy in rats exposed to chronic hypoxia and that CD117⁺ ECs undergo EndMT *in vitro* and *in vivo* (158). Hence, growing literature indicates that CD117⁺ ECs represent a stem-like EC population in the developing and adult lung (159, 160) and that CD117⁺ ECs contribute to lung vascular remodeling in PAH (158).

In PAH, not only ECs but also smooth muscle cells (SMCs) play a central role in the pathogenesis of the disease (161). Normal pulmonary-artery SMCs most abundantly expressed TLR3, TLR4, and TLR6. TLR3 and TLR4 co-activation resulted in IL-8 release whereas TLR3 activation alone promoted IP10 and endothelin 1 release (162). HMGB1-mediated activation of TLR4 in pulmonary-artery SMCs augmented SMC proliferation and migration along with a decline in bone morphogenic protein receptor 2 (BMPR2) signaling. Based on the dysfunctional

BMPR2 pathway in PAH, it is likely that the HMGB1/TLR4 pathway has a pathogenic role in PAH (149). BMPR2-deficient mice had elevated levels of TLR4 in pulmonary-artery SMCs and lungs. LPS stimulation then leads to a significant increase in IL-6 and IL-8 production (163). These data indicate that TLR4-mediated downregulation of BMPR2 plays an auto-enhancer role driving TLR4 expression in the form of a vicious circle, which contributes to the pathogenesis of PAH. However, hypoxia reduces TLR4 expression and TLR4 knockout mice spontaneously develop PH, suggesting a protective role of TLR4 in PAH (164). This work coincided with a similar observation in ECs, which showed that hypoxia exposure reduced the expression of TLR4 and downstream nuclear translocation of AP-1 in cultured HUVECs and HPAECs (165). In stark contrast, some studies suggest that TLR4-deficient mice were protected from chronic hypoxia-induced PH (166, 167).

In summary, the existing literature indicates that TLRs have functions that exceed by far the induction of an innate immune response in ECs. Instead, TLR signaling is tightly connected with crucial elements of EC function, such as proliferation, apoptosis, angiogenic sprouting, and migration. Potential therapies targeting TLR–ligand interactions or using TLR agonists have emerged in multiple vascular diseases, in

particular in PH, but careful preclinical and clinical evaluation is required when modulating TLR signaling because of the highly conserved and multifaceted effects of TLR signaling in ECs.

AUTHOR CONTRIBUTIONS

AB, AT, and LF conceived the manuscript. AB and LF wrote the initial draft of the manuscript. All authors edited the manuscript and approved the final version.

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Metformin in Pulmonary Hypertension in Left Heart Disease

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Metformin is ubiquitously used in the management of Type II Diabetes Mellitus (DMII). Over the years, our growing knowledge of its therapeutic potential has broadened its use to the treatment of infertility in polycystic ovarian syndrome, gestational diabetes, and even obesity. Recently, it has been suggested as a novel therapy in cardiovascular disease (CVD). Given that CVD is the leading cause of death in patients with DMII, with ~ 75% dying from a cardiovascular event, the intersection of DMII and CVD provides a unique therapeutic target. In particular, pulmonary hypertension (PH) related to CVD (Group II PH) may be an optimal target for metformin therapy. The objective of this review article is to provide an overview of the pathophysiology of PH related to left heart disease (PH-LHD), outline the proposed pathophysiologic mechanism of insulin resistance in heart failure and PH-LHD, and evaluate the role metformin may have in heart failure and PH-LHD.

Keywords: pulmonary hypertension, heart failure, therapeutics, metformin, HFpEF

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METFORMIN AND CARDIOVASCULAR DISEASE

Several large observational cohort studies have outlined the merits of metformin in CVD in diabetics, including improved outcomes in coronary artery disease (CAD), myocardial infarction (MI), and heart failure (HF). Furthermore, data suggests that metformin may avert the progression and/or development of HF (1). The most current data corroborating this is from supplemental analysis of the EMPA-REG OUTCOME trial, in which patients were randomly assigned to receive empagliflozin 10 mg or 25 mg or placebo daily. Outcomes from this trial include CV death, non-fatal MI or stroke, and hospitalizations. Metformin use, even in the placebo arm of the trial, was associated with improved cardiovascular (CV) outcomes including CV death, HF hospitalizations, major adverse cardiac events, and all-cause mortality (2). The benefit of metformin in CVD is so profound that the American College of Cardiology (ACC) has delineated its benefits in its 2018 ACC consensus, “New Diabetic Drugs to Reduce CV Outcomes” (3). Despite this growing evidence, data on metformin’s potential therapeutic role in specific CVD processes is limited.

PULMONARY HYPERTENSION DUE TO CARDIOVASCULAR DISEASE

Pulmonary hypertension is a complex, heterogeneous, multisystem syndrome that affects 10–20% of the general population (4). It results from a panyasculopathy that restricts blood flow in the distal pulmonary arteries with eventual maladaptive consequences on the right ventricle. It is

a rapidly progressive, debilitating disease process that leads to right ventricular failure, and ultimately death. With its heterogeneous pathophysiology and increasing prevalence, the World Health Organization (WHO) created the Dana Point Clinical Classification, most recently updated by the 6th World Symposium on Pulmonary Hypertension which catalogs the disease into 5 categories based on etiology and pathophysiology (5). The most commonly encountered form of PH is WHO group 2, also known as pulmonary venous hypertension due to left heart disease (4, 6).

PH is estimated to affect up to 35% of patients with left heart disease (LHD). PH-LHD is defined as a mean pulmonary artery pressure of > 20 mmHg with a concomitant pulmonary artery wedge pressure of > 15 mmHg (7). As we uncover more about PH-LHD, it is becoming increasingly apparent that the pathophysiologic aberrations and the phenotypic variations are highly dependent on the underlying cardiovascular disease process, e.g., heart failure with reduced ejection fraction (HFrEF) vs. heart failure with preserved ejection fraction (HFpEF) (4, 7, 8).

HFpEF is characterized by signs and symptoms of HF in the setting of abnormal left ventricle (LV) systolic function. It is defined by a LV ejection fraction (LVEF) of $\leq 40\%$, progressive LV dilation, eccentric remodeling, reduced LV systolic elastance, and increased arterial elastance. Regardless of the etiology, there are a number of beneficial therapeutic interventions available including beta-adrenergic receptor blockers, aldosterone antagonists, ace-inhibitors, angiotensin receptor blockers, hydralazine and isosorbide dinitrate, intracardiac defibrillators, and cardiac resynchronization therapy. Unfortunately, the benefits of these therapies do not translate to PH-HFrEF, as HFpEF patients with PH continue to have worse outcomes than those without (9–12).

In contrast, HFpEF is a complex, heterogeneous, systemic syndrome characterized by signs, and symptoms of HF in the setting of a normal left ventricular ejection fraction. A number of cardiac and extracardiac aberrations and comorbidities result in HFpEF's heterogeneity and phenotypic variation (13, 14). Unlike in HFrEF, there are a paucity of beneficial therapeutic interventions available in HFpEF raising the question of whether targeting specific HFpEF sub-phenotypes may lead to, as of yet, elusive successful therapies. One such unique phenotype is pulmonary hypertension (PH).

PH is prevalent in ~ 60 – 70% of HFrEF patients and 80% of HFpEF patients and is also associated with a higher morbidity and mortality (15). The mechanism and pathophysiology of PH in HF, as alluded to previously, is complex and incompletely understood. Initially elevated LV end-diastolic pressure and secondary mitral regurgitation leads to chronically elevated left atrial pressure and increased pulmonary artery wedge pressure. This contributes to both pulmonary vasoconstriction and alveolar capillary stress failure. Eventually over time there is increased pulmonary vascular remodeling, endothelial dysfunction, intimal fibrosis, and venous hypertension. The venous congestion from left atrial hypertension in conjunction with reactive arterial vasoconstriction and remodeling is referred to as combined pre- and post-capillary PH, defined hemodynamically as a mean pulmonary artery pressure > 20

mmHg with a pulmonary artery wedge pressure > 15 mmHg and pulmonary vascular resistance ≥ 3 Woods units (5, 7). Other pathophysiologic mechanisms have also been implicated in PH-LHD as well-including intrinsic myocyte stiffness, extracellular matrix alterations, microvascular dysfunction, inflammation, and metabolic dysfunction (7, 16). Given the lack of available therapies, investigating alternative pathophysiologic mechanisms such as metabolic syndrome or insulin resistance as novel therapeutic targets is paramount.

METABOLIC SYNDROME, INSULIN RESISTANCE, PH-LHD

Heart failure (HF) is a heterogeneous syndrome with regards to demographics, underlying etiology, disease course, and prognosis. To better categorize and risk stratify this complex syndrome, different schema have been developed including the American Heart Association (AHA) stages as well as the New York Heart Association classification (17). Over the last several years, greater efforts have been made to sub-phenotype HF in order to improve risk stratification and facilitate therapeutic interventions. This stratification has led to a greater awareness of the role of metabolic syndrome in HF as well as PH-LHD.

DM is closely associated with HF. A meta-analysis of HF clinical trial populations uncovered that the prevalence of DMII in HF patients approximates 30% (18). This increases to 40% in African Americans. Additionally, patients with concomitant HF and DM have increased morbidity and mortality. In the Candesartan in Heart Failure Assessment of Reduction in Mortality and Morbidity trial (CHARM), DM was an independent predictor of cardiovascular death and hospitalization in patients with HFpEF (19). More recently, DMII has been linked as an independent risk factor in the development of HF as publicized by data derived from the Framingham Risk Study. Both men and women with DMII were 5x more likely to develop HF even after taking into account incidence of CAD (20). Furthermore, 60% of HF patients with normal fasting glucose and insulin levels were found to have abnormal responses to oral glucose tolerance testing—implicating insulin resistance as an underlying pathophysiologic contributor to HF (20). Given this, traditional and novel glycemic control agents are being increasingly studied as a therapeutic agent in the management of HF. A list of such agents is summarized in **Table 1**.

It is hypothesized that insulin resistance directly affects myocardial structure and function through lipotoxicity, free fatty acid oxidation, oxidative stress, impaired nitric oxide bioavailability, and mitochondrial dysfunction and fibrosis (27–31). Impaired insulin signaling diminishes myocardial glucose extraction and uptake resulting in increased fatty acid oxidation (20). Free fatty acid oxidation in turn leads to accumulation of lipid storage molecules including triglycerides, ceramide, and diacylglycerol in cardiomyocytes. Ceramide induces apoptosis and impaired mitochondrial function, which eventually results in ventricular dysfunction and remodeling. The histologic outcome is similar to what is observed in pressure overload, volume overload, or myocardial infarction (20).

TABLE 1 | Antiglycemic agents and cardiovascular outcomes.

Agent	Trial	CV death	All-cause death	HF hospitalizations
Empagliflozin	EMPA-REG (2)	0.62 (0.49–0.77)	0.68 (0.57–0.82)	0.65 (0.50–0.85)
Canagliflozin	CANVAS (21)	0.87 (0.72–1.06)	0.87 (0.74–1.01)	0.67 (0.52–0.87)
Saxagliptin	SAVOR-TIMI53 (22)	1.03 (0.87–1.22)	1.11 (0.96–1.27)	1.27 (1.07–1.51)
Sitagliptin	TECOS (23)	1.03 (0.89–1.19)	1.01 (0.90–1.14)	1.00 (0.83–1.20)
Liraglutide	LEADER (24)	0.78 (0.66–0.93)	0.85 (0.74–0.97)	0.87 (0.73–1.05)
Semaglutide	SUSTAIN-6 (25)	0.98 (0.65–1.48)	1.05 (0.74–1.50)	1.11 (0.77–1.61)
Exenatide	EXSCEL (26)	0.88 (0.76–1.2)	0.86 (0.77–0.97)	0.94 (0.78–1.13)

All numerical values including cardiovascular death, all cause death, and heart failure hospitalizations are presented as hazard ratios with the 95% confidence interval. CV indicates cardiovascular; HF indicates heart failure.

The role of insulin resistance in HF and PH-LHD has been well-documented in preclinical models. For example, altered glucose metabolism involving downregulation of 5'adenosine monophosphate-activated protein kinase (AMPK) has been implicated in the pathogenesis of PH-LHD. Mice exposed to a high fat diet and L-NAME (NO synthase suppression) to induce metabolic syndrome and hypertension were found to have many of the hallmarks of HFpEF including exercised intolerance, lung congestion, LV hypertrophy, cardiac fibrosis, reduced myocardial capillary density, increased left ventricular filling pressure, reduced LV global longitudinal strain despite preserved LV ejection fraction (EF), and decreased contraction velocity with impaired relaxation (32).

In another preclinical model, metabolic syndrome, and left ventricular dysfunction were induced in mice through supracoronary aortic banding and a high fat diet supplemented by olanzapine. In this model, metabolic syndrome exacerbated group 2 PH via vascular remodeling. There was increased macrophage accumulation, interleukin 6 (IL-6), and signal transducer and activator of transcription 3 (STAT3). These findings were associated with increased proliferation of smooth muscle cells in pulmonary arteries, remodeling of distal pulmonary arteries, and PH (33).

Metabolic changes resulting from abnormal glucose handling and insulin resistance were found to precede the structural and functional aberrations in cardiac muscle. This was demonstrated through glucose-6-phosphate (G6P) via mammalian target of rapamycin (mTOR) and endoplasmic reticulum stress in *ex vivo* animal models, *in vivo* animal models, and in humans with HF (34). While under stress such as increased inotropic demand, cardiac myocytes undergo both metabolic, and structural remodeling. Metabolically, carbohydrates become the main energy source. Structurally, cardiac hypertrophy has been linked to increased activity of mTOR, a regulator myocardial protein synthesis (35). Rats pretreated with various energy sources including glucose, non-carbohydrate energy substrate, or glucose analog underwent *ex vivo* analysis of cardiac muscle exposed to high workloads. There was increased G6P accumulation, mTOR activity, endoplasmic reticulum stress, and impaired contractility regardless of energy source. This was not appreciated in rats pretreated with rapamycin (an mTOR inhibitor) or metformin (an AMPK activator). This was re-demonstrated with *in vivo* rat models. Rats exposed

to aortic constriction had increased glucose tracer analog uptake and contractile dysfunction utilizing micro-PET imaging (34). These changes preceded LV dilation. Furthermore, humans with heart failure were found to have decreased G6P and endoplasmic reticulum stress markers once the heart was mechanically unloaded utilizing left ventricular assist devices (34). Pathophysiologically, it is hypothesized that increasing workload surpasses cardiomyocyte glucose uptake capability and oxidative capacity. This results in G6P accumulation, a crucial mediator in load-induced mTOR activation. mTOR impairs cardiac power and induces ER stress leading to contractile dysfunction and hypertrophy. Rapamycin, metformin, and mechanical unloading reversed this metabolic cascade (34).

Impact

HF is a worldwide epidemic, affecting ~ 6 million people in the United States and 23 million people worldwide (36). Approximately 1 in 5 adults over the age of 40 will develop HF and survival estimates are as low as 70% at 1 year and 50% at 5 years. Furthermore, it costs the United States roughly 30 billion dollars per year. This is expected to more than double by 2030 (36). Morbidity and mortality of HF patients with concomitant PH is significantly higher (10, 11, 37–40). It is also apparent that there is continuum of risk with even borderline PH associated with higher mortality and morbidity (41). Unfortunately, once HF patients develop PH there are no therapies to reverse or even cease progression.

Thus far, treatment of PH-LHD is geared toward managing volume status with diuretics and dietary modifications, implementing goal directed medical therapy for left ventricular dysfunction, and treating comorbidities such as systemic hypertension. Despite medical optimization with the aforementioned therapies, elevated pulmonary pressures persist, and often worsen. Traditional PAH therapies have been studied, but unfortunately with generally negative outcomes. Endothelin Receptor Antagonists (ERA) trials such as ENABLE (Endothelin Antagonist Bosentan for Lowering Cardiac Events in Heart Failure) reported no benefit in regards to mortality and in fact found increased heart failure exacerbations and heart failure hospitalizations secondary to fluid retention (42). Prostacyclins have been associated with improved cardiac output and decreased pulmonary vascular resistance but have had a

strong trend toward higher mortality (43). Phosphodiesterase-5 inhibitors in clinical trials have had disappointing results. For example, RELAX (Phosphodiesterase-5 Inhibition to Improve Clinical Status and Exercise Capacity in Diastolic Heart Failure) showed no significant improvement in clinical outcomes (44). Uncovering novel therapies to prevent the development of PH-LHD from HF or even halt the progression of PH-LHD is paramount given the growing public health concern that HF and PH-LHD poses. Given the association of insulin resistance with HF and PH-LHD, metformin is of high interest as a novel treatment.

PRECLINICAL MODELS: METFORMIN'S MECHANISTIC EVIDENCE

Mechanistically, metformin exerts its benefit on the CV system in a pleiotropic manner (Figure 1). It is hypothesized that these pleiotropic effects are due to its role in directly activating AMPK, which is a cellular energy sensor that is intricately involved in numerous metabolic processes not limited to liver and skeletal muscle glucose metabolism (45).

In skeletal muscle, metformin enhances glucose-mediated glucose uptake resulting in improved insulin sensitivity, oxidative metabolism, mitochondrial function (46). Dysregulation of AMPK and its downstream signaling network have both been implicated in metabolic syndrome, HF, and PH. This may be further mediated upstream via sirtuin-3 (SIRT3), a mitochondrial deacetylase. When downregulated, SIRT3 contributes to the development of insulin resistance, and eventually diabetes. Furthermore, SIRT3-AMPK downregulation in pulmonary

artery smooth muscle cells is associated with the development of PH in leptin deficient, obese, diabetic rats with metabolic syndrome, and was reversed with metformin (47). Metformin restored SIRT 3 in skeletal muscles, improved insulin sensitivity, and was associated with decreased pulmonary pressures (47).

Metformin upregulates the SIRT3-AMPK-Glut4 pathway. Glucose transporter type 4 (Glut 4), an insulin-regulated glucose transporter in adipose tissue and striated muscle, improves insulin action and glucose uptake. Myokines, such as fibroblast growth factor 21 (FGF21) which regulates simple sugar intake, may be a critical link in crosstalk between skeletal muscle metabolism and pulmonary vascular smooth muscle cells, further activating AMPK, and preventing pulmonary vascular remodeling. Additional mechanisms of action of metformin on vascular remodeling include AMPK-mediated inhibition of estrogen and aromatase synthesis, inhibition of leptin secretion, and downstream STAT3 activation—all crucial mediators of group 2 PH (48).

Additionally, metformin augments weight loss, directly treating the adverse effects of obesity (19). Over 80% of HF patients in clinical trials and epidemiologic cohorts are overweight or obese. Mechanistically, obesity is hypothesized to have pleiotropic effects on HF and PH-LHD as a result of abnormal endocrine, cellular, and cardiometabolic signaling. Important mediators of this include adipokines including leptin and adiponectin. Leptin directly affects the sympathetic nervous system, aldosterone receptor signaling, janus kinase (JAK) signal transducer, STAT proteins, and p38 mitogen-activated protein kinase, all of which have been implicated in deleterious cardiac remodeling, impaired calcium handling, and impaired relaxation. Adiponectin decreases with obesity and is linked

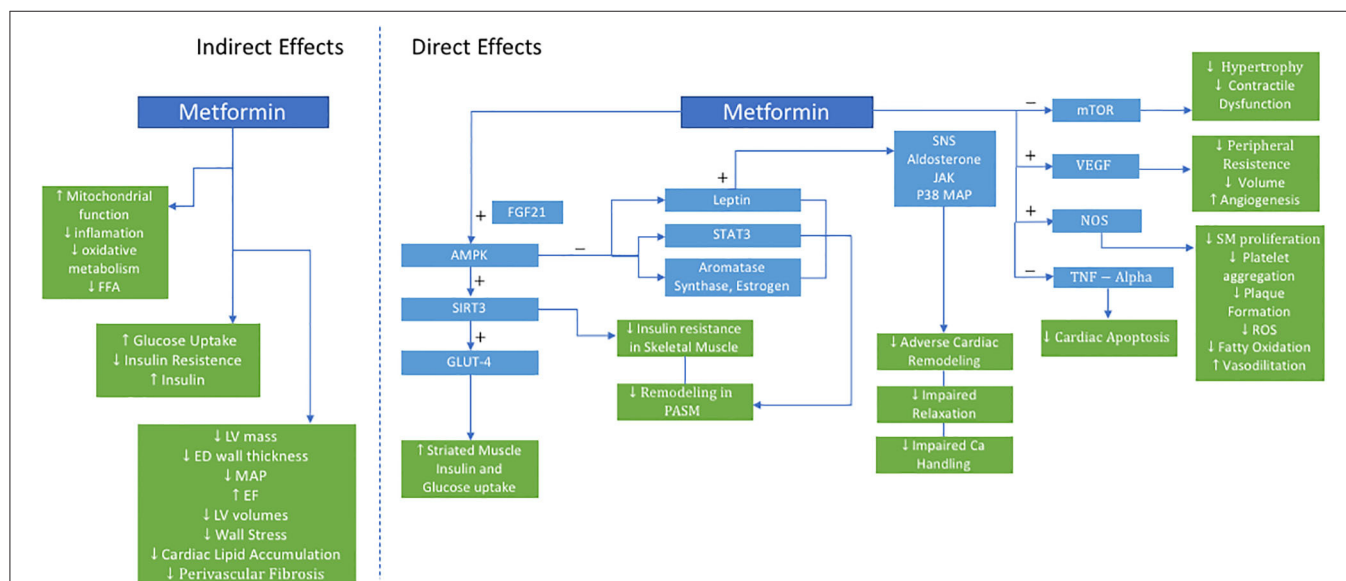


FIGURE 1 | Proposed Actions of Metformin in PH-HFpEF. AMPK indicates 5'-AMP-activated protein kinase; Ca, calcium; ED, end-diastolic; EF, ejection fraction; FGF21, fibroblast growth factor 21; FFA, free fatty acid; LV, left ventricle; GLUT4, glucose transporter type 4; JAK, janus kinase; LV, left ventricle; MAP, mean arterial pressure; mTOR, mammalian target of rapamycin; NOS, nitric oxide synthase; P38 MAP, P38 mitogen-activated protein; PASM, pulmonary artery smooth muscle; ROS, reactive oxygen species; SIRT3, sirtuin-3; SM, smooth muscle; SNS, sympathetic nervous system; STAT3, signal transducer and activator of transcription 3; TNF, tumor necrosis factor; VEGF, vascular endothelial growth factor.

to increasing inflammation, oxidative stress, as well as adverse cardiac remodeling (49). While the effect of obesity on HF has been previously described, obesity's impact on the pulmonary vasculature in the PH-LHD phenotype is still unknown.

Metformin has also been shown to have a beneficial role in chronic systemic hypertension preventing the development and progression of hypertensive LHD. Spontaneously hypertensive rats were noted to have significant myocardial metabolic changes even prior to developing cardiac dysfunction. Metformin treatment in this model lowered mean arterial pressure, lowered glucose uptake, improved ejection fraction, and decreased left ventricular mass and end-diastolic wall thickness. Furthermore, free fatty acid levels and mTOR activity were reduced as well (50). In another model, spontaneous hypertensive, insulin-resistant rats (SHHF) treated with metformin had decreased LV volumes, wall stress, cardiac lipid accumulation, and perivascular fibrosis—all markers of LV remodeling. Additional improvements were seen in indices systolic and diastolic ventricular function. Metformin activated AMPK, vascular endothelial growth factor, and nitric oxide synthase, while reducing myocyte apoptosis and tumor necrosis factor- α expression—all have which been implicated in HF pathogenesis. These metabolic, molecular, and structural changes translated to functional cardiomyocyte improvements and prevention of HF (51).

CLINICAL TRIALS

Will preclinical, observational, and *post-hoc* data translate into distinct measurable benefit for HF or PH-LHD? This has yet to be determined, but there is certainly reasonable evidence to support clinical trials (52). In a compelling pilot study recently completed, Mohan et al. (53) assessed the therapeutic benefit of metformin on LV hypertrophy in 68 patients with coronary artery disease without diabetes. They showed metformin decreased LV mass, systolic blood pressure, body weight, and oxidative stress as measured by thiobarbituric acid reactive substances in patients with preserved LVEF (53). Another trial in 62 non-diabetic insulin resistant HFREF

patients found improvements in functional class and minute ventilation/carbon dioxide production (VE/VCO₂ slope) (54). We are currently pursuing a prospective phase II clinical trial evaluating the therapeutic efficacy of metformin in PH-HFpEF (Clinicaltrials.gov identifier NCT03629340) which will evaluate exercise hemodynamics, functional capacity, skeletal muscle signaling, and insulin sensitivity.

METFORMIN IN PAH AND HFPEF

Briefly, although beyond the scope of this review, the therapeutic role of metformin is also being investigated in both pulmonary arterial hypertension (PAH) and HFpEF. There is a growing body of evidence that PAH is associated with a number of systemic metabolic derangements including metabolic syndrome, insulin resistance, and glucose intolerance. Alterations in aerobic glycolysis, tricarboxylic acid metabolism, and fatty oxidation have all been implicated in the pathophysiology. Given these findings, there is at least one currently ongoing clinical trial (NCT01884051) looking at the role of metformin in PAH (55). While in HFpEF there is significant clinical heterogeneity, there is increasing evidence illustrating metformin's therapeutic benefit across the varying phenotypes. Halabi et al. (56) recently published a systematic review and meta regression analysis analyzing the role of metformin in HFpEF. Metformin reduced mortality and morbidity in HFpEF patients even after adjustment for other HF and glycemic control therapies including beta blockers, angiotensin converting enzyme inhibitors, and insulin (56).

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All authors contributed to all aspects of the manuscript.

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Expression of a Human Caveolin-1 Mutation in Mice Drives Inflammatory and Metabolic Defect-Associated Pulmonary Arterial Hypertension

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Background: In 2012, mutations in Cav1 were found to be the driving mutation in several cases of heritable pulmonary arterial hypertension (PAH). These mutations replaced the last 21 amino acids of Cav1 with a novel 22-amino-acid sequence. Because previously only Cav1 knockouts had been studied in the context of PAH, examining the *in vivo* effects of this novel mutation holds promise for new understanding of the role of Cav1 in disease etiology.

Methods: The new 22 amino acids created by the human mutation were knocked into the native mouse Cav1 locus. The mice underwent hemodynamic, energy balance, and inflammatory measurements, both at baseline and after being stressed with either a metabolic or an inflammatory challenge [low-dose lipopolysaccharide (LPS)]. To metabolically challenge the mice, they were injected with streptozotocin (STZ) and fed a high-fat diet for 12 weeks.

Results: Very little mutant protein was found *in vivo* (roughly 2% of wild-type by mass spectrometry), probably because of degradation after failure to traffic from the endoplasmic reticulum. The homozygous mutants developed a mild, low-penetrance PAH similar to that described previously in knockouts, and neither baseline nor metabolic nor inflammatory stress resulted in pressures above normal in heterozygous animals. The homozygous mutants had increased lean mass and worsened oral glucose tolerance, as previously described in knockouts. Novel findings include the preservation of Cav2 and accessory proteins in the liver and the kidney, while they are lost with homozygous Cav1 mutation in the lungs. We also found that the homozygous mutants had a significantly lower tolerance to voluntary spontaneous exercise than the wild-type mice, with the heterozygous mice at an intermediate level. The mutants also had higher circulating monocytes, with both heterozygous and homozygous animals having higher pulmonary MCP1 and MCP5 proteins. The heterozygous animals also lost weight at an LPS challenge level at which the wild-type mice continued to gain weight.

Conclusions: The Cav1 mutation identified in human patients in 2012 is molecularly similar to a knockout of Cav1. It results in not only metabolic deficiencies and mild pulmonary hypertension, as expected, but also an inflammatory phenotype and reduced spontaneous exercise.

Keywords: Cav1, pulmonary hypertension, exercise, inflammation, monocyte, metabolism

INTRODUCTION

Caveolae are flask-shaped invaginations in the plasma membrane, typically about 50–100 nm in diameter. They are abundant in endothelial cells, smooth muscle, and adipocytes but also present in many other cell types (1). They serve many cellular functions, including buffering mechanical stress, maintaining membrane integrity, and regulating endocytosis (2). They are also involved in many signal transduction pathways, although their exact role is still under investigation.

Caveolin-1 (Cav1) is an essential component of caveolae in most tissues. It is a 178-amino-acid integral membrane protein that inserts into the membrane in a hairpin structure, generally as part of a complex of 14–16 caveolin monomers (generally incorporating both caveolin 1 and 2). The caveolae assembly also involves several accessory proteins, including cavin, PACSIN2, and EHD-2 (3).

Until 2012, the disease with the clearest association with Cav1 mutations had been lipodystrophy (4), although associations between caveolae function and disease had been drawn for multiple other conditions, including cancer, muscular dystrophy, and cardiovascular disease (5, 6). There was an extensive literature on Cav1 knockout driving pulmonary hypertension in mice (7–10), but this was largely theoretical since it had not been seen in humans. Overall, though, Cav1 knockout mice are viable and fertile, and their health problems are unobvious until in advanced age.

However, in 2012, Cav1 mutations were found as the likely causative mutations in a familial pulmonary hypertension family (11), and since then, causative mutations have been found in additional families, although it remains a rare cause. Other examples include a F160X Cav1 frame shift mutation associated with both pulmonary arterial hypertension (PAH) and congenital generalized lipodystrophy that causes premature truncation of the protein (12–14). In addition, in a recent whole-exome sequencing of 2,572 cases, potentially causative CAV1 mutations were found in 10 individuals, representing seven families or isolated cases (15).

The mutation identified in that first family was a frameshift mutation near the end of the protein P158P fsX22. This replaced

the final 21 amino acids, which were known to make up a trafficking signal (16), with an entirely novel 22 amino acids. That this resulted in pulmonary hypertension in humans as heterozygous was interesting since, in mice, only homozygous knockouts developed the disease. This implied the possibility of a dominant negative effect—since caveolins assemble into oligomers, one might imagine that the mutant proteins within these structures either retain the wild-type protein in the endoplasmic reticulum (ER) or Golgi or are trafficked to the plasma membrane and form caveolae with the wild-type, resulting in some dysfunction of the caveolae. We thus set to work in building a mouse with a knock-in of the human mutation into the native mouse Cav1 locus shortly after its initial publication in 2012.

In the interim, both we and others took a closer look at how the mutation worked in cell culture models. On its own, the P158P fsX22 mutation was trapped in the endoplasmic reticulum; however, when co-expressed with wild-type Cav1, it appeared to act as a partial dominant negative, with fewer caveolae and with compromised function as membrane reservoir, in addition to potential signaling defects (17, 18). The goal of the present study was to determine whether these defects were recapitulated *in vivo* in mice and their physiological consequences.

MATERIALS AND METHODS

Reagents and Chemicals

Streptozotocin (STZ), lipopolysaccharide (LPS), and all the other fine chemicals were purchased from Sigma Aldrich (St. Louis, MO, USA). High-fat diet (HFD) was purchased from Bio-Serv (Flemington, NJ, USA). Caveolin (610060) and Caveolin 2 (610684) antibodies were purchased from BD Biosciences (San Jose, CA, USA). Cavin1 (ab48824), EHD2 (ab23935), β -actin (ab8227), and CD68 (ab125212) antibodies were obtained from Abcam (Cambridge, MA, USA), α -smooth muscle actin was from Dako (Santa Clara, CA, USA), and PACSIN2 (AP8088b) antibody was from Abcepta (San Diego, CA, USA). Proteome Profiler Mouse Cytokine Array Kit was purchased from R&D systems (Minneapolis, MN, USA).

Animals

All animal studies were reviewed and approved by the Institutional Animal Care and Use Committee at the Vanderbilt University Medical Center (approval numbers M/12/106 and M/11/207), in compliance with the National Institutes of Health guidelines. All these studies are conducted in accordance with the ARRIVE guidelines for reporting experiments involving animals (19).

Abbreviations: AUC, area under the curve; CAV1, caveolin-1; CAV2, caveolin-2; CAVIN1, caveolae-associated protein 1; ECHO, echocardiography; EDA, end diastolic area; EF, ejection fraction; EHD2, EH domain containing 2; ER, endoplasmic reticulum; HFD, high-fat diet; IPF, idiopathic pulmonary fibrosis; LPS, lipopolysaccharides (from bacteria); MCP-1, monocyte chemoattractant protein 1; MCP-5, monocyte chemoattractant protein 5; PACSIN2, protein kinase C and casein kinase substrate in neurons 2; PAH, pulmonary arterial hypertension; PH, pulmonary hypertension; RVH, right ventricular hypertrophy; RVSP, right ventricular systolic pressure; STZ, streptozotocin.

Generation of Transgenic Mice Expressing Human CAV1 Mutation

Heterozygous ($Cav1^{+/P}$) and homozygous ($Cav1^{P/P}$) animals expressing the frame shift human caveolin 1 mutation (P158P fsX22) were generated on a mixed C57/Bl6J and FVB/N background. This was accomplished with a traditional knock-in into embryonic stem cells, with the mutation inserted into the native locus of the mouse CAV1 exon 3 and with the recombination arms extending into intron 2 and the 3' genomic sequence. Selection markers were included in intron 2 but excised during the final creation of the mouse.

Study Design

Two different animal studies were performed apart from a pilot phenotyping investigation. In the phenotyping study, we had two sets of groups. Nine- to 11-weeks-old ($Cav1^{+/+}$, $n = 4$; $Cav1^{+/P}$, $n = 4$; $Cav1^{P/P}$, $n = 2$) and 28- to 31-weeks-old ($Cav1^{+/+}$, $n = 8$; $Cav1^{+/P}$, $n = 15$) animals in the first (younger) and second (older) groups were subjected to hemodynamic assessments. In the subsequent studies, the animals were either exposed to high-fat diet + streptozotocin (diet study) or lipopolysaccharide (LPS study). Unless specified, all animals throughout the studies were exposed to 12:12-h dark/light cycle with unlimited access to regular chow and water. Both sexes (male and female) were used in all studies.

a. Diet study

In this study, 25- to 58-weeks-old animals were fed either on regular chow (13.5% fat) or 60% high-fat, high-calorie diet for 4 months. This study included six groups in two subsets. Each subset had three groups each of wild-type, heterozygous, and homozygous $Cav1$ animals—one set on regular chow and the other on HFD. In the first subset (HFD), after 4 weeks of HFD exposure, the animals were fasted for 4 h and streptozotocin (STZ) was administered (150 mg/kg, i.p. route). Following 3 days of STZ administration, blood glucose was estimated to confirm diabetes (blood glucose, >250 mg/dl). A second dose (half-dose) of STZ was administered (on the 5th week) to the animals, which did not develop diabetes following a single dose of STZ. Subsequent to STZ administration, the animals were continued on HFD for another 8 weeks. Blood glucose was monitored in all animals every 4 weeks—week 0 and end of weeks 4, 8, and 12. After a week of acclimatization in individual metabolic cages (week 10), all animals underwent an extensive indirect calorimetry energy expenditure assessment for a continuous 72 h in the Vanderbilt Mouse Metabolic Phenotypic Center. On the 11th week, an oral glucose tolerance test (OGTT) was performed on all animals, followed by 4 h of fasting. Following OGTT, the animals were continued on HFD until the experiments were terminated. Transthoracic echocardiography was performed on all animals before the hemodynamic assessment. This HFD subset included three groups, $Cav1^{+/+}$ ($n = 13$), $Cav1^{+/P}$ ($n = 15$), and $Cav1^{P/P}$ ($n = 16$). All animals, $Cav1^{+/+}$ ($n = 10$), $Cav1^{+/P}$ ($n = 8$), and $Cav1^{P/P}$ ($n = 20$), in the

next subset were exposed to regular chow throughout the study and subjected to the same experimental parameters, as mentioned previously.

b. LPS study

Similar to the diet study, we had six groups of animals spread in this study between two subsets, aged between 42 and 76 weeks. In the first subset (LPS), the animals fed on high-energy chow (25.4% fat) were challenged with LPS (7.5 μ g/dose/intratracheal route) every 4 days, for a cumulative dose of 45 μ g over 3 weeks. All the animals were provided with saline and supplemental gel food and eventually HFD, if they were sick. This subset included the following groups: $Cav1^{+/+}$ ($n = 3$), $Cav1^{+/P}$ ($n = 10$), and $Cav1^{P/P}$ ($n = 4$). After 21 days of LPS exposure, the study was terminated following a hemodynamic assessment. In the second subset (vehicle), all the experimental parameters as explained above were adopted, except LPS, which was replaced with the vehicle (PBS). This subset had three groups: $Cav1^{+/+}$ ($n = 3$), $Cav1^{+/P}$ ($n = 3$), and $Cav1^{P/P}$ ($n = 2$).

Energy Expenditure Assessment by Indirect Calorimetry Method

For the energy expenditure assessment, the animals were individually placed in metabolic cages (identical to home cages with bedding) in a 12-h light/dark cycle, temperature/humidity-controlled dedicated room in the Vanderbilt Mouse Metabolic Phenotyping Core. Energy expenditure measures were obtained by indirect calorimetry (Promethion, Sable Systems, Las Vegas, NV, USA). The calorimetry system consisted of cages identical to home cages, with the bedding equipped with water bottles and food hoppers connected to load cells for food and water intake monitoring. All animals had *ad-libitum* access to either regular diet or HFD and water. The air within the cages was sampled through microperforated stainless steel sampling tubes that ensure uniform cage air sampling. Promethion utilizes a pull-mode, negative pressure system with an excurrent flow rate set at 2,000 ml/min. Water vapor was continuously measured, and its dilution effect on O_2 and CO_2 was mathematically compensated for the analysis stream (20). O_2 consumption and CO_2 production were measured for each mouse every 5 min for 30 s. Incurrent air reference values were determined every four cages. The respiratory quotient was calculated as the ratio of CO_2 production over O_2 consumption. Energy expenditure was calculated using the Weir equation: EE (kcal/h) = $60 * [0.003941 * VO_2$ (ml/min) + $0.001106 * VCO_2$ (ml/min)] (21). Ambulatory activity was determined every second with XYZ beams. Data acquisition and instrument control were coordinated by MetaScreen v2.2.18, and the raw data were processed using ExpeData v1.7.30 (Sable Systems). Finally, the fat and the lean masses were estimated by NMR on the Bruker Minispec (Bruker Biospin Corporation, Bellerica, MA, USA). This study included the following randomly chosen animals: (i) regular chow subset— $Cav1^{+/+}$ ($n = 10$), $Cav1^{+/P}$ ($n = 8$), and $Cav1^{P/P}$ ($n = 8$) and (ii) HFD subset— $Cav1^{+/+}$ ($n = 9$), $Cav1^{+/P}$ ($n = 10$), and $Cav1^{P/P}$ ($n = 8$).

Oral Glucose Tolerance Test

On the 11th week of the study, oral glucose tolerance test (OGTT) was performed on the same animals discussed in the energy balance study. For OGTT, all animals fasted for 4 h were administered with clinical-grade dextrose (per oral, 1.5 g/kg), and blood glucose was monitored at 0, 10, 20, 30, 45, 60, 90, and 120 min after the sugar load. Almost all animals in the HFD group demonstrated more than 600 mg/dl blood glucose, and therefore the OGTT area under the curve could not be calculated for this group.

Transthoracic Echocardiography and Hemodynamic Assessment

All the animals were subjected to transthoracic echocardiography on the penultimate day of the study using Vivo 770 high-resolution image systems (Visual Sonics, Toronto, Canada). Briefly, the animals were anesthetized with 2% isoflurane-oxygen mixture, and all ventricular and pulmonary dimensions and functions were recorded. Subsequently, invasive hemodynamic assessment was performed as explained previously (22). Following the hemodynamic measurement, the animals were euthanized, blood was withdrawn, a thoracotomy was performed to exsanguinate the heart and the lung, and finally, all other required organs were collected and tissue samples were processed for RNA, protein, and histology assessments. Right ventricular hypertrophy or Fulton's index [$RVH = RV / (LV + S)$] was calculated as the ratio of wet weight of the right ventricle (RV) to the left ventricle + intra-ventricular septum (LV + S). A complete blood count analysis was performed at Vanderbilt University Medical Center Pathology Core.

Western Blot and Dot Blot Analyses

After euthanization, the superior lobe of the right lung was frozen for protein work and homogenized in radioimmunoprecipitation assay buffer supplemented with protease inhibitor cocktail (Sigma Aldrich, St. Louis, MO, USA). Following the protein estimation, an equal amount (30 µg) of protein was resolved on 4–12% NuPAGE gel (Thermo Fisher Scientific, Waltham, MA, USA) and transferred electrophoretically onto a polyvinylidene fluoride membrane, and the membranes were blocked with 5% non-fat milk solution in Tris-buffered saline with 0.1% Tween 20 for 1 h. Subsequently, the membranes were probed with the following antibodies: Caveolin (1:10,000), Caveolin 2 (1:500), Cavin 1 (1:1,000), EHD2 (1:1,000), PACSIN2 (1:1,000), and β-actin (1:5,000) overnight at 4°C. The blots were developed on the following day and visualized on the ChemiDoc MP imaging system (Bio-Rad Laboratories, Hercules, CA, USA). For the dot blot experiment, 200 µg of protein was used as per the manufacturer's instruction.

Immunohistochemical Analysis

After euthanization, the left lobe of the lung was processed and paraffin-embedded, and immunohistochemical staining was performed as explained previously (23). CD45 (1:100), CD68 (1:100), and α-smooth muscle actin (1:200) antibodies were used for the assessment of macrophage and pulmonary

vessel muscularization (24), and the images were analyzed as explained previously.

Mass Spectrometry Identification of Mutant Cav1 in Mouse Tissues

Cav1 was immunoprecipitated from mouse tissues using Dynabeads® Protein G Immunoprecipitation Kit (Thermo Fisher Scientific, Waltham, MA, USA). Then, 5 µg pAb h1-97 IgG (20 µl) was cross-linked to 50 µl Dynabeads® Protein G by BS 3 (Sulfo-DSS; Thermo Scientific™ Pierce™) according to the manufacturer's suggested protocol. Tissues were lysed at 4°C with 1,000 µl CellLytic™ M buffer (Sigma-Aldrich), and the lysed cells were centrifuged for 15 min at $12,000 \times g$ at 4°C to pellet the cellular debris. The protein-containing supernatant was transferred into a new tube containing Dynabeads®-Ab complex and incubated with rotation for 10 min at room temperature. The tube was placed on a magnet, and the supernatant ("unbound") was transferred to a clean tube for further analysis. The Dynabeads®-Ab-antigen complex was washed four times, using 200 µl washing buffer for each wash. The final wash was conducted in a new clean tube. The wash solution ("wash") was collected for further analysis. The Dynabeads®-Ab-antigen complex was resuspended with 20 µl elution buffer, 10 µl premixed NuPAGE® LDS sample buffer, and NuPAGE sample reducing agent and heated for 10 min at 70°C. The supernatant was loaded onto the gel for western blot and mass spectrometry identification. Bands corresponding to caveolin-1 were excised and subjected to in-gel trypsin digestion. The resulting peptides were analyzed by a 70-min data-dependent liquid chromatography–tandem mass spectrometry (MS/MS) analysis. Briefly, the peptides were autosampled onto a 200 × 0.1-mm (Jupiter 3 µm, 300 Å) self-packed analytical column coupled directly to an LTQ (ThermoFisher) using a nanoelectrospray source and resolved using an aqueous-to-organic gradient. A series of full scans followed by five data-dependent MS/MS was collected throughout the separation. Dynamic 17 exclusion was enabled to minimize the acquisition of redundant peptide spectra. MS/MS spectra were searched *via* SEQUEST against a mouse database (UniprotKB—reference proteome set) to which the mutant caveolin sequence had been appended and that also contained a reversed version for each of the entries (25). The identifications were filtered and collated at the protein level using Scaffold (Proteome Software).

Statistics

Graph Pad Prism, version 8.4.2 (La Jolla, CA, USA) was used for the statistical analysis. One-way ANOVA data analysis followed by Bonferroni's test for multiple comparison was performed for end-point parameters. Two-way ANOVA followed by Tukey's multiple-comparison *post-hoc* analysis was performed to compare the interaction between genotype and treatment. Pearson correlation coefficient was performed to compare the correlation between variables. The values are represented as means ± SEM; $p \leq 0.05$ were considered as statistically significant.

RESULTS

Creation of the Cav1^{P158PfsX22} Mice

Although Caveolae have been studied in the context of PAH for many years, it was studied in the context of knock-out (7–10). Before 2012, CAV1 mutations had never been identified as causal in human PAH. The initial mutations identified, a P158PfsX22 mutation, was not obviously a complete loss-of-function mutation. The mutation impacted only the final 20 amino acids, replacing the existing termination of the protein with a new rather different termination of 21 amino acids.

Our goal here was to study the physiologic impact of the human CAV1 mutations as closely as possible in a mouse model. Although the mouse and the human CAV1 sequences are very well-conserved—both are 178 amino acids, with 169/178 amino acids identical (95%) and 175/178 conserved—third base redundancy means that just inserting the frameshift would have produced a different amino acid sequence in mice. We thus replaced the end of the mouse CAV1 with the mutated human sequence rather than merely inserting the frameshift (**Figure 1A**). The final amino acid sequence of our knock-in of the human mutation into the mouse is depicted in **Figure 1B**.

These Cav1^{P158PfsX22} mice were created through routine embryonic stem cell methods with homologous recombination. The resulting mice were born with normal Mendelian proportions and were apparently healthy and bred well, even as homozygous.

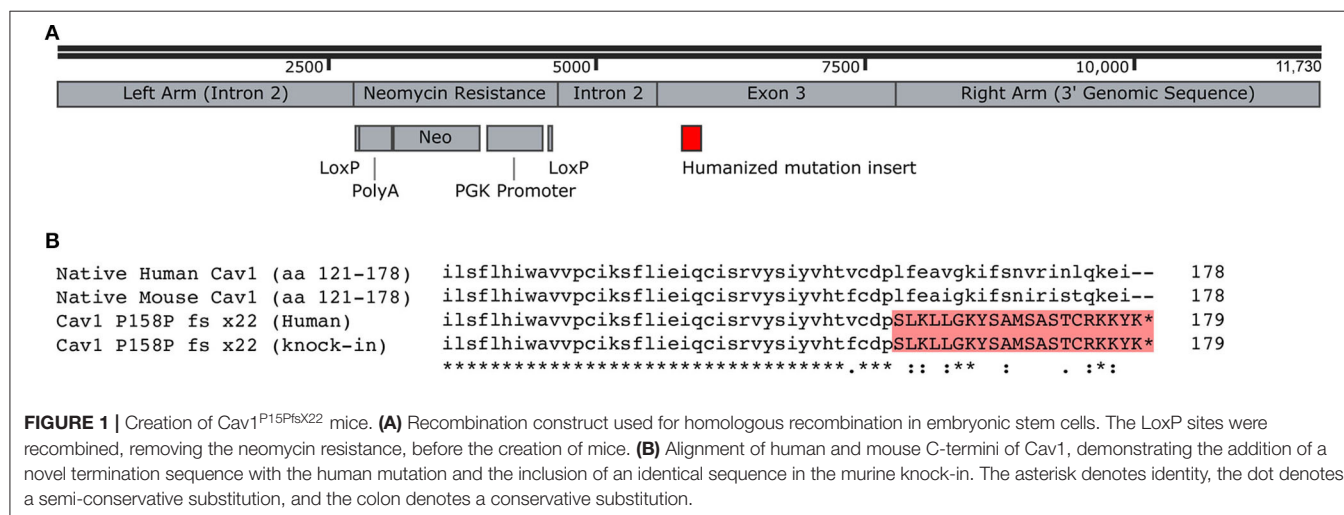
Cav1^{P158PfsX22} Protein Is Largely Degraded in Mice, With an Organ-Specific Pattern of Loss of Accessory Proteins

First, we examined the expression of both Cav1 protein in knock-in mice as well as with other proteins which are part of caveolae. These include Cav2, a homolog with which Cav1 normally hetero-oligomerizes, Cavin1, a protein that stabilizes caveolae, and EHD2 and PACSIN, which form an actin cytoskeleton-interacting complex as part of caveolae (26).

By quantitative RT-PCR using primers that spanned exons 2 and 3, we found that the spliced RNA expression of the knock-in was comparable to that of the wild-type, as expected (not shown). However, in the lungs, we found that the overall Cav1 levels were reduced by about 20% in heterozygous animals and undetectable in homozygous ones (**Figure 2A**). The levels of Cav2 were decreased at approximately the same extent as Cav1, in accord with prior literature indicating that the expression of Cav1 is required for the stability of Cav2 (27, 28). We also saw a decrease in accessory proteins EHD2 and Cavin1, but not PACSIN2, which increased as Cav1 decreased.

Since the native and the mutant proteins are of the same size, and the antibody recognizes both (there is no antibody to the mutant section of the protein, and we did not tag it in an attempt to preserve the function), we could not readily determine how much of this was from the native and how much was from the mutant protein. Although the homozygous animal had no visible Cav1 band by western blot, this is presumably because it is retained in the ER and degraded, as has been previously reported (17, 18). However, at least when overexpressed, it has been reported that native Cav1, by complexing with a mutant, can assist with the transport to caveolae, which implies that perhaps the mutant might be stabilized in the heterozygous mice. We thus assessed the levels of mutant compared to wild-type protein by using mass spectrometry. We assessed the levels of four different peptides common between the mutant and the wild-type versions of Cav1 and three different peptides specific to the mutant version. We found that, even in heterozygous animals, the expression of the mutant peptides was very low—on average only 1.4% of that of the wild-type Cav1 (**Figure 2B**).

The pattern in other organs was distinctive—in the heart, most proteins were similar to the lung, but there was no increase in PACSIN2 levels in the homozygous animals (**Figure 2C**). In the liver, there was no decrease in Cav2 or EHD2, but Cavin1 was still decreased and PACSIN2 appeared to trend down (**Figure 2D**). The kidney was similar to the liver, except that Cav2 decreased moderately in the homozygous animals (**Figure 2E**). This preservation of Cav2 in some tissues when Cav1 is missing



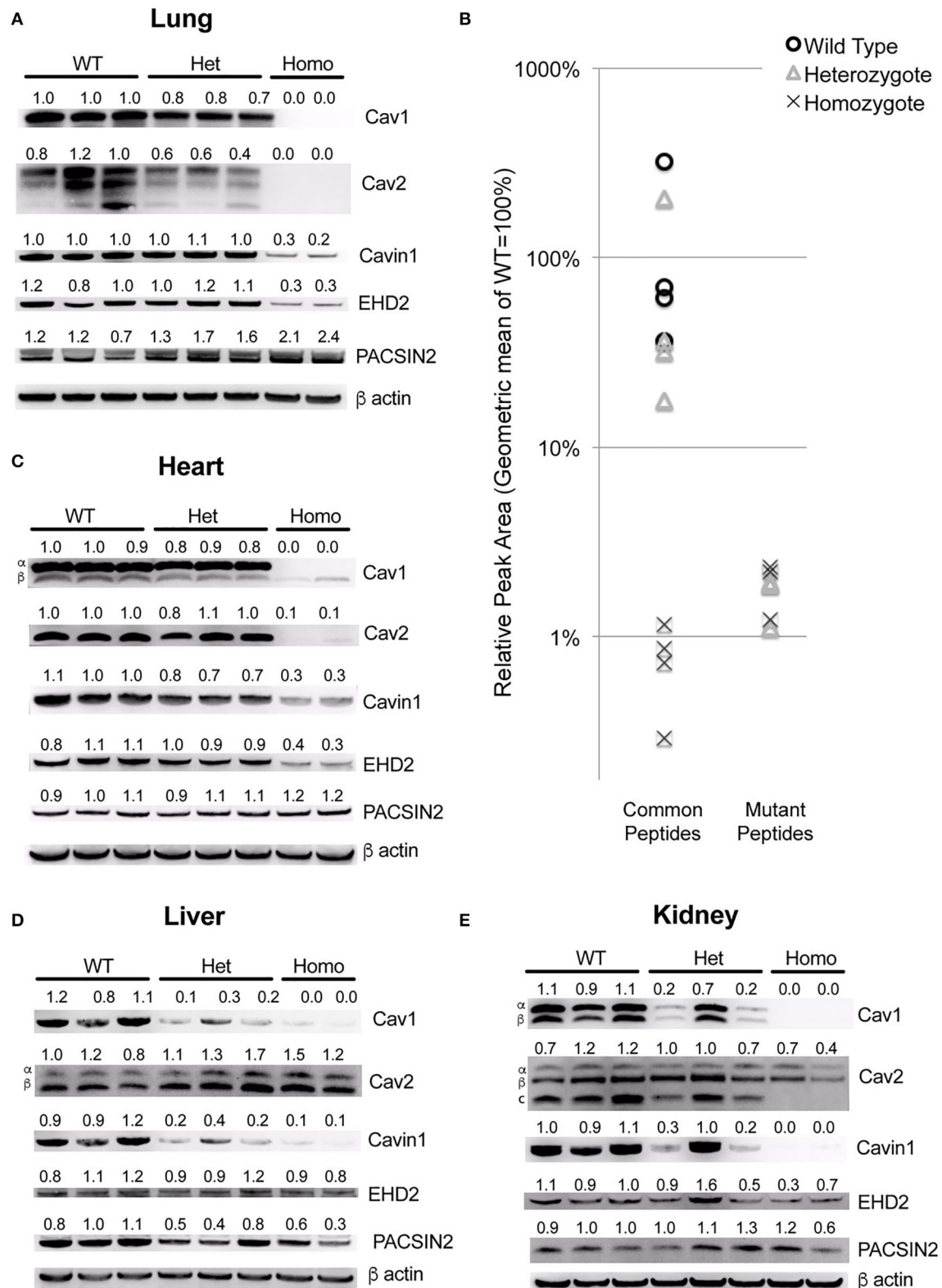
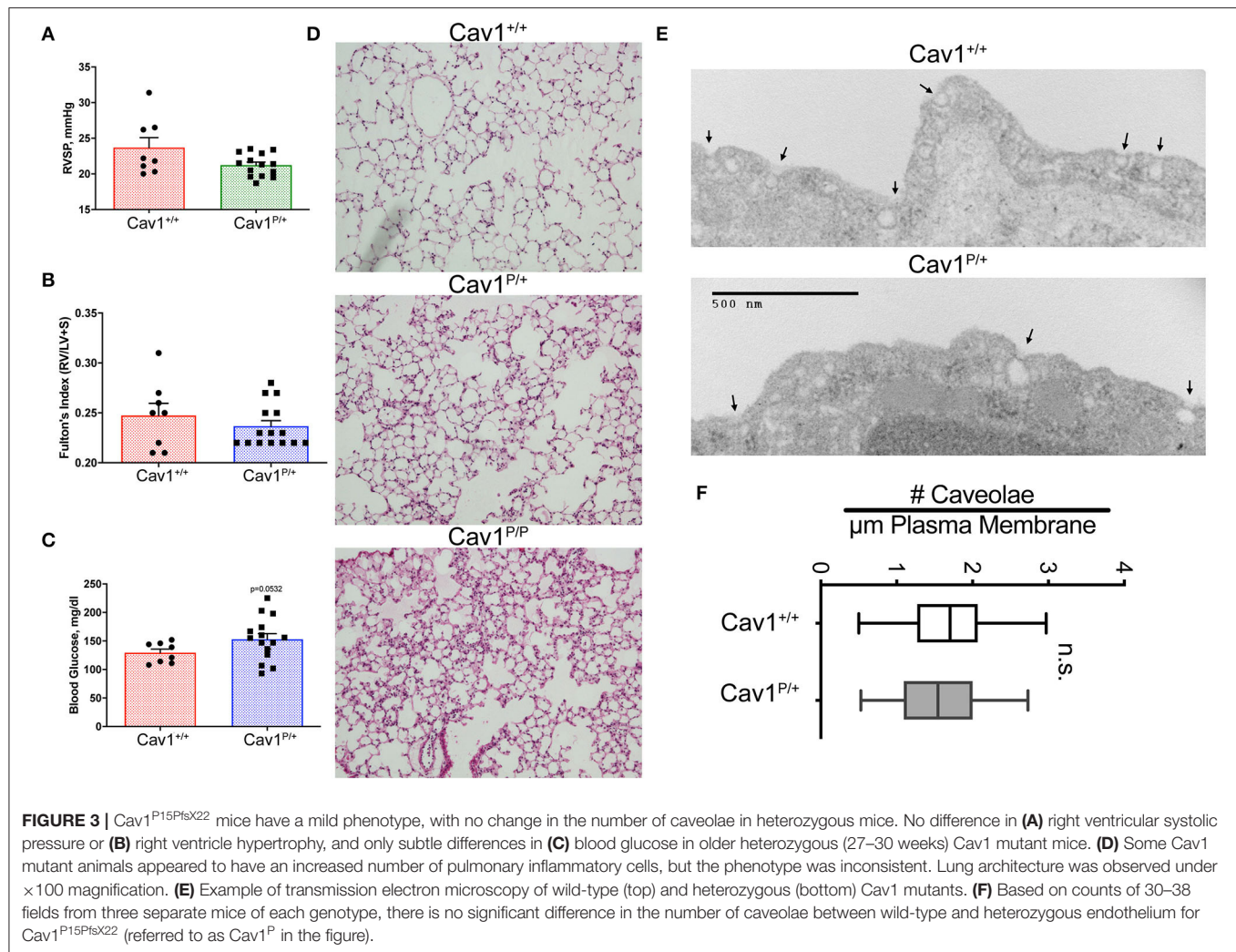


FIGURE 2 | The Cav1^{P15PfsX22} protein is largely degraded in mice, with an organ-specific pattern of loss of accessory proteins. **(A)** Western blots for the indicated proteins from whole mouse lung from wild-type, heterozygous and homozygous mice for the Cav1^{P15PfsX22} mutation—the numbers indicate densitometry normalized to wild type and to beta-actin. **(B)** The mass spectrometry quantification results of proteins from wild-type, heterozygous, and homozygous mice suggest that very little mutant protein is retained even in heterozygous mice. **(C)** Western blots for the indicated proteins from whole mouse heart. **(D)** Western blots for the indicated proteins from whole mouse liver. **(E)** Western blots for the indicated proteins from whole mouse kidney.



contradicts some literature but is supported by others, which have found that Cav2 can form homo-oligomers (29). Broadly, these data suggest that, in all circumstances, the mutant Cav1 is rapidly degraded, and the effect on accessory proteins is very organ specific.

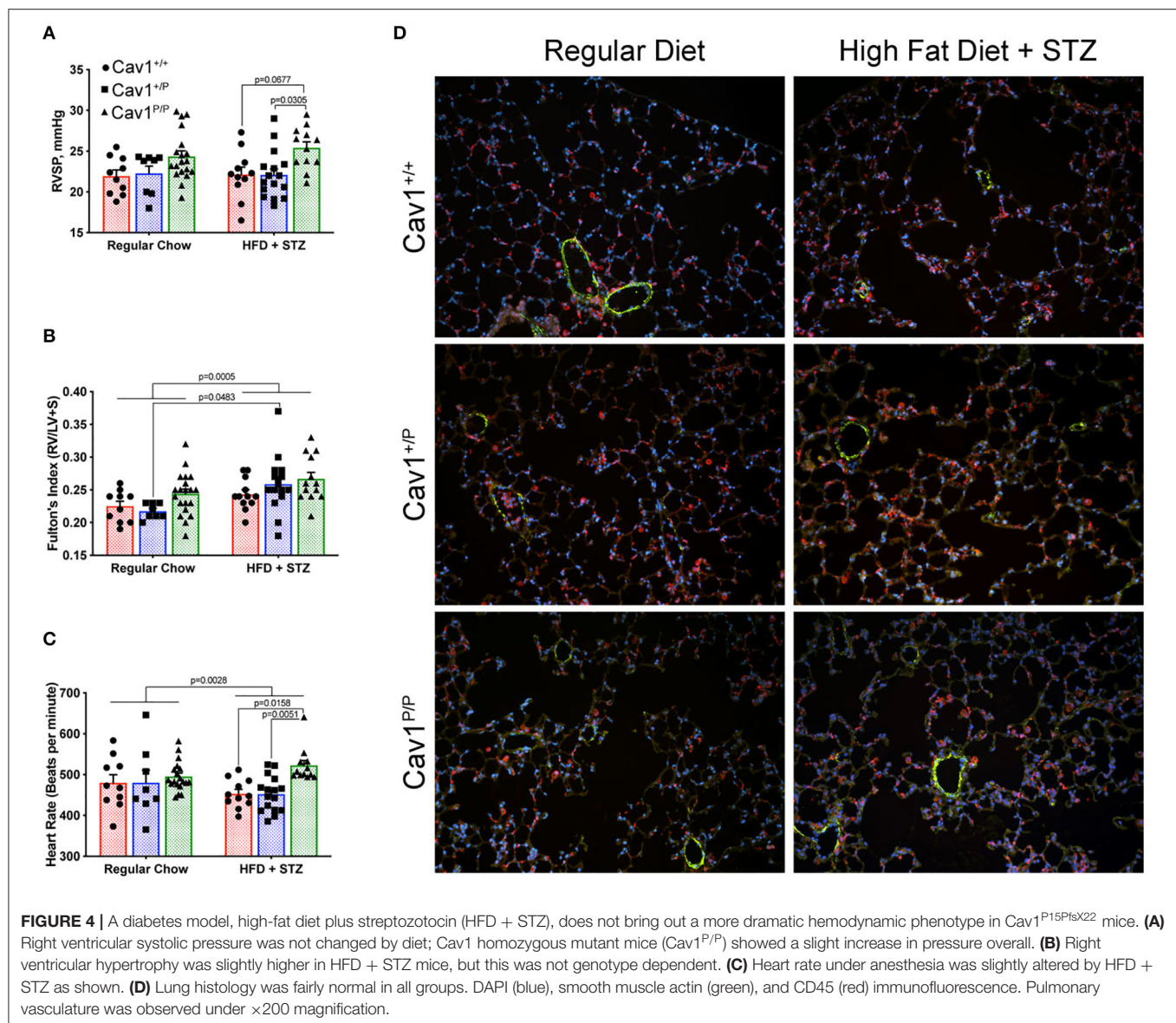
Initial Phenotyping of Cav1^{P158Pfsx22} Mice Showed a Moderate Phenotype Consistent With Prior Reports on Cav1 Knockout Mice

As an initial examination of the phenotype in these mice, we performed hemodynamic and histologic phenotyping on 27- to 30-weeks-old heterozygous Cav1 mutant and littermate control animals. The heterozygous were our focus because the heterozygous mutation results in pulmonary arterial hypertension in human patients. We found no significant difference in right ventricular systolic pressure (RVSP) (Figure 3A) or right ventricular hypertrophy (Figure 3B) and only a slight trend toward increased blood glucose (Figure 3C, $p = 0.0532$, mean 153 vs. 129 mg/dl). The results in a smaller experiment with younger animals (8- to 10-weeks-old mice)

which included a few homozygous animals showed a slight increase in RVSP in homozygous of about 30 mmHg, consistent with previous reports (Supplemental Figure 1A), but neither Fulton index (right ventricular hypertrophy) nor blood glucose changed in the mutant mice (Supplemental Figures 1B,C). By histology, the Cav1 mutant animals (Cav1^{P158Pfsx22}, hereafter referred to as Cav1^P), appeared to have increased inflammatory cell burden in the lungs (Figure 3D, Supplemental Figure 2 for higher-resolution pictures), but this pattern was inconsistent and not apparent in the older mice initially. We performed a transmission electron micrograph of lung sections and found no difference in the number of caveolae between wild-type and heterozygous endothelium (Figures 3E,F).

Stressing Cav1^{P158Pfsx22} Mice With a Diabetes Model Results in Only Subtle Changes in Hemodynamic Phenotype

Cav1 mutation in humans is more commonly associated with lipodystrophy, and Cav1 knockout in mice has previously been shown to cause broader metabolic dysfunctions (30), in addition



to a failure to gain weight under a high-fat diet (31). We thus hypothesized that the pulmonary phenotype might be more strongly brought out by metabolically stressing the mice. We used a high-fat diet with streptozotocin (STZ) to metabolically stress the Cav1^{P15Pfsx22} mice. STZ is a naturally occurring alkylating agent which is extremely toxic to insulin-producing beta cells; two doses of STZ should thus result in insulin deficiency, whereas the high-fat diet results in insulin resistance.

The inclusion of high-fat diet plus streptozotocin (HFD + STZ) made a little difference to the hemodynamics. Although the mice homozygous for mutation once again had slightly higher pressures, the increase was lower than what we had previously seen in younger mice; HFD + STZ did not cause meaningful changes in the pressures of any group (Figure 4A). HFD + STZ slightly increased RV mass across groups—there was no genotype-specific effect (Figure 4B). There were small

but statistically different changes in heart rate under anesthesia, with the control and the heterozygous mice moving downwards slightly and the homozygous mice moving upwards slightly (Figure 4C), but the changes probably are not large enough to be meaningful. The lung histology across groups was fairly normal (Figure 4D)—these pressures are not high enough to drive muscularization (see Supplemental Figure 3 for detailed counts of muscularized vessels, which did not change). We also observed a slight increase in CD45⁺ inflammatory cells in the homozygous group, but it was too patchy within and between mice to be statistically significant (not shown).

We also measured a large number of cardiac indices—left ventricular fractional shortening, left ventricular ejection fraction, left ventricular cardiac output, cardiac index, pulmonary artery area, right ventricular cardiac output, and right ventricle internal dimension during diastole. There were no meaningful

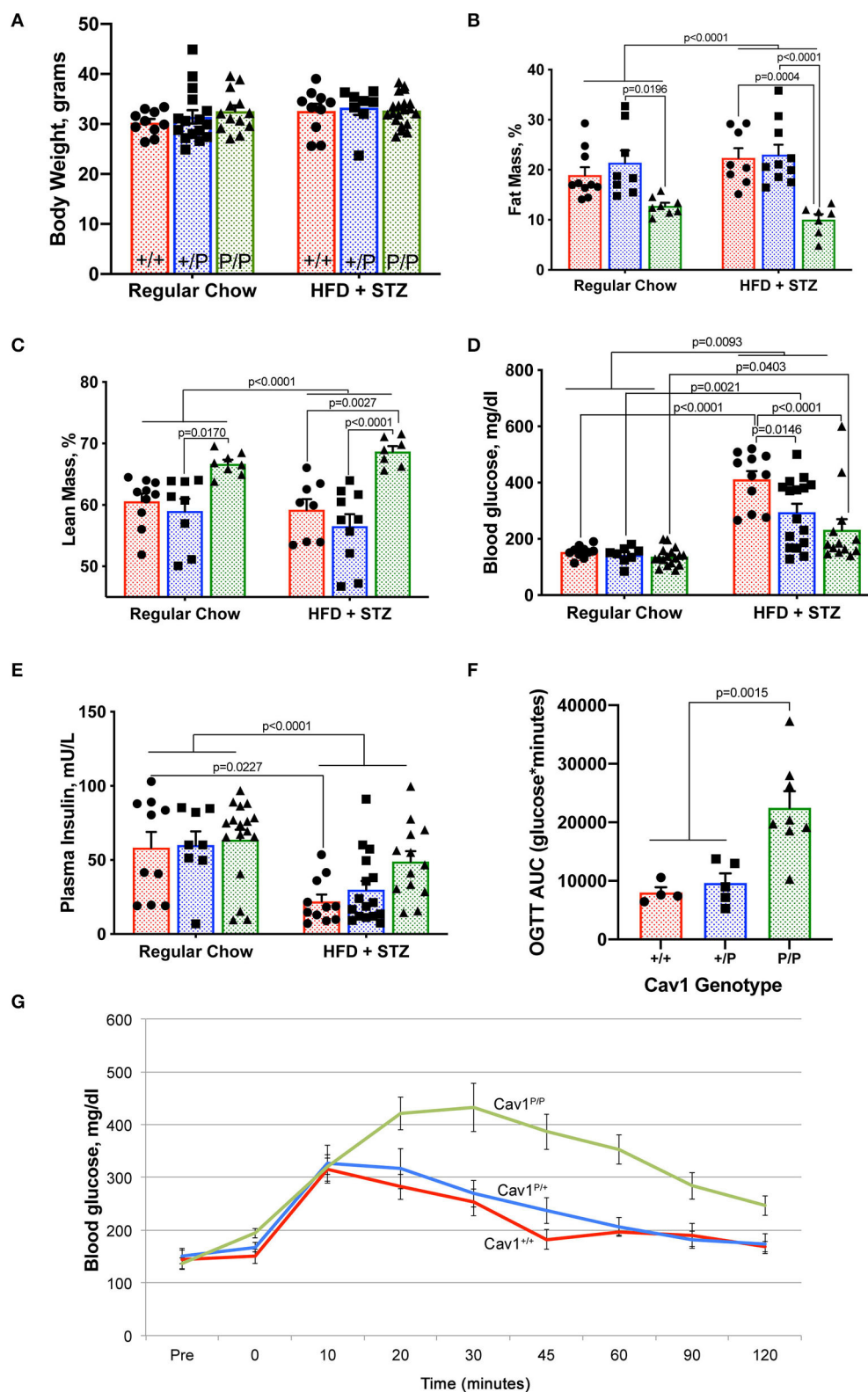


FIGURE 5 | Cav1^{P158Pfsx22} homozygous mice exhibit altered body composition and glucose tolerance. For all figure parts, red shading/circles indicate wild-type mice, blue shading/squares indicate heterozygous mice, and green shading/triangle indicates homozygous mice. Each symbol is the measurement from one animal, with (Continued)

FIGURE 5 | the bar showing mean and SEM, except for (G). Statistics were performed by ANOVA with *post-hoc* tests. (A) Body weight is not impacted by genotype or diet. (B) Homozygous mice have a lower percentage of body fat. (C) Homozygous mice have a higher percentage lean mass. (D) Although blood glucose is increased significantly in all groups with high-fat diet plus streptozotocin (HFD + STZ), it is increased to a lower extent in homozygous Cav1^{P158Pfsx22} mice. (E) Insulin is reduced in HFD + STZ mice but to a lesser extent in Cav1 homozygous mice. (F) Although they can produce more insulin, homozygous mice take a much longer time to clear oral glucose. OGTT, oral glucose tolerance test; AUC, area under the curve. These are regular chow-fed mice (HFD + STZ are off-scale for OGTT). (G) Average blood glucose at each time point during the oral glucose tolerance test by genotype. The error bars are standard error of the mean.

differences between groups in any of these measurements (not shown).

Cav1^{P158Pfsx22} Homozygous Mice Have Altered Body Composition and Glucose Tolerance Compared to Wild-Type or Heterozygous Mice

Mice on either regular diet or high-fat diet plus streptozotocin were subjected to metabolic phenotyping. Although mice on high-fat diet alone tend to have an increased body weight, HFD + STZ did not result in significant differences in body mass, either by diet or by genotype (Figure 5A). However, mice on HFD + STZ had a slightly higher body fat on average, except for Cav1^{P/P} mice, which had a significantly lower proportion of body fat and a higher proportion of lean mass (Figures 5B,C) at the same overall weight. Caveolae are abundant in adipose tissues, and adipose tissues in Cav1 knockout mice lack perilipin, which is necessary to stabilize adipocyte oil droplets (32). In contrast, caveolae in the skeletal muscle is produced by Cav3 (28), whose action is independent of Cav1 and thus likely still functional in these mice. These findings are thus consistent with the literature.

By measurements of blood glucose (Figure 5D) and plasma insulin (Figure 5E), we found that the Cav1^{P/P} mice retained greater insulin production than the other genotypes. It is possible that streptozotocin is reliant on caveolae for its function. The heterozygous mice appeared to have an intermediate phenotype. Cav1^{P/P} mice had a significantly delayed glucose clearance, with the area under the curve of the oral glucose tolerance test being nearly three times as high in homozygous as in heterozygous or wild-type mice (Figures 5F,G). The data shown are from regular chow-fed animals; OGTT was not possible in HFD + STZ mice because their glucose readings were higher than the measurable level throughout the first 90 min of the OGTT test, regardless of genotype. This lower insulin sensitivity is consistent with earlier studies on caveolin 1, which directly regulates insulin receptors (33, 34).

Cav1^{P158Pfsx22} Mice Have Much Lower Waking Locomotion but Are Similar to Wild-Type in Energy Balance Measurements

We measured a variety of metabolic parameters in heterozygous and homozygous Cav1^{P158Pfsx22} mice, both during the night (waking) and during the day (sleeping). These include food uptake, locomotion, energy expenditure, O₂ consumption, CO₂ production, and respiratory coefficient (Figure 6). For most of these, there was a diet-specific effect but no genotype-specific effect. The exception to this was locomotion—homozygous

Cav1^{P158Pfsx22} mice move less than half as much on average as wild-type mice during their waking hours (Figure 6D). The heterozygous mice were intermediate—this is one of only a small number of parameters in which the heterozygous mice show any difference compared to the wild-type mice. This is consistent with previously published results, in which Cav1 knockout mice had reduced endurance in a forced swim test (27); however, the current data indicate that this reduced physical ability is true of normal activity as well.

The respiratory coefficient, the ratio of CO₂ exhaled to O₂ consumed, is indicative of the preferred metabolic substrate, with pure carbohydrate metabolism producing a number of “1” and pure fat metabolism producing a number of “0.7.” That similarity of respiratory coefficient between homozygous and wild-type animals (Figures 6K,L) is moderately unexpected—cav1 depletion might be expected to interfere only with certain metabolic pathways. In shifting to a high-fat diet, though, it appears that Cav1^{P158Pfsx22} mice shift to a higher proportion of fat burning in the same proportion as wild-type mice.

Mice Homozygous for the Cav1^{P158Pfsx22} Mutation Have Elevated Circulating Monocytes and Elevated Monocyte Chemoattractant Proteins in the Lungs

We performed complete blood counts on mice from the dietary study (Figure 7). We found that circulating monocytes were increased in homozygous Cav1^{P158Pfsx22} mice overall, but without a clear effect of diet (Figure 7A). In contrast, there were no significant changes in lymphocytes, overall white blood count, or neutrophils (Figures 7B–D). We tested the cytokine levels in plasma and the lung by dot blot array. Although we found no significant differences in cytokines in plasma (not shown), we found increases in monocyte chemoattractant proteins 1 and 5 (MCP1/CCL2, MCP5/CCL12) in the lung (Figure 7E). There were otherwise few differences between groups (Figure 7F). There are a few previous publications linking Cav1 to monocyte attraction or function (35, 36), but this is not typically thought of as a primary role for caveolae.

These data, combined with the trend toward increased CD45⁺ cells that we saw in previous figures (Figure 4D), suggest that Cav1^{P158Pfsx22} mutation in homozygous mice is associated with moderate infiltration of inflammatory cells in the lung, but more specifically the circulating monocytes. CD68 staining for monocytes/macrophages found a trend toward an increase in the lungs of Cav1 mutant mice, but the level of variability was too high at the current *n* for it to be significant (Supplemental Figures 4A,B). In particular, there

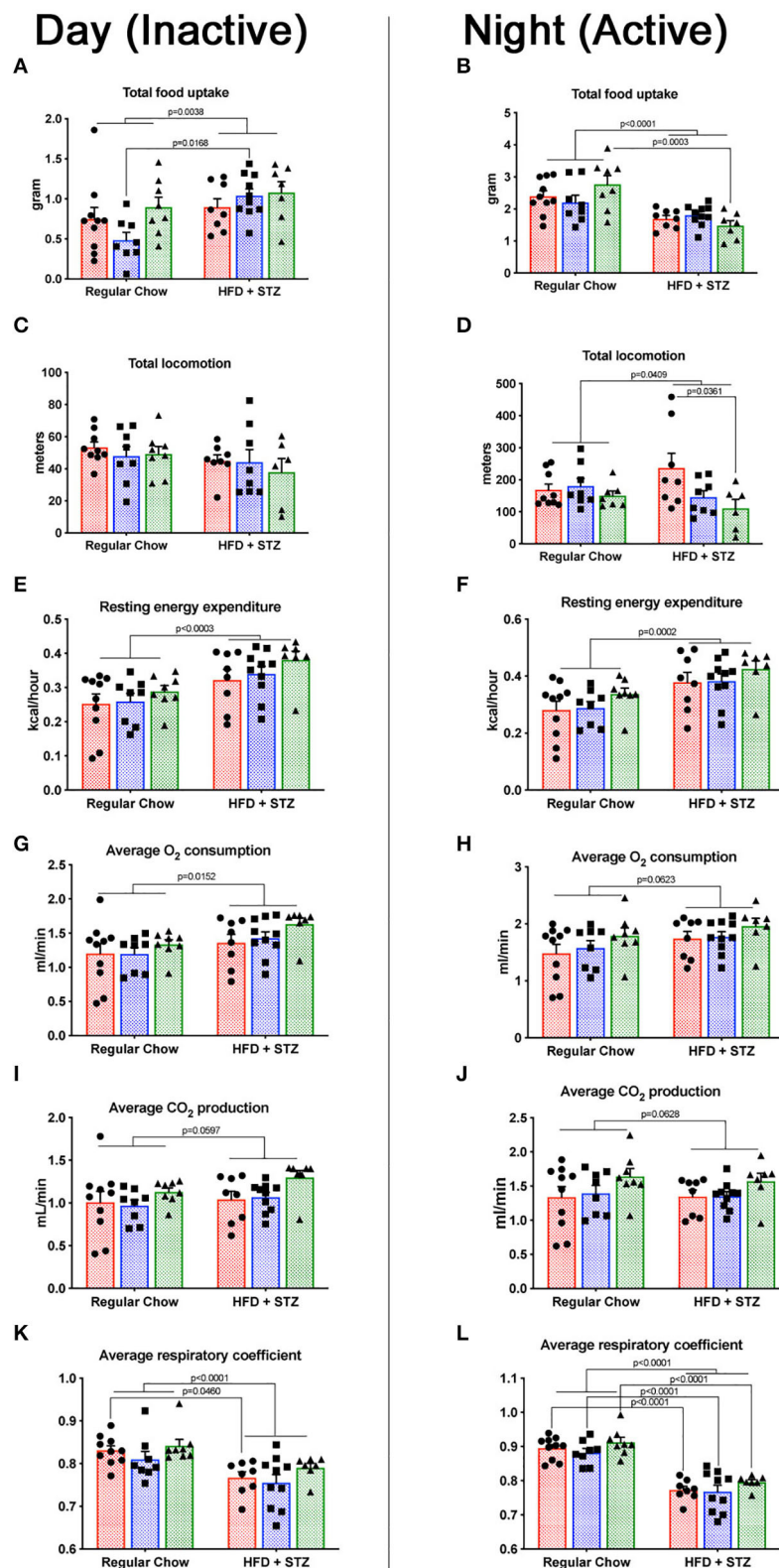


FIGURE 6 | *Cav1^{P158Pfsx22}* mice have lower waking locomotion but are similar to wild-type mice in energy balance measurements. For all figure parts, red shading/circles indicate wild-type mice, blue shading/squares indicate heterozygous mice, and green shading/triangles indicate homozygous mice. Each symbol is (Continued)

FIGURE 6 | the measurement from one animal, with the bar showing the mean and SEM. The left half of the figure (**A,C,E,G,I,K**) are measurements made during the day, when mice were inactive; the right half of the figure (**B,D,F,H,J,L**) are measurements made during the night, when mice were active. (**A,B**) Food consumed during the day and the night, respectively. (**C,D**) Total locomotion (this includes climbing as well as horizontal locomotion but is nearly identical to horizontal locomotion alone. (**E,F**) Resting energy expenditure is significantly higher with high-fat diet plus streptozotocin (HFD + STZ) in both day and night. (**G,H**) The oxygen consumption trends were higher. (**I,J**) CO₂ production. (**K,L**) The respiratory coefficient is significantly lower in HFD + STZ, as expected.

were frequently large numbers immediately surrounding large vessels (**Supplemental Figure 4C**).

Heterozygous Cav1^{P158Pfsx22} Mice Do Not Appear to Have a Hemodynamic Susceptibility to LPS but May Have a Susceptibility to Body Weight

The trends in histology shown in **Figures 3, 4**, combined with the mild suggestion of increased circulating monocytes and increased lung monocyte chemoattractant proteins in **Figure 7**, made us wonder if the Cav1^{P158Pfsx22} mice would be particularly susceptible to an inflammatory insult. We thus performed a small study using low-dose intratracheal installation of LPS, focusing on heterozygous animals, since human patients with this mutation are heterozygous.

We utilized a low dose of LPS with repeated dosing, which we had previously determined in other models to have a little effect on wild-type mice but which brought out a phenotype in susceptible animals (37). It did not, however, appear to have an impact on RVSP in either heterozygous or homozygous mice (**Figure 8A**—compare to **Figure 4A**). Fulton index is a little higher in all LPS-treated animals, but this did not differ by genotype (**Figure 8B**). Heart rate under anesthesia was similar across groups (**Figure 8C**). The only element here in which there was an apparent susceptibility was body weight; we would have anticipated that body weight would increase after 3 weeks on breeder chow, and it generally did for WT mice. However, the weight stayed the same or decreased in LPS-treated heterozygous mice (**Figure 8D**). By ANOVA, genotype was significant at $p = 0.0446$.

DISCUSSION

This study established that the new human CAV1 termination sequence, when knocked in to the native locus in mice, results in the degradation of a large majority of mutant proteins, in both heterozygous and homozygous animals, resulting in the loss of CAV2 and accessory proteins in some, but not all, tissues (**Figure 2**). Heterozygous mutants have no change in their hemodynamic phenotype, and their number of caveolae in the pulmonary vascular endothelium is unchanged (**Figure 3**)—there appears to be a compensatory upregulation of wild-type Cav1. Furthermore, they have no particular hemodynamic susceptibility to either metabolic challenge (**Figure 4**) or inflammatory challenge (**Figure 8**). For most energy balance measurements, the heterozygous mice

were more similar to the wild-type mice rather than to the homozygous—they did not share the homozygous increase in lean mass or their poor oral glucose tolerance (**Figure 5**). In other aspects, though, the heterozygous animals did show interesting differences. They appeared to have an intermediate phenotype in blood glucose and plasma insulin between wild-type and homozygous mice (**Figures 5D,E**). This was also the first study to show that the loss of Cav1 results in reduced spontaneous locomotion; the heterozygous mice appeared to have an intermediate phenotype (**Figure 6D**). Finally, the heterozygous Cav1 mutants appeared to have an intermediate phenotype in the increase in circulating monocytes seen in homozygous animals (**Figure 7A**), and they had the same increase in monocyte chemoattractant proteins (**Figure 7E**). While the LPS challenge did not increase their right ventricular systolic pressure, it did appear to make them susceptible to weight loss (**Figure 8D**).

The substantial degradation of the mutant protein in the homozygous mice is consistent with prior findings in cell culture that this mutation causes the protein to accumulate in the endoplasmic reticulum as the result of the introduction of an ER retention signal (17). Notably, the levels of Cav1 protein detected in heterozygous animals varied across tissues; they were only modestly decreased in the lung and the liver, whereas substantially less Cav1 was detected in the liver and the kidney. This suggests that in some tissues the wild-type copy of Cav1 may assist in the translocation of the mutant copy to the plasma membrane, as has been observed in cell culture studies (17, 18). Alternatively, the compensatory upregulation of WT Cav1 may occur in a tissue- or cell type-specific manner.

The metabolic findings in this study are largely supportive of the existing literature on the effects of the loss of Cav1. Both the lower fat mass (32) and the increased lean mass (28) are consistent with prior reports in mice and with the known association with lipodystrophy in human patients (4). The poor insulin sensitivity and the oral glucose tolerance AUCs are also consistent with earlier reports (33, 34). The previous publication showing that Cav1 knockout mice had difficulty with the forced swim test (27) makes sense because of their reduced ability to rapidly internalize glucose; the present study is the first to show that this is true even for a routine activity—in homozygous and, to a lesser extent, in heterozygous animals. The mechanism is presumably the same.

The inflammatory phenotype that we observed, while mild, is particularly interesting in the context of a recent publication in which reciprocal bone marrow transplants between wild-type and Cav1^{-/-} animals suggested that the

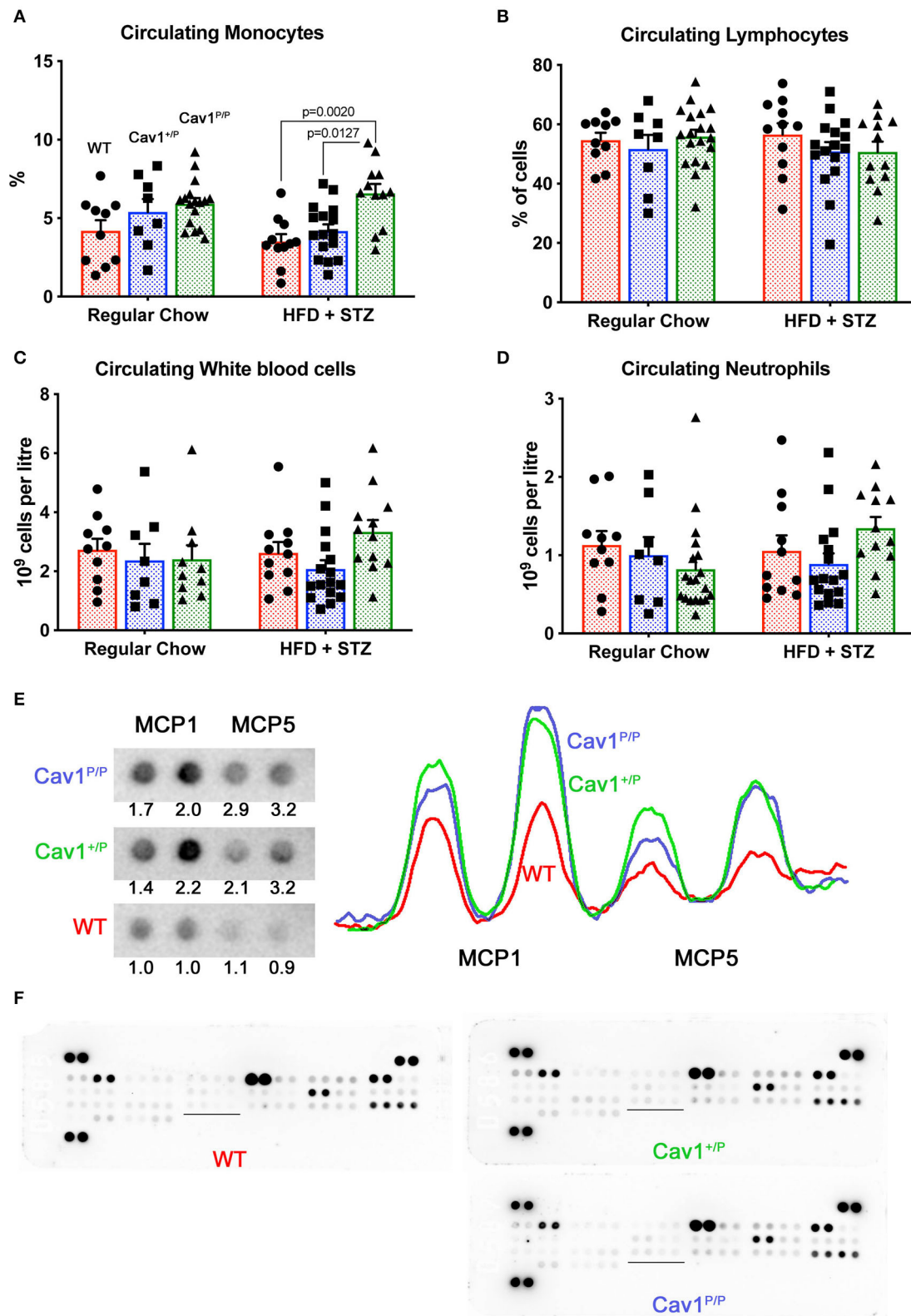


FIGURE 7 | Mice homozygous for the Cav1^{P158Pfsx22} mutation have elevated circulating monocytes and elevated monocyte chemoattractant proteins in the lungs. For figure parts (A–D), red shading/circles indicate wild-type (WT) mice, blue shading/squares indicate heterozygous mice, and green shading/triangles indicate (Continued)

FIGURE 7 | homozygous mice. Each symbol is the measurement from one animal, with the bar showing the mean and SEM. The statistics are derived by ANOVA, with *post-hoc* tests to compare individual groups. **(A)** The percentage of circulating monocytes is increased in homozygous Cav1 mutant animals, although diet has no significant effect. **(B)** The percentage of circulating lymphocytes is not altered by genotype or diet. **(C)** White blood count is not altered by genotype or diet. **(D)** Circulating neutrophils are not altered by genotype or diet. **(E)** MCP1 and MCP5 protein are increased in both heterozygous and homozygous mutant animals. The tracing is densitometry for each of the blots at the left; the numbers are areas under each curve, normalized to the average WT level of each protein. **(F)** Overall cytokine blots on lung proteins from WT, heterozygous, and homozygous mutants have few differences aside from MCP1 and MCP5 (the underlined section is blown up in **(E)** above).

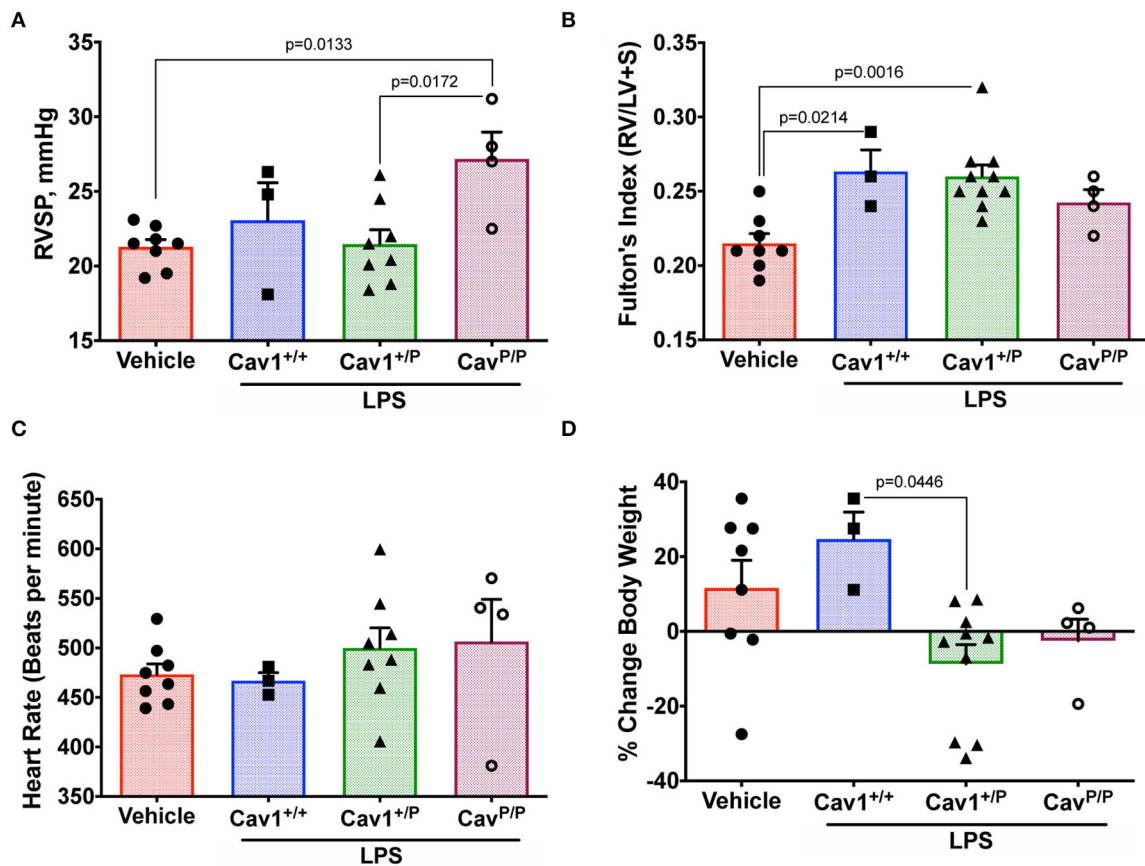


FIGURE 8 | Heterozygous Cav1^{P158Pfsx22} mice do not appear to have a hemodynamic susceptibility to lipopolysaccharide (LPS) but may have a susceptibility to body weight. For all figure parts, blue shading/squares indicate wild-type mice, green shading/triangles indicate heterozygous mice, and maroon shading/open circle indicate homozygous mice in the LPS group. Red shading/circles in vehicle control mice indicate a mixture of genotypes (three wild type, three heterozygous, and two homozygous mice). Each symbol is the measurement from one animal, with the bar showing the mean and SEM. The mice given a chronic low-dose of LPS through installation for 3 weeks had **(A)** no changes in right ventricular systolic pressure (compared to **Figure 4A**). **(B)** No genotype-dependent changes in right heart hypertrophy. **(C)** No differences in heart rate under anesthesia. **(D)** Heterozygous animals may be sensitive to LPS in weight gain, by ANOVA.

pulmonary hypertensive phenotype of Cav1^{-/-} was being driven by circulating cells (38). We are not the first to see an association between macrophages and Cav1. Cav1 knockouts showed increased adventitial macrophages in aortic transplants (39). The overexpression of Cav1 reduced the infiltration of monocytes/macrophages in a mouse model of pulmonary fibrosis (40). Reduced Cav1 levels in monocytes specifically lead to increased chemotaxis activity (36). Furthermore, the loss of Cav1 in psoriatic skin as well as peripheral blood mononuclear cells and CD14⁺ monocytes results in skin inflammation and an increase in monocyte–macrophage lineage

activities (36). In scleroderma patients, the decreased levels of Cav1 serve as a risk factor for the differentiation of monocytes to spindle-shaped fibrocytes (41). Potentially, these and other studies suggest that the loss of Cav1 may have a direct impact on monocyte/macrophage recruitment, function, or both.

Even the homozygous mice in the present study had less of a pulmonary hypertensive phenotype than had previously been published for Cav1 knockouts, although the level of pulmonary hypertension in other studies has also been variable, with some studies showing no increased pressure at baseline (42). These

differences are probably just strain differences—the original knockouts were created on a mixed 1295SVJ × Black Swiss background, whereas ours were on a mixed C57/Bl6J × FVB/N background. Differences in background genotype can make a large difference in the pulmonary hypertensive phenotype. The same is true, really, of the patients. Mutations that seem functionally similar in other individuals produce lipodystrophy, not pulmonary hypertension, which makes it seem likely that the reason that we did not see pulmonary hypertension in these mice is that they lacked the underlying polymorphisms associated with disease in the patients, although it could be due to differences between human and mouse anatomy or Cav1 function.

Hypoxia is a standard “second hit” for pulmonary hypertension genetic models, and Cav1 has previously been extensively associated with vasoreactivity, particularly through nitric oxide (10). However, the susceptibility of Cav1 knockout mice to hypoxia has been previously published, with results suggesting no particular pulmonary susceptibility but an increased rate of right ventricular failure (42). Because this model is likely to be molecularly similar to knockouts (Figure 2), we did not pursue susceptibility to hypoxia in this model.

Whether the inflammatory or the metabolic abnormalities seen in these mice are actually drivers of the hemodynamic phenotype is currently unknown. However, these defects are causally associated in some other models, and so overall, the present work adds to the body of literature associating insulin resistance and increased inflammation with pulmonary hypertension.

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DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The animal study was reviewed and approved by Institutional Animal Care and Use Committee at the Vanderbilt University Medical Center.

AUTHOR CONTRIBUTIONS

AK, AH, AR, CC, EA, EC, and JW conceived and designed the experiment. AC, AR, CC, CM, SG, and SS performed the experiment. AR, AK, AH, CC, and JW analyzed the data. AR and JW wrote the paper. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmed.2020.00540/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Current Understanding of Circulating Biomarkers in Pulmonary Hypertension Due to Left Heart Disease

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Pulmonary hypertension due to left heart disease (PH-LHD; Group 2), especially in the setting of heart failure with preserved ejection fraction (HFpEF), is the most frequent cause of PH. Despite its prevalence, no effective therapies for PH-LHD are available at present. This is largely due to the lack of a concise definition for hemodynamic phenotyping, existence of significant gaps in the understanding of the underlying pathology and the impact of associated comorbidities, as well as the absence of specific biomarkers that can aid in the early diagnosis and management of this challenging syndrome. Currently, B-type natriuretic peptide (BNP) and N-terminal proBNP (NT-proBNP) are guideline-recommended biomarkers for the diagnosis and prognosis of heart failure (HF) and PH. Endothelin-1 (ET-1), vascular endothelial growth factor-D (VEGF-D), and microRNA-206 have also been recently identified as new potential circulating biomarkers for patients with PH-LHD. In this review, we aim to present the current state of knowledge of circulating biomarkers that can be used to guide future research toward diagnosis, refine specific patient phenotype, and develop therapeutic approaches for PH-LHD, with a particular focus on PH-HFpEF. Potential circulating biomarkers identified in pre-clinical models of PH-LHD are also summarized here.

Keywords: pulmonary hypertension, group 2 PH, PH-HFpEF, HFpEF - heart failure with preserved ejection fraction, biomarkers

INTRODUCTION

Pulmonary hypertension due to left heart disease (PH-LHD, Group 2) is the most common cause of PH and is a growing public health problem with high morbidity and mortality (1–3). In fact, the presence of PH in patients with LHD has been associated with up to 5.6 times higher mortality compared to patients without PH (4). Left ventricular (LV) systolic dysfunction and diastolic dysfunction, left-sided valvular disease (aortic and mitral valve disease), and metabolic dysregulation are all known contributing factors that lead to increased LV filling pressure and the subsequent development of PH. Among these, PH attributed to LV diastolic dysfunction, also referred to as PH associated with heart failure with preserved ejection fraction (PH-HFpEF), is the most common form, although the reported prevalence varies from 23 to 83% due to variable definitions and diagnostic methods used to date (5–7). In addition to the already existing ambiguity in definition of this challenging syndrome, the 6th World Symposium on PH (Nice, 2018) has

recently redefined PH-LHD as consisting of a mean pulmonary artery pressure (mPAP) greater than 20 mm Hg (revised from at least 25 mm Hg) and a pulmonary artery wedge pressure (PAWP) greater than 15 mm Hg (**Figure 1**) (8–11). Although this change reflects recent reports describing increased risk of disease progression in patients with a mild elevation in mPAP (21 to 24 mm Hg), this revision may lead to further confusion among clinicians when trying to diagnose PH in patients (9, 12).

Abnormally elevated LV filling pressures in HFpEF, heart failure with reduced ejection fraction (HFrEF, systolic heart failure), and valvular disease can all lead to elevated left atrium (LA) pressure, increased LA volume, and reduced LA compliance (5, 13). Basic cellular mechanisms affecting LV and LA remodeling include myocyte hypertrophy, up-regulated myocardial brain/B-type natriuretic peptide (BNP) expression in the failing ventricular and atrial myocardium in response to pressure increases and/or volume overload, endothelial dysfunction, vascular oxidative stress, inflammation, interstitial fibrosis, metabolic abnormalities, etc. (14–18). Increased LA pressure can then back up into the pulmonary circulation, leading to increased pulmonary venous pressure, which is in turn transferred to pulmonary capillaries, causing damage to the alveolar-capillary barrier (also known as alveolar-capillary stress failure). As the disease progresses, structural and functional changes regulated by chronic elevation in capillary pressure may trigger pulmonary vasoconstriction, reduce nitric oxide (NO) bioavailability, increase endothelin-1 (ET-1) production, and promote remodeling in the pulmonary arteries and veins, with various combinations of intimal proliferation, medial hypertrophy, and adventitial thickening (13, 17, 19–22). These pulmonary vascular abnormalities may then lead to a further increase in mPAP in addition to PAWP elevation, resulting in elevated right ventricular (RV) afterload and ultimately causing right-side heart failure. Depending on the extent of progressive pulmonary vascular abnormalities (also known as the pre-capillary component) to the underlying LHD, the backward transmission of elevated filling pressure can lead to an increase in pulmonary artery pressure proportionally (1:1 ratio) or disproportionately ($> 1:1$ ratio) (17, 23–25). Patients with no significant pulmonary vasoconstriction or intrinsic pulmonary vasculopathy often exhibit proportional PH (isolated post-capillary PH, IpcPH). On the other hand, if chronically elevated LV filling pressure triggers pulmonary vasoconstriction and pathological pre-capillary remodeling to the point of having a high transpulmonary gradient (TPG, defined as mPAP–PAWP that exceeds 12 mmHg), elevated pulmonary vascular resistance (PVR, defined as TPG/cardiac output that exceeds 3 Wood units), and/or a high diastolic pulmonary gradient (DPG, defined as diastolic PAP–PAWP that equals or exceeds 7 mm Hg), patients exhibit out-of-proportion PH (combined pre-capillary and post-capillary PH, CpcPH). The severity of pre-capillary involvement can be established by the measured TPG, PVR, and/or DPG during right heart catheterization. In the past, $TPG \leq 12$ mm Hg, $PVR \leq 3$ Wood units, and/or $DPG < 7$ mm Hg suggested IpcPH, and when elevated ($TPG > 12$ mm Hg, $PVR > 3$ Wood units, and/or $DPG \geq 7$ mm Hg), suggested CpcPH (8). These features have been associated with mortality and cardiac hospitalizations

in patients with CpcPH associated with HFpEF (CpcPH–HFpEF) (26, 27). However, the new definition from the 6th World Symposium on PH only includes PVR of less than 3 Wood units for IpcPH and PVR of at least 3 Wood units for CpcPH (**Figure 1**) (9–11).

Despite its significant prevalence, along with high morbidity and mortality, there is no Food and Drug Administration (FDA)-approved treatment for PH-LHD at present. While the use of pulmonary vasodilators has been proven to be effective in the treatment of pulmonary arterial hypertension (PAH; Group 1), the use of these agents in patients with PH-LHD has been shown to be ineffective or even harmful (see Fernandez et al. and Vachiéry et al. for recent reviews in detail) (22, 28). This is at least in part due to the lack of a concise and uniform definition for hemodynamic phenotyping, poor understanding of the underlying pathology and the impact of associated comorbidities, as well as the absence of specific biomarkers that can aid in the early diagnosis and management of this heterogeneous disease. Currently, BNP and N-terminal proBNP (NT-proBNP) are guideline-recommended biomarkers for the diagnosis and prognosis of HF and PH (29–31). However, these are not specific for PH-LHD and vary significantly with age, sex, body mass index (BMI), and renal function (30, 31). Several biomarkers have also been shown to be complimentary to the established natriuretic peptides in guiding disease management and have proven to be diagnostically valuable in distinguishing between IpcPH and CpcPH. In this review, we aim to provide an updated overview of the current state of knowledge regarding circulating biomarkers that may be used to guide future research toward diagnosis, refinement of specific patient phenotypes and development of therapeutic approaches for PH-LHD, with a particular focus on PH–HFpEF. We will first focus on the well-established natriuretic peptides, then we will review known biomarkers proposed for PAH and their new roles in PH-LHD. Finally, we will highlight some of the emerging circulating biomarkers, including those identified recently in pre-clinical models of PH-LHD.

WELL-ESTABLISHED BIOMARKERS: BNP AND NT-proBNP

As LV dysfunction progresses, diastolic wall stress is the primary stimulus for myocardial BNP expression, which plays an important role in the regulation of cardiac remodeling, blood pressure, and intravascular volume (14–16). BNP is transcribed and produced primarily in the cardiomyocytes of the ventricles as prohormone proBNP, which is then cleaved into BNP (biologically active with a half-life of 20 min) and its N-terminal fragment NT-proBNP (biologically inactive with a half-life of 70 min). The precursor, BNP, and NT-proBNP are secreted directly into circulation in response to increased myocardial stretch mediated by pressure or volume overload (14–16).

BNP/NT-proBNP in HFpEF and HFrEF

Plasma levels of BNP and NT-proBNP are elevated in patients with HF and increase in proportion to the degree of LV dysfunction and the severity of symptoms of HFpEF, HFrEF, and

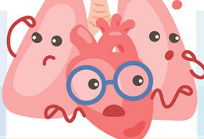
Old						New (Based on the 6th WSPH; Nice 2018)				
mPAP	PAWP	PVR	DPG	TPG		mPAP	PAWP	PVR	DPG	TPG
≥25 mm Hg	≤15 mm Hg	—	—	—	Pre-capillary PH	>20 mm Hg	≤15 mm Hg	≥3 WU	—	—
≥25 mm Hg	>15 mm Hg	≤3 WU	<7 mm Hg	≤12 mm Hg	Isolated post-capillary PH (IpcPH)	>20 mm Hg	>15 mm Hg	<3 WU	—	—
≥25 mm Hg	>15 mm Hg	>3 WU	≥7 mm Hg	>12 mm Hg	Combined post-capillary PH (CpcPH)	>20 mm Hg	>15 mm Hg	≥3 WU	—	—

FIGURE 1 | Updated hemodynamic definition and clinical classification of pulmonary hypertension (based on the 6th World Symposium on PH, Nice 2018). mPAP, mean pulmonary artery pressure; PAWP, pulmonary artery wedge pressure; TPG, transpulmonary gradient (defined as mPAP–PAWP); PVR, pulmonary vascular resistance (defined as TPG/cardiac output); DPG, diastolic pressure gradient (defined as diastolic PAP–PAWP).

valvular disease (15, 32, 33). In fact, BNP and NT-proBNP are among the first circulating biomarkers included in the current guidelines for the diagnosis and risk stratification of HF (8, 31). While definitive cutoffs of BNP and NT-proBNP are not well-established for HFpEF or HFrEF (*see* Ponikowski et al. and Pieske et al. for the most recent recommended values) (18, 30), and various studies have used different thresholds, it is generally recognized that their levels are, on average, lower in HFpEF compared to HFrEF (30, 33). This may be due to the high prevalence of obesity in HFpEF populations, as disproportionately low BNP and NT-proBNP levels have been reported in obese patients, which may be related to the mechanisms involving natriuretic peptide degradation in adipose tissue, insulin resistance, and enhanced pericardial restraint (32, 34–37). Adipocytes strongly express the natriuretic peptide receptors NPR-A and NPR-C. As BNP binds to both NPR-A and NPR-C, it has been suggested that increased adipose tissue can lead to more efficient clearance of BNP in obese patients. However, lower plasma NT-proBNP levels seen in patients with obesity-related HF cannot be explained by the same mechanism because NT-proBNP does not bind to NPR-A or NPR-C (38, 39). Additionally, Obokata et al. found that patients with obesity-related HFpEF display more concentric LV remodeling, greater epicardial fat thickness, higher PAWP, RV dilation and dysfunction, elevated right atrial (RA) pressure, and increased total heart volume accompanied by lower NT-proBNP levels compared to non-obese HFpEF patients (32). In this study, PAWP was higher for any given NT-proBNP level in patients with obesity-related HFpEF compared to non-obese HFpEF patients; however, the relationship between NT-proBNP and LV transmural pressure, defined as intracavitary pressure

(PAWP) minus the external pressure applied to the LV from pericardium and the right side of the heart (RA pressure), did not change in both obese and non-obese patients with HFpEF (32). As increased epicardial fat and higher heart volume exacerbate pericardial restraint, which may reduce wall stress as external pressure applied to the ventricle increases, these data may provide an alternative mechanism for the lower concentration of NT-proBNP observed in patients with obesity-related HFpEF (32, 37). Nevertheless, even mild elevations in NT-proBNP have been found to correlate with an increased risk of HF among individuals with obesity (40).

BNP/NT-proBNP in PH-HFpEF and PH-HFrEF

Using a BNP cutoff value of > 100 pg/ml, an analysis from the Northwestern University HFpEF Program reported that up to 70% of patients with confirmed HFpEF have elevated BNP, which is associated with significantly higher mPAP, RA pressure, and PVR, along with higher rates of HF hospitalization, cardiovascular hospitalization, and death compared to HFpEF patients with “normal” BNP (≤ 100 pg/ml) (41). Elevated BNP levels were also found to be associated with an increase in PA systolic pressure (PASP) in PH patients associated with HFpEF, HFrEF, or valvular disease (42). Significantly higher BNP levels were observed in patients with CpcPH (50% of whom have LV ejection fraction <50%) compared to that of patients with PAH (25). Similarly, CpcPH-HFpEF patients were found to have about 1.8-fold higher levels of NT-proBNP compared to IpcPH-HFpEF patients, which was associated with an increased frequency of HF hospitalizations, reduced RV-vascular coupling (as measured by the ratio between tricuspid valve annular plane systolic

excursion and PASP; TAPSE/PASP), impaired RV function, and severely depressed exercise capacity (43). Hussain et al. reported that NT-proBNP values were overall higher in HFpEF patients with PH, as compared to patients without PH. NT-proBNP levels were also higher in HFpEF patients with RV dysfunction (decreased TAPSE) despite the absence of PH. A moderate negative correlation between NT-proBNP and TAPSE/PASP ratio was found in this study cohort (44). A retrospective analysis performed by Hoepfer et al. in 108 patients diagnosed with PH-HFpEF also revealed that each 100 ng/l increase in NT-proBNP is associated with an increased risk of death (45). Guazzi et al. also demonstrated that NT-proBNP values were higher in non-survivors among PH-HFpEF and PH-HFrEF patients, as compared to survivors, although this difference was not deemed significant in the multivariate analysis of survival (46). **Table 1** summarizes recent findings of BNP/NT-proBNP in PH-LHD.

In contrast, Miller et al. reported that NT-proBNP levels were lower in systemic sclerosis patients with early PH due to LHD than that due to PAH (47). Mazurek et al. showed that NT-proBNP levels were similar in both PH-HFpEF and PAH patients (48). The reason for these discrepancies is not immediately clear. While the levels of BNP and NT-proBNP have not yet been compared within all PH-patient populations and cannot yet be used to differentiate between pre-capillary and PH-LHD, the increase in circulating levels of these biomarkers appears to correlate well with the worsening of clinical outcomes, particularly in PH-HFpEF patients. Whether or not the monitoring of either BNP or NT-proBNP would be meaningful and effective in identifying higher risk patients or in guiding specific therapy to improve outcomes in patients with PH-LHD needs to be further investigated.

KNOWN BIOMARKERS ASSOCIATED WITH LHD: ET-1

As mentioned above, chronic elevation in pulmonary capillary pressure may result in reduced pulmonary vascular compliance, structural abnormalities, and vasoconstriction in the pulmonary vasculature. Endothelial dysfunction, which leads to a reduction in NO production and an increase in ET-1 levels, is thought to be a significant contributor to these changes (13, 17, 19, 21, 22). ET-1 is the most potent endogenous vasoconstrictor known at present and has been frequently reported to be elevated in patients with HFpEF, HFrEF, and left-sided valvular disease (49–53). It is derived from prepro-ET-1, which is first proteolytically cleaved to yield a 39-amino acid intermediate Big ET-1, followed by a subsequent production of the 21-amino acid vasoactive peptide by endothelin converting enzymes (ECEs) (**Figure 2**) (54). The production and release of both Big ET-1 and ET-1 are promoted in response to increased myocardial stress, shear stress, low levels of estrogen, hyperglycemic conditions, oxidized LDL cholesterol, elevated proinflammatory cytokines, and other conditions that are commonly involved in the progression of LHD, though Big ET-1 has at least two orders of magnitude less vasoconstrictor potency than the mature ET-1 (55, 56). ET-1 acts at two different G protein-coupled receptors, ET_A and ET_B. ET_A

receptors are located predominantly in vascular smooth muscle cells and myocytes, and are known for their potent and long-lasting vasoconstrictive and proliferative responses to ET-1. ET_B activation, on the other hand, exerts vasoconstriction in smooth muscle cells but induces transient vasodilation in endothelial cells by releasing NO and prostacyclin. ET-1 levels were found to be increased in blood of HF patients (51, 52). Increased plasma levels of Big ET-1 were reported in PAH patients as well (57). Recently, Meoli et al. reported that ET-1 is higher in plasma samples collected from the wedge position of CpcPH-HFpEF patients compared to HFpEF patients with IpcPH or without PH (58). Although the sample size was small, increased wedge plasma ET-1 concentration in this study was reported to correlate strongly with PVR in patients with CpcPH-HFpEF. A slightly larger prospective cohort study performed by Chowdhury et al. showed that regardless of being associated with CpcPH or IpcPH, the wedge plasma concentration of ET-1 in PH-HFpEF patients is higher and is associated with PH, PVR, and 1-year heart failure hospitalization compared to HFpEF patients without PH (59). Similarly, Obokata et al. reported that patients with confirmed HFpEF display elevated levels of C-terminal pro-ET-1 (CT-proET-1), a stable circulating precursor of ET-1, the magnitude of which is associated with higher pulmonary artery pressure and worse pulmonary artery compliance at rest and during exercise (52). Increased plasma CT-proET-1 levels in this study also showed high diagnostic accuracy in identifying patients with abnormal pulmonary vascular reserve during exercise. Although ET-1 may play a role in the pathophysiology of PH-HFpEF, the results of using FDA-approved endothelin receptor antagonists for the treatment of PAH have been rather disappointing in the treatment of patients with PH-HFpEF. The pilot study of bosentan, a dual ET_A and ET_B receptor antagonist, showed no signs of benefit in PH-HFpEF patients (60). Moreover, this treatment may even be detrimental in the combined populations of CpcPH- and IpcPH-HFpEF patients. Bosentan demonstrated similar harmful effects during a clinical trial for the treatment of PH-HFrEF patients (61). The MELODY-1 trial, which randomized patients with CpcPH (> 75% normal ejection fraction), to a 12-week treatment with macitentan, a dual ET_A and ET_B receptor antagonist, or a placebo showed an increased incidence of side effects in the macitentan group (mainly fluid retention), without any major improvements in any of the exploratory endpoints (62). Currently, the SERENADE trial, a phase IIb clinical trial, is recruiting patients with HFpEF associated with pulmonary vascular disease or RV dysfunction to evaluate the long-term (24–52 weeks) treatment effect of macitentan (NCT03153111).

EMERGING BIOMARKERS: VEGF-D/FIGF

Beyond vasoconstrictive and proliferative responses mediated by ET-1, increased secretion of inflammatory cytokines and growth factors (e.g., transforming growth factor alpha 1, TGF- α 1; vascular endothelial growth factor, VEGF; and interleukin 1, IL1) have also been associated with structural and functional changes in response to the retrograde increase in pulmonary

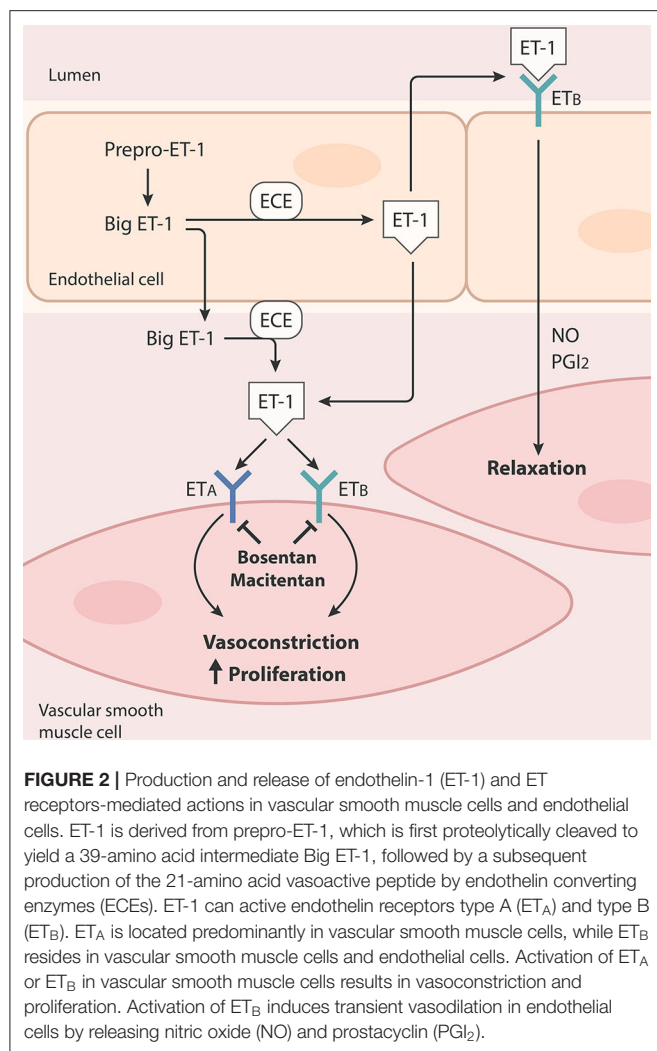
TABLE 1 | Collected studies of BNP/NT-proBNP in PH-LHD.

Authors (ref #)	Biomarker	Patients (n)	Controls (n)	Primary findings	Correlation of BNP/NT-proBNP with hemodynamics
Anjan et al. (41)	BNP	HFpEF (113), w/ BNP > 100 pg/ml, prospectively enrolled	HFpEF (46), w/ BNP ≤ 100 pg/ml	Normal BNP associated w/ lower age, female gender, obesity; Elevated BNP predicts CV hospitalization/death	RV wall thickness, RV size, LA/RA size, mPAP, RAP, PVR, ↗ Diastolic function, RV systolic function ↘
Jin et al. (42)	BNP	LHD (47), PH-LHD (35), prospectively enrolled	Healthy volunteers (36)	BNP elevated in PH-LHD patients compared to LHD patients	PASP ↗
Assad et al. (25)	BNP	CpcPH (364), lpcPH (1456), retrospective analysis	PAH (564)	BNP elevated in CpcPH and lpcPH patients compared to PAH patients	N/A
Gorter et al. (43)	NT-proBNP	lpcPH-HFpEF (46), CpcPH-HFpEF (30), prospectively enrolled	Non-PH HFpEF (21)	NT-proBNP elevated in PH-HFpEF patients compared to non-PH HFpEF patients. CpcPH-HFpEF patients have elevated NT-proBNP compared to lpcPH-HFpEF patients.	TAPSE/PASP ↘
Hussain et al. (44)	NT-proBNP	PH-HFpEF (76), retrospective analysis	Non-PH HFpEF (61)	NT-proBNP elevated in PH-HFpEF patients compared to HFpEF patients	PASP ↗ TAPSE ↘
Hoepfer et al. (45)	NT-proBNP	PH-HFpEF w/ DLCO < 45% predicted value (52), retrospective analysis	PH-HFpEF w/ DLCO ≥ 45% predicted value (56)	Elevated NT-proBNP associated with increased risk of death in univariate model	N/A
Guazzi et al. (46)	NT-proBNP	Survivors of PH-HFpEF and PH-HFrEF (246), prospectively enrolled	Nonsurvivors of PH-HFpEF and PH-HFrEF (47)	Elevated NT-proBNP not associated with increased risk of death in multivariate model	N/A
Miller et al. (47)	NT-proBNP	SSc PH-LHD (15), SSc PAH (9), retrospective analysis w/ prospective followup	SSc (19)	NT-proBNP significantly higher at diagnosis of SSc PAH compared to SSc PH-LHD	N/A
Mazurek et al. (48)	NT-proBNP	PH-HFpEF (39), prospectively enrolled	PAH (37)	No significant difference in NT-proBNP between PH-HFpEF and PAH	N/A

BNP, B-type natriuretic peptide; CpcPH, combined pre- and post-capillary pulmonary hypertension; CV, cardiovascular; DLCO, diffusion capacity of the lung for carbon monoxide; HFpEF, Heart failure with preserved ejection fraction; HFrEF, heart failure with reduced ejection fraction; lpcPH, isolated post-capillary pulmonary hypertension; LA, left atrium; LAD, left atrial diameter; LHD, left heart disease; LVEDP, left ventricular end-diastolic pressure; LVEF, left ventricular ejection fraction; mPAP, mean pulmonary arterial pressure; NT-proBNP, N-terminal fragment of pro-B-type natriuretic peptide; PAH, pulmonary arterial hypertension; PASP, pulmonary arterial systolic pressure; PH, pulmonary hypertension; PH-LHD, pulmonary hypertension associated with left heart disease; PVR, pulmonary vascular resistance; RA, right atrium; RAP, right atrial pressure; RV, right ventricle; RVSP, right ventricular systolic pressure; SSc, systemic sclerosis; TAPSE, tricuspid annular plane systolic excursion. ↗ indicates positive correlation with the corresponding biomarker; ↘ indicates an inverse correlation with the corresponding biomarker.

capillary pressure (22). Among them, the role of vascular endothelial growth factor-D (VEGF-D) has been recently evaluated in PH-LHD. VEGF-D is a secreted factor that regulates angiogenesis, lymphangiogenesis and vascular permeability (63, 64). It is synthesized and secreted as a large precursor, which is subsequently proteolytically processed at both N- and C-termini to yield the mature forms. Unprocessed VEGF-D is selective for vascular endothelial growth factor receptor 3 (VEGFR-3), which is mainly expressed in lymphatic endothelial cells, whereas the mature VEGF-D activates both VEGFR-3 and VEGFR-2,

the latter of which is found in both vascular and lymphatic endothelial cells (64). Besides modulating the growth of blood and lymphatic vessels, VEGF-D has been shown to induce cardiac fibrogenesis by stimulating myofibroblast growth, migration, and type I collagen synthesis (65). VEGF-D has also been shown to be up-regulated by mechanistic (formerly mammalian) target of rapamycin (mTOR), a master regulator of cell growth, proliferation, and survival that has been implicated in PAH, lymphangioleiomyomatosis (LAM), and cancer (66). While adenoviral delivery of a gene encoding mature VEGF-D has been



shown to improve myocardial perfusion in pigs and has now advanced through phase I/IIa studies in patients with refractory angina (67, 68), elevated circulating levels of VEGF-D have been reported in patients with HF, atrial fibrillation, ischemic stroke, PAH, chronic thromboembolic pulmonary hypertension (CTEPH) and LAM (69–73). Circulating VEGF-D levels were found to be higher in HF patients with pulmonary congestion (70). Elevated VEGF-D levels were also found to be a predictor of all-cause mortality in patients with coronary artery disease (CAD) and enabled differentiation of subjects with HF from patients with acute dyspnea (70, 74). Recently, Säleby et al. reported that VEGF-D levels were elevated in patients with PAH, CTEPH, PH-HFpEF, and PH-HFrEF compared to control subjects and HF patients without PH, with its levels in PH-HFrEF patients being significantly higher compared to all other etiologies of PH (72). Of note, plasma levels of VEGF-A were higher in patients with PAH, CTEPH, and PH-LHD compared to controls, although some controversy exists regarding the presence or absence of significant differences in VEGF-A between PH etiologies (71, 72). However, data are more consistent for

VEGF-D, showing increased levels in PH-LHD compared to PAH and CTEPH patients. Furthermore, levels of soluble fms-like tyrosine kinase 1 (sFlt-1, also known as soluble VEGFR-1 or sVEGFR-1) were significantly elevated in PH-LHD patients compared to controls, PAH, and no-PH LHD patients (71). These results are in agreement with a study by Houston et al., in which the authors showed that higher VEGF-D levels correlate with increased PAWP, reduced cardiac output, and higher BNP levels in patients with heart failure (75). A lower ejection fraction (30 ± 18 vs. $49 \pm 22\%$) was also found in subjects with higher VEGF-D levels and elevated PAWP compared to those with lower VEGF-D levels in this study cohort. While there is no data available regarding the effect of therapies targeting VEGFR3 inhibitors has been shown to improve the lumen obliteration and pulmonary pressures in Sugen/hypoxia rat models of PAH (76). Collectively, the available evidence suggests that VEGF-D could be a potential biomarker for distinguishing PH-HFrEF from other etiologies of PH, but this needs to be further evaluated in large multi-center studies.

POTENTIAL BIOMARKERS: microRNA-206

microRNAs (miRNAs) have been the subject of much excitement in molecular biology in the last few years. Composed of single-stranded non-coding RNAs that self-pair into a stem-and-loop structure, miRNAs act as post-transcriptional modifiers of their messenger RNA (mRNA), inducing their degradation and/or translational repression. While some controversy still exists as to their exact mechanism in identifying gene targets and roles in regulating diseases, miRNAs have emerged as promising biomarkers due to their relatively high stability (77). They can be detected extracellularly and have been implicated in a wide range of diseases. Several miRNA microarray profiling and quantitative PCR array studies have reported candidate miRNAs with potential utility as biomarkers in HF and PAH (see Fernandez et al., Wong et al., and Boucherat et al. for recent reviews in detail) (22, 78, 79); however, only a few miRNAs have been studied in PH-LHD. Among them, circulating levels of muscle-specific miR-204 were reported to be uniquely elevated across the pulmonary vasculature in patients with PAH, but not in patients with PH-LHD (80). On the other hand, muscle-specific miR-206 levels were found to be reduced in serum of patients with PH-HFpEF, PH-HFrEF, and valvular disease-associated PH (42). Notably, decreased miR-206 levels in this study cohort correlated with an increase in PASP, BNP, and left atrial longitudinal diameter (LAD). In addition, the authors found that the predictive value of miR-206 in detecting PH in LHD was greatly improved when combined with BNP and LAD. The exact role of miR-206 in PH-LHD is currently unknown; however, reduction of miR-206 has been shown to stabilize hypoxia-inducible factor-1 α (HIF-1 α) and to induce proliferation in cultured PASMCs (81, 82). In contrast, miR-206 levels were found to be elevated in lungs of rats with monocrotaline (MCT)-induced PAH, and cardiac-specific overexpression of miR-206 also led to cardiac hypertrophy (83, 84). While the reduction of

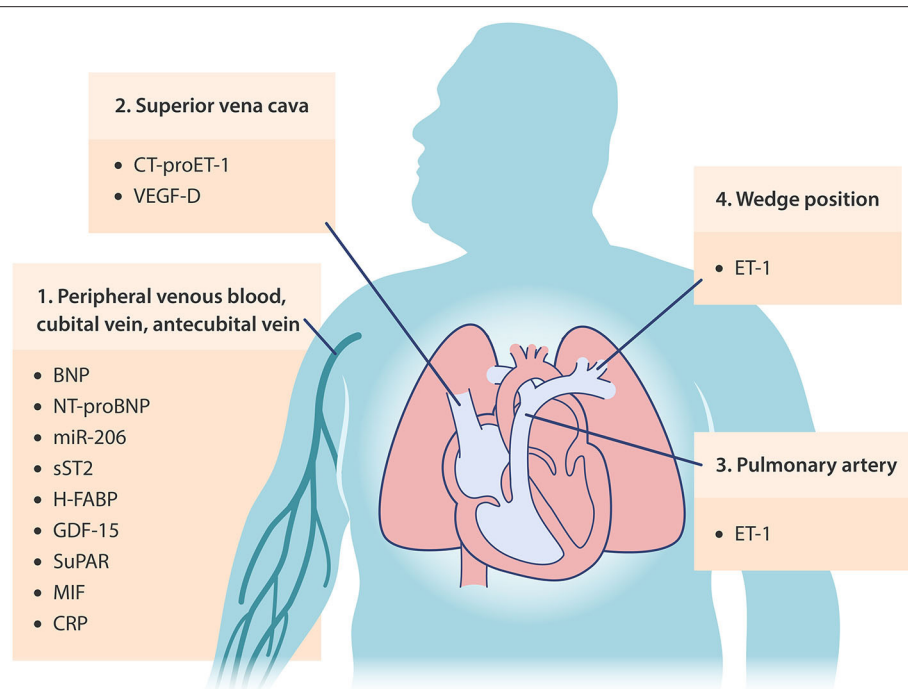


FIGURE 3 | Occurrence locations of potential circulating biomarkers in PH-LHD. BNP, brain/B-type natriuretic peptide; NT-proBNP, N-terminal proBNP; miR-206, microRNA-206; sST2, soluble suppression of tumorigenicity 2; H-FABP, heart type fatty acid binding protein; GDF-15, growth differentiation factor 15; suPAR, soluble urokinase plasminogen activator receptor; MIF, macrophage migration inhibitory factor; CRP, C-reactive protein; ET-1, endothelin-1; CT-proET-1, C-terminal pro-ET-1; VEGF-D, vascular endothelial growth factor-D.

circulating miR-206 levels may be compensated for by an increase in the intracellular miR-206 concentration, further research is needed to determine whether circulating miR-206 can be useful in the early identification of PH in LHD patients.

FUTURE CANDIDATE BIOMARKERS

In addition to the emerging biomarkers discussed above, several biomarkers of distinct processes in cardiopulmonary regulation may aid in the characterization, differentiation, and/or determination of disease management of PH-LHD (**Figure 3** and **Table 2**). Soluble suppression of tumorigenicity 2 (sST2), also known as interleukin-1 receptor-like 1 (IL1RL1), is one such example. sST2 is a marker of mechanical stress and ventricular remodeling, which identifies patients with HFpEF and has been shown recently to be elevated in plasma of patients with Group 2 PH, compared to subjects admitted for elective coronary angiography (in whom CAD was excluded), in a single-center retrospective study performed by Mirna et al. (85). While cautious interpretation of data is needed and detailed information regarding baseline characteristics, especially for Group 2 patients, is lacking, Mirna et al. also showed that Group 2 PH patients had higher circulating levels of growth differentiation factor 15 (GDF-15; a marker of cell injury and inflammation), heart type fatty acid binding protein (H-FABP; a marker of ongoing myocardial damage), and soluble urokinase plasminogen activator receptor (suPAR; a marker

of ongoing inflammation) (85). In addition, higher plasma levels of macrophage migration inhibitory factor (MIF), a proinflammatory cytokine, were found to be associated with greater PASP and natriuretic peptide levels in HFpEF patients (86). Mazurek et al. also showed a trend toward higher levels of galectin-3, a beta-galactoside-binding protein which has been implicated in inflammation and fibrosis, in serum collected from pulmonary artery of patients with PH-HFpEF relative to those with PAH (48). While this may be associated with greater systemic fibrosis and inflammation seen in HFpEF populations, no correlations between galectin-3 levels and any measured hemodynamic endpoints were found in PH-HFpEF or PAH patients in this study (48). The common biomarker of inflammation and cardiovascular distress, C-reactive protein (CRP), has also gained some interest as a potential biomarker of PH-LHD. Using a CRP cutoff value of > 3 mg/dl, the RELAX trial reported that approximately 60% enrolled HFpEF patients have elevated CRP (87). Higher CRP levels were associated with younger age, higher BMI, COPD, atrial fibrillation, RV dysfunction, reduced exercise tolerance, higher circulating levels of ET-1, aldosterone, and NT-pro BNP. While this led the authors to conclude that high CRP levels may identify a unique HFpEF phenotype that is associated with comorbidity-driven systemic inflammation, CRP levels were not associated with PH or LV function in this study cohort (87). Lastly, platelets from patients with PH-HFpEF have also been shown to exhibit a distinct metabolic phenotype compared to patients with PAH (88).

TABLE 2 | Collected studies of future candidate biomarkers in PH-LHD.

Authors (ref #)	Biomarkers	Patients (n)	Controls (n)	Primary findings	Correlation with hemodynamics
Mirna et al. (85)	sST2, GDF-15, H-FABP, suPAR, BNP	Group 1 PH (13), Group 2 PH (35), Group 3 PH (7), Group 4 PH (12), Group 5 PH (19), retrospective analysis	Elective coronary angiography w/o CAD (74)	BNP, GDF-15, H-FABP, and suPAR are elevated in Group 2 PH patients and correlate with some hemodynamic measurements	H-FABP w/ RVEDD, RAA, TR: ↗ sST2 w/ RVEDD, RAA, TR, PASP: ↗ sST2 w/ TAPSE: ↘ GDF-15 w/ RVA, RAA: ↗ suPAR w/ RVEDD, RAA, TR: ↗ suPAR w/ TAPSE: ↘
Luedike et al. (86)	MIF	HFpEF w/ MIF > 51.58 ng/ml (31), prospectively enrolled	HFpEF w/ MIF ≤ 51.58 ng/ml (31)	High MIF associated w/ 180-day mortality/hospitalization and more severe HFpEF symptoms	MIF w/ PASP: ↗
Mazurek et al. (48)	Galectin-3	PH-HFpEF (39), prospectively enrolled	PAH (37)	Galectin-3 positively correlated with mortality	N/A
DuBrock et al. (87)	hs-CRP	HFpEF w/ CRP > 3 mg/l (121), retrospective analysis	HFpEF w/ CRP ≤ 3 mg/l (93)	High CRP associated with increased comorbidity burden and presence of RV dysfunction	N/A
Nguyen et al. (88)	Platelet bioenergetics	PH-HFpEF (20), retrospective analysis	Healthy volunteers (20)	PH-HFpEF patient platelets display increased maximal OCR and reserve respiratory capacity	Reserve respiratory capacity w/ RV SWI: ↘

CAD, coronary artery disease; GDF-15, growth differentiation factor 15; H-FABP, heart-type fatty acid binding protein; hs-CRP, high-sensitivity C-reactive protein; MIF, macrophage inhibitory factor; OCR, platelet oxygen consumption rate; RAA, right atrial area; RVEDD, right ventricular end diastolic diameter; sST2, soluble suppression of tumorigenicity 2; suPAR, soluble urokinase plasminogen activator receptor; SWI, stroke work index; TR, severity of tricuspid regurgitation. ↗ indicates positive correlation with the corresponding biomarker; ↘ indicates an inverse correlation with the corresponding biomarker.

Unlike platelets from PAH patients, only an increase in maximal respiratory capacity, but not in glycolytic rate, was observed in platelets from PH-HFpEF patients relative to healthy controls. The enhancement of platelet maximal respiratory capacity was found to be associated with RV dysfunction, but not with mPAP or PVR in patients with PH-HFpEF (88). Whether or not a measurement of these factors can provide a method for distinguishing PH-LHD patient phenotypes, identifying at-risk subjects, and predicting PH-LHD patient outcomes requires additional research.

Apart from clinical studies, recent advances in pre-clinical models have provided new insights in the development of future biomarkers. Plasma levels of leptin, an adipokine known to induce secretion of proinflammatory cytokines and reactive oxygen species (ROS) generation, were found to be associated with PH severity in rats with PH-HFpEF induced by high-fat diet (HFD) in combination with supra-coronary aortic banding (SAB) and the antipsychotic olanzapine (89). Decreased plasma levels of the cardioprotective adipokine adiponectin after Sugan-mediated induction of PH-HFpEF were also observed in leptin receptor-deficient obese ZSF1 rats (90). Interestingly, metformin (the first-line drug for type 2 diabetes) was found to improve metabolic syndrome and pulmonary pressures in these models through modulation of leptin, adiponectin, as well as other mechanisms involving, at least in part, suppression

of interleukin-6-associated inflammation and activation of siruin-3-mediated skeletal muscle glucose uptake (89, 90). Moreover, metformin has recently been shown to prevent RV dysfunction via improved insulin resistance and reduced RV lipid accumulation in HFD-treated mice (91). These findings have led to an ongoing trial designed to evaluate the effect of metformin in patients with PH-HFpEF (NCT03629340). Nitrite, a dietary precursor of NO, has displayed similar qualities in a rat model of PH-HFpEF where it functions to increase adiponectin levels and improve skeletal muscle insulin resistance (90). A recent study by Simon et al. reported that inhalation of nitrite reduced pulmonary, RA, and pulmonary capillary wedge pressures in patients with PH-HFpEF (92). A clinical trial is now underway to examine the effect of oral nitrite in PH-HFpEF patients (NCT03015402). Moreover, therapies targeting ET-1 and NO with a Rho-kinase (ROCK) inhibitor, fasudil, have also been shown to be effective in rats with end-stage PH-LHD induced by SAB and in patients with PH-HFpEF, with more favorable results in the CpcPH-HFpEF population (93, 94).

SUMMARY AND FUTURE PERSPECTIVES

While many challenges still exist, researchers in the field are encouraged to further expand the growing array of circulating

biomarkers to guide future research toward facilitating screening, diagnosis, refinement of specific patient phenotypes, and development of therapeutic approaches. Consensus to concise and standardized definition for hemodynamic phenotyping is required, along with longitudinal data consisting of multiple time points and evaluation in large, multi-center studies to validate the viability of the aforementioned potential biomarkers and to explore future candidates. The ongoing PVDOMICS study supported by the National Institutes of Health/National Heart, Lung, and Blood Institute (NIH/NHLBI) may also reveal new relevant information (95). Since HFpEF and PH are recognized as multiorgan and systemic disorders (96, 97), the quest toward identification of meaningful extra-cardiopulmonary biomarkers may aid in providing important new insights into this clinically challenging syndrome and driving PH-LHD research forward.

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Unexpected Acceleration in Treprostinil Delivery Administered by a Lenus Pro® Implantable Pump in Two Patients Treated for Pulmonary Arterial Hypertension

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Intravenous treprostinil administration by an implantable pump is an attractive option for pulmonary arterial hypertension (PAH) treatment and is the subject of recent publications. Short-term studies are promising, but there is still a lack of long-term prospective data. We analyzed the treprostinil flow rate administered by the Lenus Pro® implantable pump in 2 patients suffering from PAH during follow-up times of respectively 4.2 and 3 years. The flow rate delivered by the pumps in these 2 patients exceeded the manufacturer admitted margin of error within 2 years and continued to increase to reach, respectively, 158 and 120% of the expected flow rate at the end of the follow up. In one case, the implantable pump had to be removed for this reason. The *ex-vivo* flow rate of the withdrawn pump determined in the laboratory reached 173% of the predicted value. This correlated with the *in-vivo* measurement, which suggests a continuous flow increase even after pump removal and without treprostinil use. Spontaneous flow increase from such an implantable pump is a potentially major pitfall, which needs to be identified and actively managed by the responsible clinicians.

Keywords: pulmonary arterial hypertension, prostacyclin analogs, implantable pump, treprostinil delivery, internal device

INTRODUCTION

The prostacyclin pathway is a major target for treatment of pulmonary arterial hypertension (PAH), recommended for high-risk patients (1). Due to their short half-lives, prostacyclin analogs are usually administered by continuous intravenous (IV), subcutaneous (SC) or intermittent inhaled routes. Treprostinil has a longer half-life and is much more stable at room temperature than epoprostenol permitting oral, SC or IV route administration. Its use has shown improvement in 6-min walk distance, functional class and pulmonary hemodynamics compared to placebo (2–4). The use of oral treprostinil is limited to first line therapy in non-high risk patients, due to its lesser efficacy than parenteral treprostinil (4). Unfortunately SC treprostinil administration is often

limited by significant local side effects (2). Continuous IV administration is traditionally performed by central venous catheter connected to an external pump.

The development of an implantable pump for continuous IV treprostinil administration by Tricumed Medizintechnik GmbH (LenusPro[®] pump, approved in Europe in 2009) has attracted scientific attention as it may offer less restrictive use by patients as well as reducing catheter-related bloodstream infection rates (5–7). The Lenus Pro[®] pump contains a micro-infusion system that operates by pressurized gas storage in a titanium tank which is accessible by a puncturable silicone interface directly under the skin. Two pump models are available, one with a 20 ml capacity, the other with a 40 ml capacity. The device is usually implanted in the abdominal subcutaneous tissue and connected to a tunneled catheter accessing the superior vena cava (**Figure 1**). The flow is regulated by a glass capillary chip intended to guarantee a constant flow rate (1.3 ml/day), irrespectively of gas pressure. As the flow rate is fixed, drug titration is performed by adjusting treprostinil concentration. Percutaneous refills are usually performed every 14–28 days, depending on the size of the pump.

Two small prospective studies assessed periprocedural and 6-month safety, respectively, reporting good clinical results when using IV treprostinil administered by this pump (5, 6). We

share here our experience with 2 cases receiving intravenous treprostinil via a LenusPro[®] implantable pump during a follow-up of, respectively, 4.2 and 3 years.

MATERIALS AND METHODS

The residual treprostinil volume in the pump was measured by a PAH specialized nurse before each refill procedure for the two patients treated by IV treprostinil administered by an implantable pump, as recommended by the manufacturer. The time between refill procedures and the volume administered were also reported in the registry by the nurse. Using these data we calculated the treprostinil flow rate between each refill procedure, for both patients during the follow-up period.

The *ex-vivo* flow rate of the withdrawn pump was determined in the laboratory by refilling it with saline solution (0.9 % NaCl) and keeping it in a constant 37°C water bath for 7 days. The flow rate was established by weighting (to 100 µg precision) at 24 h intervals a 5 ml polystyrene tube (352058 Becton-Dickinson) filled by the pump.

Written informed consent was obtained from the individuals for the publication of any potentially identifiable images or data included in this article.



FIGURE 1 | The pump (Lenus Pro[®], 20 ml) implanted in the left hypochondria for intravenous treprostinil delivery (patient 1).

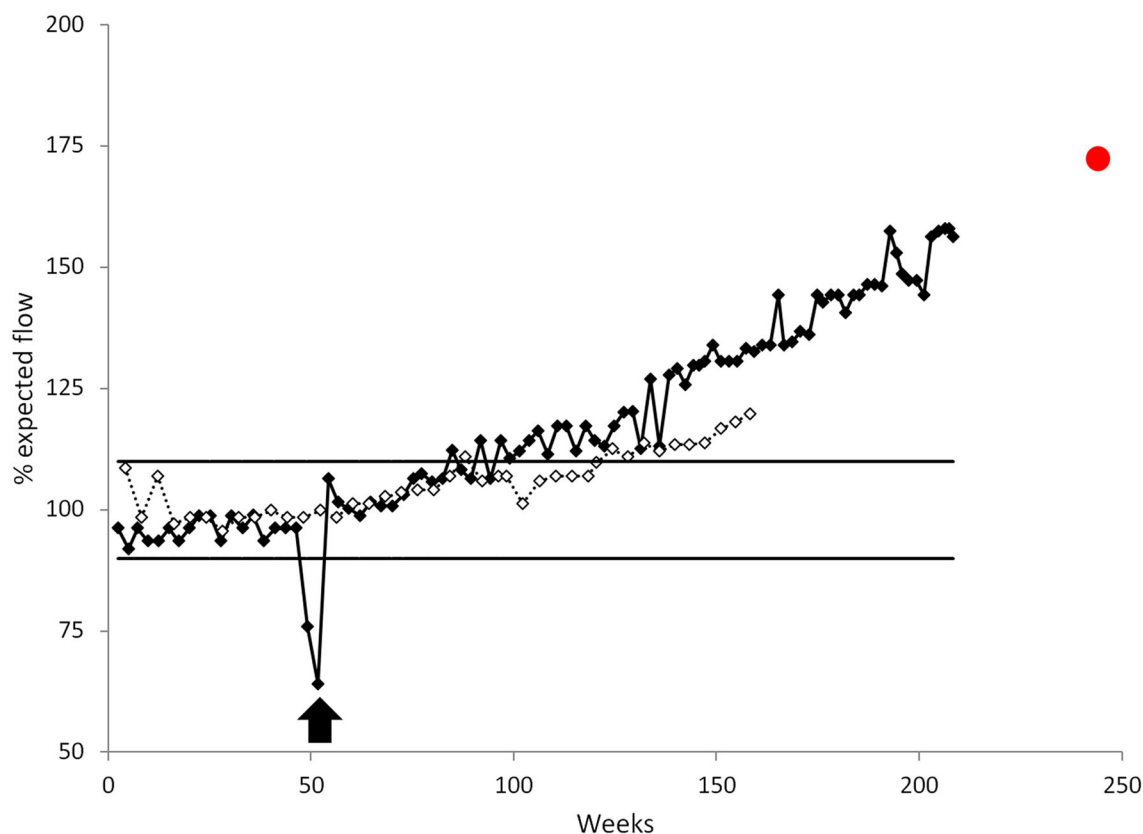


FIGURE 2 | Treprostinil delivery flow rate by the LenusPro® pump for patient 1 (black) and patient 2 (white) expressed relative to the expected calibrated flow. For patient 1, a catheter thrombosis resulted in a dramatic flow decrease that was treated by a standard repermeabilization procedure (black arrow). Horizontal lines represent the $\pm 10\%$ accepted flow error. The red dot represents the *ex-vivo* flow rate of the withdrawn pump determined in experimental condition by refilling it with a 20 ml saline solution.

CASES

The first case is a 47-year-old female patient suffering from drug-induced PAH. Under combined dual therapy (sildenafil, bosentan) the initial evolution was favorable. Due to clinical and hemodynamic deterioration, treatment of SC treprostinil was introduced later. The tolerance of the latter quickly became limited by pain at the injection site, justifying a switch to IV treatment. The patient being reluctant to IV access with external pump an implantable pump was proposed (LenusPro® 20 ml, **Figure 1**). The target dosage was reached in 2 months (47 ng/kg/min) while the clinical improvement allowed stabilization in functional class II. One year after implantation a central catheter thrombosis occurred resulting in a 40% flow rate decrease (see **Figure 2**). This was only discovered by the observation of an increased residual volume in the pump whereas the electronic flow rate sensor should have triggered an alarm in such a case. The catheter thrombosis was successfully treated by gentle rinsing with a heparin solution under fluoroscopic control by contrast injection, according to the manufacturer's guide. During the 4 years of follow-up we observed a gradual decrease in the residual treprostinil

volume at refilling resulting in a shorter period between refill procedures.

The pump is calibrated by the manufacturer to a constant flow rate ($1.04 \text{ ml/day} \pm 10\%$) whereas 4.2 years after implantation the observed flow rate had increased to 1.64 ml/day (77.4 ng/kg/min) which is a 58% flow increase above the predicted value (**Figure 2**). The increase in the treprostinil delivery flow rate resulted a proportional drug-dose increase that was, fortunately, well tolerated but induced a significant cost increase. The pump was removed and replaced by a 40 ml model due to the patient still being reluctant to other modalities of treatment and still being stabilized on this therapy. The *ex-vivo* flow rate of the withdrawn pump was established at $1.797 \pm 0.017 \text{ ml/day}$, corresponding to a 73% flow increase over the expected value (**Figure 2**). This suggests a continuous flow increase even after pump removal and independent of treprostinil use. Considering this particular case, it was considered that the dose increase may have contributed to the progressive need of the PAH evolution and that it may have participated to maintain the patient in a stable condition.

The second case is a 22-year-old male patient followed since childhood for idiopathic PAH. The clinical course was initially favorable under inhaled iloprost monotherapy

sequentially combined with bosentan and, later, sildenafil due to hemodynamic deterioration. Despite the triple therapy the clinical and hemodynamic response was unsatisfactory. The iloprost treatment was successfully changed from inhaled to IV route delivered by a central venous catheter. Following that time the patient developed numerous catheter-related infections requiring hospitalizations and central line replacements. In order to reduce the risk of infection an implantable pump (LenusPro[®] 40 ml) was fitted.

As with the first case we observed a progressive increase in treprostinil flow reaching 1.52 ml/day which is a 20% increase over the calibrated value of 1.27 ml/day ($\pm 10\%$) (**Figure 2**). The drug dilution was adjusted at each refill procedure in order to counterbalance the increased flow rate and maintain a stability of treatment at 35–41 ng/kg/min without any adverse event or side effect related to the treatment. Despite these measures, further hemodynamic evaluation concluded that the PAH was worsening under maximal treatment. The patient was referred for a lung transplant but died suddenly whilst still on the waiting list.

DISCUSSION

The currently available studies of the Lenus Pro[®] pump have drawn justifiable attention with good results reported and without significant short-term complications or safety issues (5, 6). Nonetheless the studies included small collectives and only two of them were prospective.

Interestingly a recent retrospective study including 129 patients with pulmonary hypertension (PH) treated by IV treprostinil administered by the Lenus Pro pump[®] has shown 82 complications after a median follow-up of 19 months. Most of these events were non-infectious catheter-related complications. As in the cases previously described, flow rate increases occurred in five patients. Four of them required pump replacement for this reason. Following treprostinil flow rate increase, two patients developed acute cardiac decompensation requiring intensive care unit admission. One patient underwent pump replacement. In the other patient, the treprostinil dose was adjusted to the higher flow rate without explantation (7). An unexpected increase in the flow rate administered by a LenusPro pump was also reported in a 14 years old patient (8).

In their multicentric study, Ewert et al. reported a septum defect of a Lenus Pro pump in one patient (6). This resulted in a severe overdose with subsequent hypotension, necessitating pump replacement. This septum defect was considered by the authors likely to be due to improper use of non-approved needles for refilling the pump.

This highlights the importance of maintaining a constant prostacyclin analog flow rate for patient stability and improvement as previously described in another study (9).

A warning has been published in 2013 by the manufacturer reporting occurrences of increased flow of the LenusPro[®] pump after several years of use. It is speculated that the molecule or its solvent provokes an increase in the cross-section of the glass capillary canal of the chip and therefore an augmentation of flow

by 10%, especially when pumps are used for longer periods of time (2–4 years). The corrective action was introduced by the new development of a chip canal with a more resistant glass (10).

Nevertheless, an unexpected increase in the drug administration rate was reported with a Lenus Pro implantable pump coming from upgraded series, i.e., after the 2013 safety note of the manufacturer (11). This increase from 1.3 to 1.7 ml/day resulted in symptoms of deterioration of exercise capacity and fatigue without hemodynamic collapse. The second pump of the first described patient of the present work was also introduced after 2013, demonstrating that the corrective action introduced may not have been sufficient.

Recently, the variance of the fixed flow rate during long-term follow-up has been addressed on 126 patients (12) during a median follow-up of 12 months. Based on 2,853 refills, the relative flow rate deviation between each individual refill was between -10% and $+10\%$ in 94.5% of cases. However, three refill cases (0.1%) had a relative flow rate deviation of more than 40% from the previous refill (one case of -40% and two cases of $+40\%$).

SynchroMed[®] II (Medtronic) is the second implantable pump currently approved for IV treprostinil administration in patients with PH. Unlike the Lenus Pro[®] pump, SynchroMed[®] II pump is battery-driven and has an adjustable flow rate, between 0.048 and 24 ml/day. Nonetheless, recent studies reported decreased flow rate and flow rate accuracy over time with this device (13, 14). Gomberg-Maitland et al. reported also eight pump failures events (13). One of them was undetected, leading to subsequent death of the patient. After this, pump design have been implemented, addressing many of the observed pump failures and motor stalls. Later, a programmable battery-driven model (Siromedes[®] pump) with adjustable flow has been developed by the German company Tricumed and approved in 2014. The three devices have similar use and implantation procedures.

The use of an implantable pump for IV treprostinil administration in PAH management is attractive, but the unexpected increase in the rate of treprostinil delivery by the Lenus Pro[®] pump is a major issue.

Such an increase is potentially life threatening and must be detected and actively managed by the responsible clinicians. This therapy should be administered in PH specialized centers by clinicians familiarized with the device. Reporting the time between refill procedures, the residual volume and the volume administered at each refill procedure should allow early detection of any derivation from the target flow rate and anticipate potential pump failure.

As an alternative option of removing and replacing the defective pump, adjustment of the drug dilution has to be considered, as long as the flow deviation from the factory setting can be anticipated.

This report is limited to the description of only two cases. Nevertheless, the frequency of adverse events such as previously described in our cases could have been underestimated in the studies actually available, due to the retrospective bias. Prospective studies are still needed to assess the long-term safety and efficacy of treprostinil administration by this device.

DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author.

ETHICS STATEMENT

We hereby confirm that written informed consent was obtained from the individuals for the publication of any potentially identifiable images or data included in this article.

AUTHOR CONTRIBUTIONS

GK, MB, and FL contributed conception and design of the article. AB and FL performed the analysis of the

withdrawn pump. GK wrote the first draft of the manuscript. MB, AB, and FL wrote sections of the manuscript. All authors contributed to the article and approved the submitted version.

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The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Portopulmonary Hypertension: From Bench to Bedside

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Portopulmonary hypertension (PoPH) is defined as pulmonary arterial hypertension (PAH) associated with portal hypertension and is a subset of Group 1 pulmonary hypertension (PH). PoPH is a cause of significant morbidity and mortality in patients with portal hypertension with or without liver disease. Significant strides in elucidating the pathogenesis, effective screening algorithms, accurate diagnoses, and treatment options have been made in past 20 years. Survival of PoPH has remained poor compared to IPAH and other forms of PAH. Recently, the first randomized controlled trial was done in this patient population and showed promising results with PAH specific therapy. Despite positive effects on hemodynamics and functional outcomes, it is unclear whether PAH specific therapy has a beneficial effect on long term survival or transplant outcomes. In this review, we will discuss the epidemiology, pathophysiology, clinical and hemodynamic characteristics of PoPH. Additionally, this review will highlight the lacunae in our current management strategy, challenges faced and will provide direction to potentially useful futuristic management strategies.

Keywords: portopulmonary hypertension, pulmonary arterial hypertension, portal hypertension, liver transplant, MELD exception

INTRODUCTION AND BACKGROUND

Pulmonary arterial hypertension (PAH) is a chronic progressive disease characterized by elevated pulmonary artery pressure and pulmonary vascular resistance (PVR) eventually leading to right heart failure and premature death. PAH associated with portal hypertension is called portopulmonary hypertension (PoPH), and is a subset of Group 1 PH (1). Similar to other causes of PAH, PoPH is characterized by the presence of intimal proliferation, medial hypertrophy, adventitial fibrosis of the muscular arteries and plexiform arteriopathy, and is thought to be histopathologically indistinguishable from other PAH phenotypes (2–4). Importantly, there does not seem to be a link between the presence or severity of PoPH with the degree of underlying liver dysfunction or hepatic venous pressure gradient (5, 6).

Patients with PoPH may be treated with PAH specific therapies, which may decrease the severity of the disease, improve functional parameters and hemodynamics, and allow for liver transplant (LT) (7). Despite a rapidly evolving understanding of PoPH, challenges remain in the diagnosis, treatment and transplantation of patients with PoPH. In this review, we will discuss the latest expert opinion in diagnosis and management of PoPH, as well as identify key areas of future research focus.

DEMOGRAPHICS AND CLINICAL CHARACTERISTICS OF PATIENTS WITH PoPH

The exact prevalence of PoPH is difficult to determine. The annual incidence of all types of PAH is <10 patients per million population, and PoPH patients are a subset of this group (8). PoPH was the third most common form of PAH in a population-based epidemiologic study in France (9). Historically, PoPH was thought to be 5–10% of all patients with PAH (10, 11). More recently, French authors have reported that the proportion of newly diagnosed (incident) patients with PoPH is as high as 15% of all patients with PAH, and is continuing to rise as wider screening practices are adopted (12). PoPH is thought to be present in anywhere between 2 and 10% of patients with portal hypertension (6, 13). Furthermore, while the vast majority of cases of PoPH are in patients with portal hypertension related to cirrhosis, non-cirrhotic causes of portal hypertension (including portal vein thrombosis, granulomatous disease, auto-immune disorders, drug reactions, infections, and congenital abnormalities) are also important contributors (14, 15). McDonnell et al. showed a prevalence of histopathologic changes of PAH of 0.61% in autopsies of patients with cirrhosis (16), and small cohort studies have shown prevalence of PoPH in patients with cirrhosis undergoing transplant evaluation to be between 5 and 6% (17–19).

Although the hemodynamic profile of patients with PoPH is better than in patients with idiopathic/familial PAH (IPAH/HPAH), their overall mortality is similar or worse (12, 20–22). The French Pulmonary Hypertension Registry (FHPR) shows a 1, 3, and 5-years survival of 84, 69, and 51%, respectively, for patients with PoPH (12), which is similar to the survival data for patients with IPAH, HPAH and anorexigen associated PAH (23). In contrast, analysis of the US based REVEAL registry showed that patients with PoPH had significantly worse survival when compared to patients with IPAH/FPAH: 67 vs. 85% at 2 years, and 40 vs. 64% at 5 years (**Figure 1**) (20). These findings were similar in the Spanish REHAP registry, where the 5 year mortality was 49 and 69%, for PoPH and IPAH/HPAH, respectively (**Figure 2**) (24, 25). Furthermore, severe PoPH is associated with significantly decreased survival in patients undergoing LT (7, 26), and severe PoPH precludes liver transplantation and significantly affects the course of hepatic failure in these patients (26, 27).

The lack of better outcomes in patients with PoPH as compared to their IPAH/FPAH counterparts is not entirely clear, especially in light of their relatively better hemodynamic profile and functional status (20, 25). One possible explanation is increased deaths from liver related events, which account for 25–33% of deaths in patients with PoPH, as compared to only 5% in patients with IPAH (22, 25). In patients with liver disease, higher Child-Pugh Scores are associated with worse outcomes (14, 22), which makes it clear that the severity of liver disease (rather than PoPH) is the main factor in worse outcomes in this population. Another possibility is a relative delay in PAH-specific treatments within the PoPH cohort. PoPH patients started on

PAH-specific therapy live longer, despite worse hemodynamics, when compared to PoPH patients not on therapy (25). However, the percentage of PoPH patients on pulmonary vasodilators at both time of enrollment and at 90-days post enrollment into the REVEAL registry was significantly lower than their IPAH/FPAH counterparts (20). The cause of this delay is likely multifactorial, and includes milder symptoms, better hemodynamics, and lack of definitive data to suggest the ideal agent in patients with PoPH.

Likewise, the difference in survival seen between different PoPH cohorts are also likely due to a variety of factors, which include differences in regional screening practices for the presence of PoPH. Patients in the US and Spain with cirrhosis are typically screened at time of liver transplant evaluation, which likely skews the diagnosis of PoPH to patients with more severe liver disease. In contrast, all patients with cirrhosis are screened for PoPH in France, which likely leads to increased rates of diagnosis of PoPH in patients with milder liver disease, which in turn increases the overall survival rates. Rates of referrals of patients to PH specific centers may differ by region (the REVEAL registry included PAH patients treated in the community in addition to academic PH-centers). There may also be differences in baseline characteristics of patients enrolled in each respective registry (including New York Heart Association Functional Class [NYHA FC] and severity of underlying liver disease at the time of enrollment). Additional prospective studies are needed as PoPH-specific management guidelines are established and standardized, to see if this gap can be improved.

PATHOGENESIS OF PoPH

The underlying mechanisms for the development of PoPH are poorly understood and continue to be an area of active research. Liver cirrhosis and portal hypertension lead to splanchnic vasodilatation and formation of portosystemic shunts, which are thought to contribute to PoPH pathogenesis in multiple ways (**Figure 3**). Mechanically, increased intrahepatic resistance to flow from cirrhosis results in an increased portal pressure gradient and portosystemic collateralization through the reperfusion/dilation of preexisting vessels and through the generation of new vessels (28). On a molecular level, portosystemic shunting enables blood to bypass the liver, and thus evade hepatic metabolism of vasoactive substances. This direct reduction in the peripheral vascular resistance, combined with indirect vasodilation via intestinal vasoactive substances that are now able to bypass the liver and reach systemic circulation, ultimately result in a hyperdynamic state. This phenomenon has been seen in patients immediately after undergoing a transjugular intrahepatic portosystemic shunt (TIPS) placement, and persisted even a month out (29). Likewise, levels of vasoactive substances acting in the pulmonary circulation (including pulmonary vasodilators nitric oxide and prostacyclin, as well as pulmonary vasoconstrictors endothelin-1, thromboxane A₂, and serotonin), are similarly dysregulated in cirrhosis (30–32). The imbalance of these vasoactive substances in the pulmonary vasculature results in net vasoconstriction and pulmonary vascular resistance elevation.

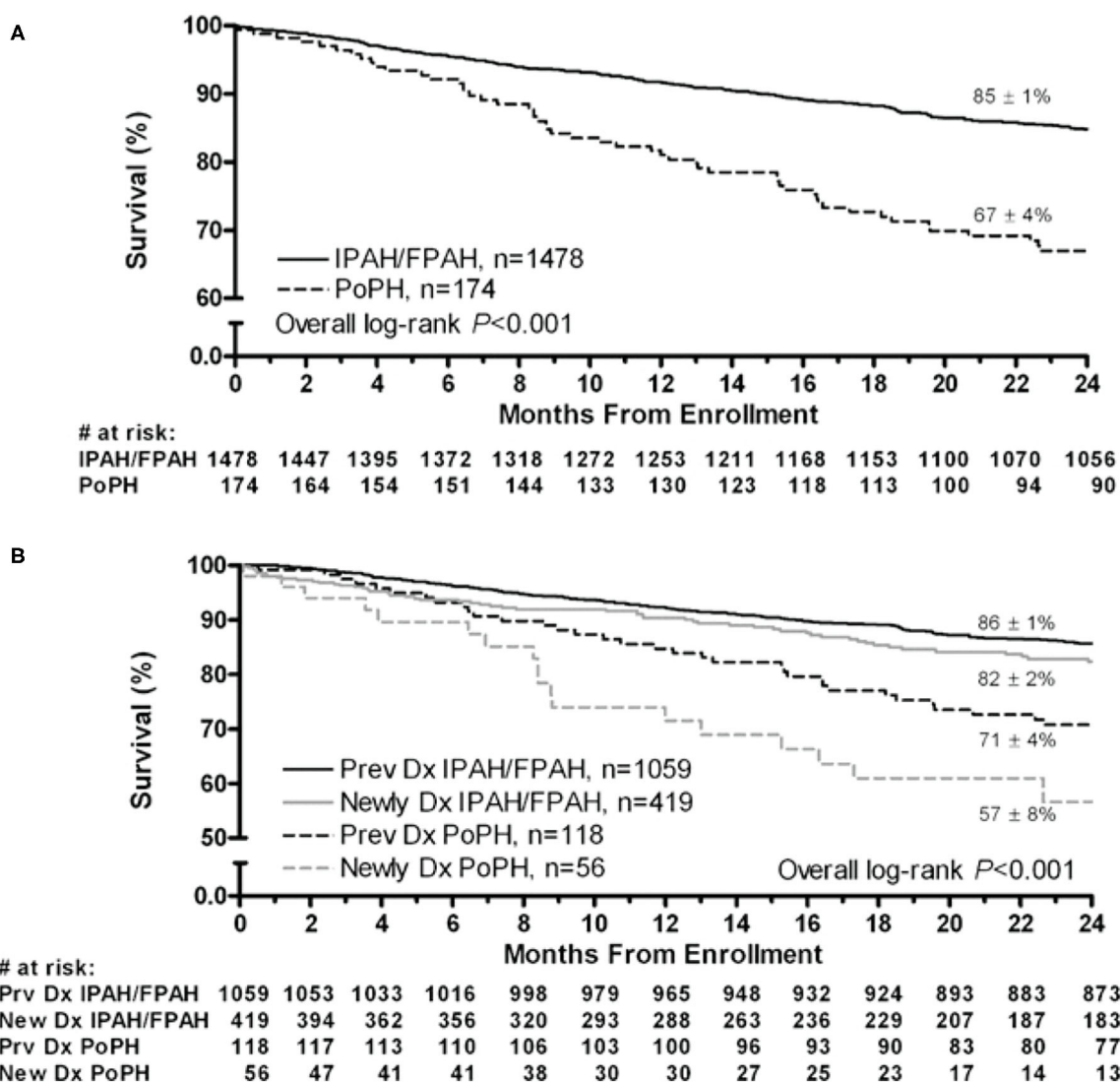
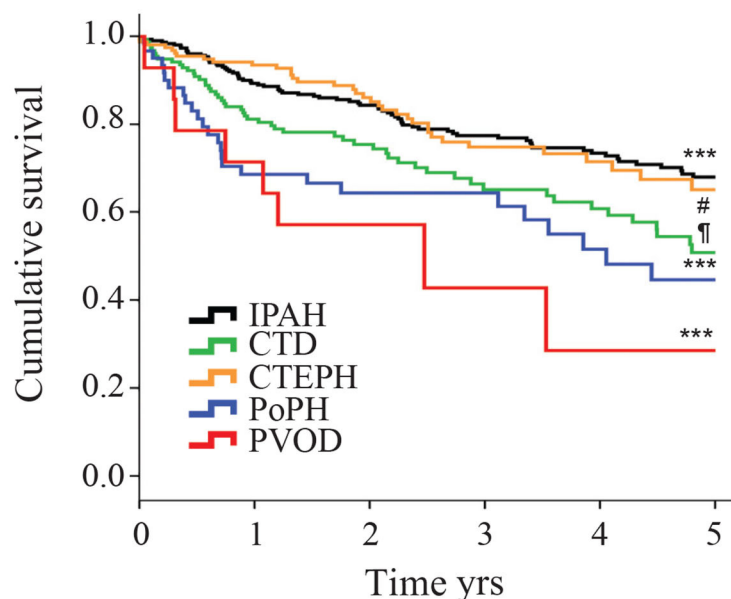


FIGURE 1 | Two-year survival from enrollment. **(A)** Patients with PoPH and IPAH/FPAH. **(B)** Stratified by duration of disease from enrollment. Dx, diagnosis; FPAH, familial pulmonary arterial hypertension; IPAH, idiopathic pulmonary arterial hypertension; PoPH, portopulmonary hypertension. Reproduced with permission of Elsevier: Chest 141 (4) 906-915; doi: 10.1378/chest.11-0160 Epub 21 July 2011.

Shear stress from persistently high flows leads to endothelial cell injury and the activation and repression of genes that participate in the vascular remodeling process, and is believed to play a role in various forms of PAH (33–35). Exposure of and damage to the underlying arterial smooth muscle results in smooth muscle proliferation and thickening of the tunica intima, media, and adventitia within the pulmonary vasculature (4). The arterial wall thickening, in turn, can result in more sluggish pulmonary blood flow, platelet aggregation, and thrombus formation, which can then become recanalized over time (3). Autopsy data of patients with PoPH confirms that the end stage of this process is medial and intimal thickening, plexiform lesions and fibrotic venular obstruction (36). Together, these vascular lesions result in permanently elevated PVR that distinguishes

PoPH from other causes of elevated mPAP in patients with portal hypertension (**Figure 5**).

Signaling mediated by bone morphogenic protein receptor 2 (BMPR2), a member of the TGF- β superfamily, has been shown to play a key role in familial pulmonary hypertension, and mutations in the gene encoding this receptor have also been found in 15–40% of idiopathic cases of PAH (37, 38). Bone morphogenic protein 9 (BMP9) is a circulating factor produced by hepatic stellate cells, which serves as a ligand for BMPR2 signaling, and has been shown to have a protecting effect against hepatic fibrosis (39). Recent studies have shown that patients with PoPH have significantly lower circulating levels of BMP9 as compared to control patients with advanced liver disease and no evidence of pulmonary hypertension (40). BMP9



At risk n							
IPAH	314	304	236	179	136	108	p-value [#]
CTD	157	149	98	67	46	37	0.023
CTEPH	162	153	117	82	51	34	0.95a1
PoPH	61	58	34	26	21	14	<0.001
PVOD	15	14	8	4	3	2	<0.001

FIGURE 2 | Kaplan-Meier estimates of 5-years survival from time of diagnosis in different pulmonary hypertension subtypes. IPAH, Idiopathic Pulmonary arterial hypertension; CTD, connective tissue disease; CTEPH, chronic thromboembolic pulmonary hypertension; PoPH, portopulmonary hypertension; PVOD, pulmonary veno-occlusive disease. The *p*-value for the overall comparison is <0.001. # compared with IPAH. Reproduced with permission of the © ERS 2020: European Respiratory Journal 40 (3) 596-603; doi: 10.1183/09031936.00101211 Published 31 August 2012.

levels were likewise significantly lower in patients with PoPH as compared to patients with other etiologies of group 1 PAH (41). Moreover, selective enhancement of endothelial BMPR2 with exogenous BMP9 has been shown to reverse the presence of PAH in multiple mouse models of PAH (42). Together, these findings suggest that BMP9 and BMPR2 associated downstream signaling pathways also likely play a role in the pathogenesis of PoPH.

More recently, activation of toll-like receptors (TLRs) by bacterial lipopolysaccharides (LPS) from bacterial translocation has been implicated in the pathogenesis of PAH. Mouse models deficient in BMPR2 exposed LPS were shown to develop evidence of pulmonary hypertension, which did not occur in their wild-type counterparts (43). The underlying mechanism for this is thought to be rooted in increased IL-6 production and activation of the STAT3 transcription factor, which is associated with increased expression of TLR4 (43). TLR4 is the main receptor for LPS and has been previously shown to promote the development of PAH (44–46). Together, these data suggest that exposure to bacterial endotoxins, which occurs more frequently in cirrhotic patients due to recurrent gut bacterial translocations, may

provide a “second hit” toward the development of PoPH in an otherwise predisposed host.

Lastly, there is growing evidence that altered sex hormone also contributes to the pathogenesis of PoPH. Estrogen binds to the promoter region of the BMPR2 gene and regulates its expression (47). Polymorphisms in CYP1B1 which metabolizes estrogen were associated with penetrance of PAH in women with BMPR2 mutations, but not men (48). Expression of CYP1B1 has also been shown to be increased in experimental models and human PAH (49). A candidate gene study to identify genetic risk factors in PoPH found single nucleotide polymorphisms (SNPs) in the genes coding for estrogen receptor 1 and aromatase, a critical enzyme in the estrogen metabolism (50). Recently, Al-Naamani et al. showed that in PoPH patients, the risk allele rs7175922 in CYP19A1 was associated with significantly higher levels of estradiol, urinary 2-hydroxyestrogen/16- α -hydroxyestrone (2-OHE/16 α -OHE1), plasma levels of dehydroepiandrosterone-sulfate and plasma levels of 16- α -hydroxyestradiol (16 α -OHE2) compared to the patients with liver disease but no PoPH (51). 16 α -OHE1 leads to the development of PAH in animals via upregulation

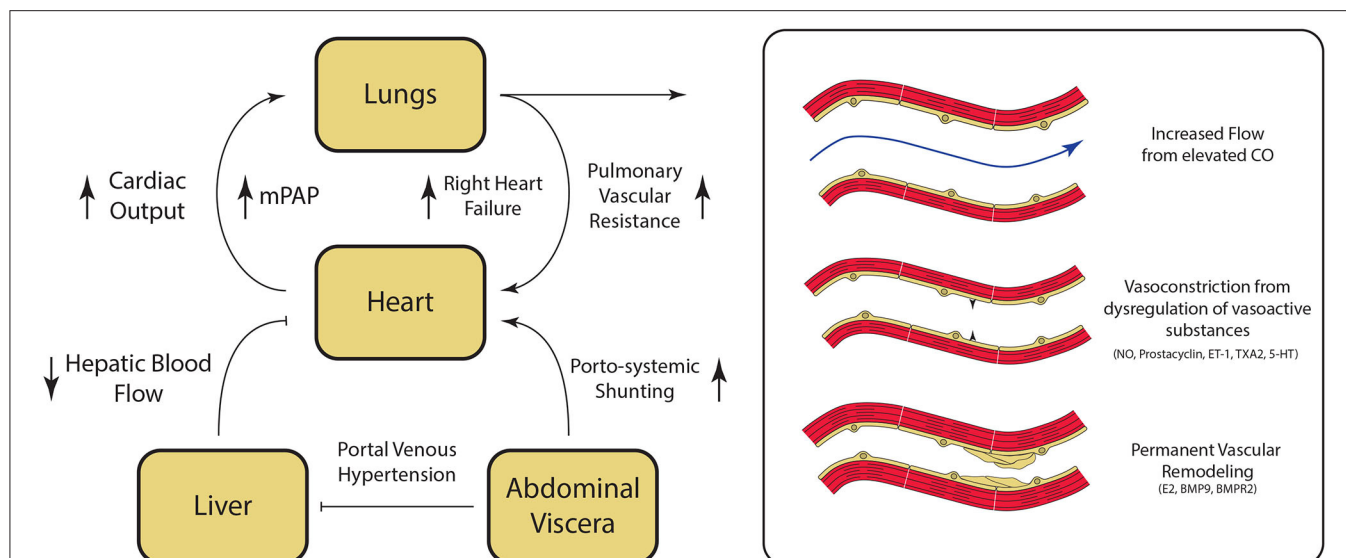


FIGURE 3 | Pathogenesis of Portopulmonary Hypertension (PoPH). Hepatic fibrosis results in portal venous hypertension, increased resistance to hepatic blood flow, and splanchnic vasodilation. Splanchnic vasodilation results in increase in overall circulating volume, and the diversion of blood flow from the liver to the heart via porto-systemic shunting resulting in an overall hyperdynamic state. Dysregulation of key regulator of pulmonary vascular tone associated with cirrhosis results in pulmonary vasoconstriction. At the same time, damage to the pulmonary endothelium and the underlying smooth muscle results in permanent vascular remodeling, which ultimately leads to the development of pulmonary hypertension. CO, cardiac output; NO, nitric oxide; ET-1, endothelin-1; TXA2, thromboxane A2; 5-HT, serotonin; E2, estrogen; BMP9, Bone Morphogenic Protein 9; BMPR2, Bone Morphogenic Receptor 2. Please refer to the text for additional details.

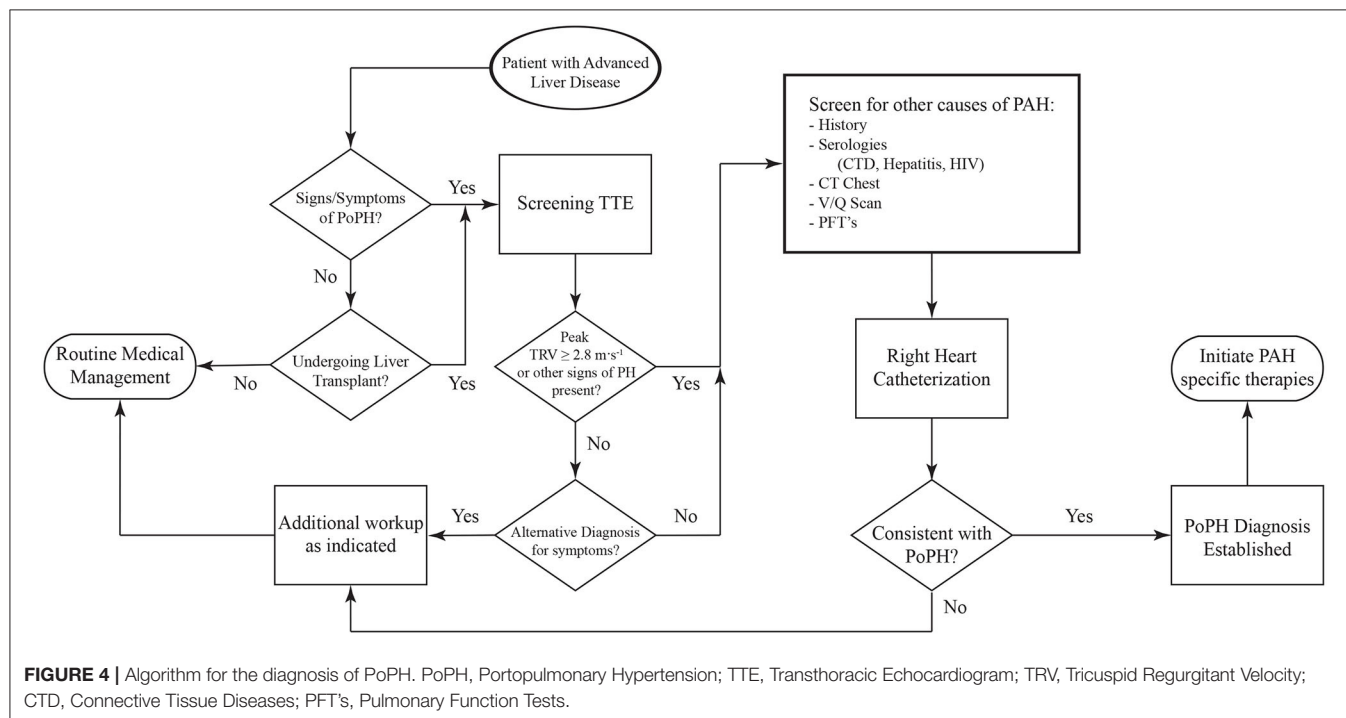


FIGURE 4 | Algorithm for the diagnosis of PoPH. PoPH, Portopulmonary Hypertension; TTE, Transthoracic Echocardiogram; TRV, Tricuspid Regurgitant Velocity; CTD, Connective Tissue Diseases; PFT's, Pulmonary Function Tests.

of miR-29 and has been associated with abnormal markers of insulin resistance (52). 16 α -OHE1 has also been shown to increase oxidative-stress related proliferation in pulmonary arterial smooth muscle cells from PAH patients (53). Similarly, Sahay et al. performed qualitative immunohistochemical (IHC)

staining of the explanted livers of PoPH patients and compared them with normal and cirrhotic livers. However, limited by the small sample size, they did not find any significant difference between the PoPH and cirrhotic livers, but aromatase expression was increased compared to the normal liver (54). More research is

TABLE 1 | Common clinical and diagnostic findings in patients with Pulmonary Hypertension.

Symptoms	Physical exam	CT findings	TTE findings (57)
Dyspnea on exertion	Accentuated P2	PA diameter ≥ 29 mm	Peak TR velocity > 2.8 m/s
Fatigue	Right sided S3	PA to Ascending Aorta	RV/LV basal diameter > 1.0
Weakness	Right sided S4	diameter ratio of ≥ 1	Flattening of interventricular septum
Orthopnea	TR Murmur	RA Dilation	RV outflow Doppler acceleration
Near-syncope	Pulmonic Insufficiency Murmur	RV Dilation	< 105 msec
Syncope	Jugular Venous Distension		Early diastolic pulmonary regurgitation
Palpitations	Ascites		velocity > 2.2 m/s
Chest Pain	Lower Extremity Edema		PA diameter > 25 mm
	Cyanosis		IVC diameter > 21 mm
			RA end-systole area > 18 cm ²

TR, Tricuspid Regurgitation; PA, Pulmonary Artery; RA, Right Atrium; RV, Right Ventricle; LV, Left Ventricle; IVC, inferior vena cava.

needed to explore the links between altered estrogen metabolism and the development of PoPH in patients with liver disease.

SCREENING AND DIAGNOSIS OF PORTOPULMONARY HYPERTENSION

Screening for PoPH should begin with a detailed history and physical exam (**Figure 4**). Common symptoms of PoPH include unexplained dyspnea on exertion, fatigue, weakness, orthopnea, near-syncope, syncope, palpitations, and chest pain, with dyspnea being by far the most common. On physical exam, these patients often have an accentuated pulmonic component of the second heart sound (P2), a right sided S3, and a right sided S4. Additionally, the presence of a tricuspid regurgitant murmur or a pulmonic insufficiency murmur can be heard in 40% and 13% of patients, respectively (55). Other signs of right heart failure, including jugular venous distention, ascites, and bilateral lower extremity edema, may also be seen (56). Additional history looking for other known World Health Organization (WHO) Group 1 PAH risk factors should be taken. This includes use of drugs such as methamphetamines, use of dietary supplements, prior history of chemotherapy, and history of autoimmune diseases. Clinicians should also review the medical history for any other conditions known to be associated with PH, such as left heart disease, chronic lung disease, obstructive sleep apnea, venous thromboembolism, and hematologic disorders. If the history and physical arouses any suspicion for PoPH, the patient should undergo an echocardiogram.

Laboratory workup should include serological screening tests for connective tissue diseases, hepatitis and HIV. Additional evaluation with pulmonary function tests (including spirometry, lung volumes, diffusion capacity, and a 6 min walk test) and a CT scan of the chest should be obtained to screen for the presence of significant lung disease, and to establish a baseline which can then be used to track disease progression (11, 57). A ventilation/perfusion (V/Q) scan should also be done to screen for the presence of chronic thromboembolic pulmonary hypertension (CTEPH) (especially in the setting of portal vein thrombosis). See **Table 1** for more symptoms, physical exam and imaging findings.

Transthoracic echocardiography (TTE) is the best screening test for PoPH, as it is for other forms of PAH. All patients with portal hypertension should be screened for PoPH with a

TTE. A list of the most common echocardiographic findings that are seen in patients with pulmonary hypertension can be seen in **Table 1**. Although traditionally used throughout the literature to screen for the presence of pulmonary hypertension, right ventricular systolic pressure (RVSP) calculations using the simplified Bernoulli equation are prone to error, stemming from the difficulty in accurately assessing a patient's right atrial pressure. In their analysis of echocardiographic screen of patients undergoing liver transplant evaluation, Krowka et al. demonstrated only moderate correlation between the RVSP and the PA systolic pressures that were measured on right heart catheterization (RHC) (19). Moreover, this discrepancy increased with increasing values of RVSP. Likewise, cutoff values of RVSP used for ruling out the presence of PoPH have differed between studies, and the ideal threshold value for a positive screen has been a matter of debate. As such, it has been suggested that the use of RVSP should be replaced by continuous wave Doppler measurements of peak tricuspid regurgitant velocity, with cutoff values of $2.8 \text{ m}\cdot\text{s}^{-1}$ for intermediate and $3.4 \text{ m}\cdot\text{s}^{-1}$ for high probability for the presence of PH (58). Issues with calculating RVSP notwithstanding, current guidelines from the American Association for the Study of Liver Diseases (AASLD) recommend evaluation with right heart catheterization in all patients being considered for LT with an RVSP ≥ 45 mmHg on screening echocardiogram (59). The International Liver Transplant Society (ILTS) guidelines recommend a RHC in patients with RVSP > 50 mmHg and/or evidence of right ventricular (RV) hypertrophy or dysfunction on TTE (60). Clinicians should understand that when lower RVSP cutoffs are used, the sensitivity will approach 100%, but the specificity will decrease, resulting in a significant number of false positives (17). However, given the high mortality associated with severe PoPH and LT, performing an unnecessary RHC is preferable to “missing” PoPH. In practice, all patients in whom there are TTE findings suspicious for PoPH (elevated RVSP or signs of RV dysfunction) should undergo RHC.

RIGHT HEART CATHETERIZATION HEMODYNAMICS INTERPRETATION IN PoPH

The diagnosis of PoPH includes hemodynamic evaluation of both the pulmonary and hepatic circulations. Portal hypertension can be challenging to diagnose non-invasively. Esophageal varices

	<u>Mean PA Pressure</u>	<u>Cardiac Output</u>	<u>Pulmonary Vascular Resistance</u>	<u>Pulmonary Arterial Wedge Pressure</u>
Hyperdynamic State	↑	↑	↓	↓
Volume Overload	↑	↑	↑/N	↑
Portopulmonary Hypertension	↑	↓/N	↑	↓/N

FIGURE 5 | Right heart catheterization profiles in patients with elevated mean PA pressures. Up arrow, increased; Down arrow, decreased; N, normal.

and skin collaterals are suggestive of portal hypertension. On ultrasound, decreased portal vein flow velocity, and portal vein biphasic or flow reversal are suggestive of portal hypertension (61). A hepatic venous pressure gradient (HVPG) (wedge hepatic vein pressure [WHVP]—free hepatic vein pressure [FHVP]) of >5 mmHg is diagnostic of sinusoidal portal hypertension (62).

On RHC, the patient must have both an elevated mPAP and PVR and a normal pulmonary artery wedge pressure (PAWP) for a positive diagnosis of PoPH. During the most recent World Symposium on Pulmonary Hypertension, the hemodynamic criteria for the diagnosis of pre-capillary pulmonary hypertension (which includes WHO Group 1) were revised. With this change, patients with a mPAP > 20 mmHg (previously > 25 mmHg), a PVR of ≥ 3 Wood units (WU) (≥ 240 dynes), and a PAWP < 15 mmHg meet the diagnostic criteria for the presence of PAH (1).

Careful evaluation of a patient's hemodynamic profile on RHC is crucial in order to distinguish PoPH from other clinical conditions that are common in patients with liver disease that can also result in elevated mPAPs. These conditions can be classified into three broad categories: hyperdynamic state, volume overload and true portopulmonary hypertension (63). See **Figure 5** for comparisons between these hemodynamic profiles.

Patients with decompensated liver disease may present with an elevated mPAP, PAWP, and PVR. This may be due to a combination of volume overload, hyperdynamic state, and PoPH (perhaps all three). Volume overload is managed with careful diuresis, as patients with liver disease are prone to kidney injury (64). A hyperdynamic state can be intrinsic to liver disease, but may also be due to other treatable medical comorbidities (such as anemia, arteriovenous shunts, and hyperthyroidism) (65–67). This common clinical scenario speaks to the challenge of diagnosing and treating PoPH, which

is best done at expert centers which are most familiar with these complicated patients.

TREATMENT OF PoPH

The importance of screening for PoPH was established by Krowka et al. in a retrospective review of case reports and case series, which looked at the relationship between the presence of untreated PoPH and cardiopulmonary related mortality in patients undergoing LT (7). In this seminal work, Krowka et al. examined 43 patients with PoPH (confirmed by RHC) who were not on medical treatment and underwent LT. The patients were grouped based on severity of pulmonary hypertension. Patients with a mPAP between 25 and 35 mmHg were considered to have mild disease, those with a mPAP between 35 and 50 mmHg were considered moderate, and those with a mPAP >50 mmHg were considered severe. In these groups, 100% of the patients with severe and 50% of the patients with moderate PoPH died due to cardiopulmonary related events (the majority died during the transplantation hospitalization). No mortality was reported among the patients with mild PoPH (mPAP < 35 mmHg). As a result of this study, and others, routine screening for the presence of pulmonary hypertension is now recommended in all patients undergoing evaluation for LT.

After this landmark study established that PoPH patients with higher mPAP are at high risk of mortality with LT, it prompted the question of whether medical therapy could mitigate some of those risks. Treatment of PoPH prior to or instead of liver transplantation remains challenging and is fraught with a paucity of data and large knowledge gaps. Randomized controlled trials of pulmonary hypertension specific medical therapies for patients with portopulmonary hypertension have been limited by concerns regarding adverse hepatic effects of PAH specific medications (68). We will review select clinical trials of PAH-specific therapy in PoPH, as well as the indications and contraindications for liver transplantation.

Retrospective data on treatment of PoPH is quite heterogeneous, with some studies showing improvement in hemodynamics and outcomes, and others showing no treatment effect. There is higher quality data from prospective non-randomized studies, from which a few themes emerge. A recently published prospective cohort study from the French Pulmonary Hypertension Registry (FPHR) examined data on 637 patients with PoPH, 90% of whom were started on PAH-specific therapies (12). These patients experienced significant improved WHO functional class, 6-min walk distance (6MWD) and hemodynamic parameters. The ANGEL study is a recent single arm prospective trial of ambrisentan in 31 patients with PoPH (69). This trial showed improved hemodynamics (PVR, mPAP, cardiac index [CI]) and functional class, but no change in 6MWD. From these prospective studies, it is reasonable to conclude that PAH directed therapy has favorable effects on hemodynamics and functional outcomes, but the effects on mortality and candidacy for LT are unclear.

As previously stated, the vast majority of randomized controlled trials of PAH-specific therapies in PAH patients have excluded patients with PoPH. One notable exception is the PATENT-1 trial by Ghofrani et al. (phase 3, double-blind, randomized controlled trial), which randomized 443 patients with various etiologies of WHO group 1 PAH to receive either placebo or two different doses of riociguat (70). This trial is significant in that it included 13 patients with PoPH, 11 of whom received the highest dose of study medication (2.5 mg daily). The primary end point of the PATENT-1 trial was 6MWD, which was significantly improved in the riociguat 2.5 mg group. Secondary hemodynamic (PVR, mPAP, and CO) and functional (WHO functional class, Borg dyspnea scale and time to clinical worsening) endpoints were also significantly improved in the riociguat 2.5 mg daily group. The PATENT-2 trial was an extension of the PATENT-1 study, in which all eligible patients from the first study were given riociguat and followed for 2 years. PATENT-2 showed sustained improvements in functional class and 6MWD (71). A recently published subgroup analysis of the 13 patients with PoPH who were enrolled in PATENT-1 and 2 showed similar improvements in WHO functional class, 6MWD and PVR (72). Importantly, riociguat was well-tolerated by the PoPH patients.

The first ever randomized controlled trial of PAH specific therapy in patients with PoPH was published in 2019. The PORTICO study, which randomized 85 patients with PoPH to receive either macitentan or placebo for 12 weeks, found that the macitentan group had a 35% reduction in PVR, the primary endpoint (73). While there were significant improvements in some secondary endpoints (mPAP and CI), there were no significant improvements in WHO functional class and 6MWD. Importantly, macitentan was well-tolerated in this population, with no hepatic side effects.

A notable limitation of PORTICO is that it excluded patients with Child-Pugh class C liver disease and patients with a MELD score of >19, which are groups with known poor survival (12).

It is still unclear whether this significant reduction in PVR could translate into more meaningful outcomes for patients

(improved functional capacity, decreased mortality) or alter candidacy for liver transplant. However, a recently published *post-hoc* analysis of the PORTICO study found that patients in the macitentan treatment group had significantly decreased waitlist and perioperative mortality risk categories as compared with the placebo arm (74).

Taken together, these studies show that PAH specific therapies can improve hemodynamics and functional outcomes in patients with PoPH. Importantly, these medications seem to be well-tolerated in patients with liver disease and PoPH. By extension, they can help patients become candidates for LT and potentially live longer. In the absence of PoPH specific treatment guidelines, clinicians should follow the general treatment principles of PAH, with special attention to the unique considerations for patients with PoPH. More research is needed to elucidate the nuances of PAH specific therapies in the PoPH population.

LIVER TRANSPLANTATION

As previously discussed, the severity of PoPH is an established risk factor for liver transplant waitlist mortality (75) and poor perioperative outcomes (7). As such, decisions about candidacy for liver transplantation in patients with PoPH are complex and best done by experienced centers.

The following general principles are recommended by the AASLD (59) for management of patients with liver disease who are being considered for LT. All patients who are undergoing evaluation for LT should be screened for PoPH with transthoracic echocardiography, and patients with RVSP > 45 mmHg should be further evaluated by RHC. Patients with mPAP < 35 mmHg do not require treatment with PAH specific therapies and can proceed with further evaluation for LT. Patients with mPAP > 35 mmHg should be evaluated by an experienced PH specialist and be considered for PAH specific therapy. If the mPAP can be lowered to <35 mmHg and the PVR can be lowered to <5 WU with medical therapy, LT can be pursued.

For some patients with mild liver disease and PoPH, their higher risk for waitlist mortality is not accurately captured by the MELD score (76). LT candidates with PoPH are eligible for MELD exception points if the following criteria are met: initial mPAP > 35 and PVR > 3 WU; patients are treated with PAH specific therapy; and post-treatment mPAP is lowered to < 35 mmHg (76).

Giving MELD exception points to patients with PoPH is controversial for a number of reasons. The U.S. based Organ Procurement and Transplant Network (OPTN) requires a RHC and resubmission of full hemodynamic data every 3 months (76) to maintain the MELD exception points, which is burdensome and often impractical for both patients and institutions. Despite having to submit full hemodynamic data every 3 months, only mPAP < 35 mmHg is used to maintain eligibility MELD exception points. Some patients may achieve a lower PVR with PAH directed therapy, but continue to have an elevated mPAP from volume overload and/or a high flow state (60). In an analysis of patients with PoPH on the LT waitlist, initial PVR

was an independent predictor of waitlist mortality, while pre- and post-treatment mPAP were not (75). This suggests that a broader hemodynamic picture should be used to maintain MELD exception points eligibility.

In one analysis of 155 patients on the LT waitlist with approved PoPH MELD exception points, less than half of the patients actually met the standardized OPTN MELD exception criteria. Furthermore, almost one third of patients in this study lacked sufficient hemodynamic data to diagnose PoPH or had hemodynamic data that was inconsistent with PoPH (77). These researchers found that overall mortality was higher for patients given PoPH MELD exception points (as compared to patients listed without exception points), which shows the need for more research into methods of risk stratification for patients with PoPH.

In the era of the AASLD/ILTS guidelines and MELD exception points, data on the natural course of PoPH after LT is limited. In one retrospective cohort study, hemodynamics and survival data for 35 patients with PoPH were examined (78). After LT, 6 month, 1 year, and 3 year survival rates were 80, 77, and 77%, respectively. Of the 27/35 patients who survived more than 6 months after LT, all were able to be weaned from intravenous epoprostenol. Furthermore, most of the patients had improvement in their hemodynamics after LT, with 30% having a mPAP < 25 mmHg at last follow up. This data suggests that carefully selected patients with PoPH who undergo LT will have significant improvement in their hemodynamics.

CHALLENGES IN THE DIAGNOSIS AND MANAGEMENT OF PoPH

There are a number of barriers to the effective diagnosis and treatment of patients with PoPH. As previously discussed, accurate diagnosis of PoPH is challenging, and the differential for an elevated mPAP in the setting of liver disease is broad. There is only one randomized controlled trial for PAH specific therapy in patients with PoPH. Advanced liver disease symptoms have significant overlap with the side effects of PAH specific therapies (e.g., nausea, fatigue, and fluid overload), which can make the initiation and maintenance of treatment challenging (67).

Many of the drugs used to treat PoPH are metabolized in the liver, and drug metabolism is altered by liver disease and other medications that patients may be taking (67). Additionally, patients may have hepatic encephalopathy or lack the social support needed to safely administer parenteral therapy or take medications multiple times per day (67). More generally, many of the treatment barriers that apply to all patients with PAH apply to patients with PoPH: late referrals, lack of an expert center in

close proximity to where the patient lives, and insurance issues. There is significant variability in the screening and management practices of PoPH, discordance between published guidelines and actual practice patterns, and disagreements about the role of LT in PoPH (79).

CONCLUSIONS AND AREAS OF FUTURE FOCUS

PoPH remains a challenging disease entity with many facets in need of further research. More research is needed into which types of PAH specific therapies will benefit patients with varying severities of PoPH. The first randomized control trial in PoPH was recently published, which showed that macitentan significantly lowered PVR in patients with PoPH. It unfortunately remains unclear if hemodynamics improvements translate into successful LT or substantial survival benefit. While it is clear that patients with severe disease have high mortality with or without LT, their long-term response to treatment is unknown. It is unknown if mild PoPH needs to be treated or given MELD exception points, and the risk for progression to more severe disease is unknown.

Pre-operative risk stratification of patients prior to LT remains a challenge, as the traditional cut off of a mPAP of 35 mmHg is somewhat arbitrary and is based on a small retrospective review of case reports and series. There are no published guidelines on initiation of PAH specific therapies in PoPH, or whether to continue PoPH treatment post-LT. Finally, there is no centralized national registry of PoPH patients who undergo LT; this would serve as a foundation for focusing on the aforementioned areas of need.

While there are indeed many barriers to and challenges in caring for patients with PoPH, it remains a field with many areas of active investigation. As the mechanisms and treatment continue to be explored, we are confident that care for these complex patients will continue to improve.

AUTHOR CONTRIBUTIONS

CT and VG: primarily responsible for writing the manuscript. VJ and SS: editing and revision and mentoring. All authors contributed to the article and approved the submitted version.

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Epidemiology, Pathogenesis, and Clinical Approach in Group 5 Pulmonary Hypertension

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Pulmonary hypertension (PH) is recognized to be associated with a number of comorbid conditions. Based on these associations, PH is classified into 5 groups, considering common pathophysiologic drivers of disease, histopathologic features, clinical manifestations and course, and response to PH therapy. However, in some of these associated conditions, these characteristics are less well-understood. These include, among others, conditions commonly encountered in clinical practice such as sarcoidosis, sickle cell disease, myeloproliferative disorders, and chronic kidney disease/end stage renal disease. PH in these contexts presents a significant challenge to clinicians with respect to disease management. The most recent updated clinical classification schemata from the 6th World Symposium on PH classifies such entities in Group 5, highlighting the often unclear and/or multifactorial nature of PH. An in-depth review of the state of the science of Group 5 PH with respect to epidemiology, pathogenesis, and management is provided. Where applicable, future directions with respect to research needed to enhance understanding of the clinical course of these entities is also discussed.

Keywords: multifactorial, pulmonary hypertension, Group 5, hematologic, sickle cell, sarcoidosis, metabolic, CKD - chronic kidney disease

INTRODUCTION

Pulmonary hypertension (PH) is characterized by elevated mean pulmonary artery pressure (mPAP) of >20 mmHg as determined by right heart catheterization (RHC). PH is divided into 5 groups based on the underlying mechanism using the original World Health Organization classification system framework. In 2019, the proceedings of the World Symposium on Pulmonary Hypertension published the updated and revised classification based on the 6th World Symposium on PH (1). According to the new classification, causes of PH include pulmonary arterial hypertension (PAH, Group 1 PH), pulmonary venous hypertension due to elevated filling pressure of the left heart (Group 2 PH), PH due to chronic pulmonary disease or hypoxemia (Group 3 PH), chronic thromboembolic disease (Group 4), whereas PH that develops due to multiple or unclear mechanisms is referred to as Group 5 PH (Table 1). Within Group 5, several clinical disorders are implicated in association with the development of PH (Table 2). The most commonly seen clinical entities in Group 5 PH (Groups 5.1–5.3) are covered in detail: PH in the setting of hematologic disorders, sarcoidosis, and chronic renal failure. A brief overview of some of the remaining, less common but clinically relevant entities in Group 5 is also provided. Complex congenital heart disease (Group 5.4) is outside the scope of this manuscript.

TABLE 1 | Current clinical classification of pulmonary hypertension (1).

Group	Description
1	Pulmonary Arterial Hypertension 1.1 Idiopathic PAH 1.2 Heritable PAH 1.3 Drug- and toxin-induced PAH 1.4 PAH associated with: 1.4.1 Connective tissue disease 1.4.2 HIV infection 1.4.3 Portal hypertension 1.4.4 Congenital heart disease 1.4.5 Schistosomiasis 1.5 PAH long-term responders to CCB 1.6 PAH with overt signs of venous/capillaries (PVOD/PCH) involvement 1.7 Persistent PH of the Newborn syndrome
2	PH Due to Left Heart Disease 2.1 PH due to heart failure with preserved EF 2.2 PH due to heart failure with reduced EF 2.3 Valvular heart disease 2.4 Congenital/acquired CV conditions leading to post-capillary PH
3	PH Due to Lung Diseases and/or Hypoxia 3.1 Obstructive lung disease 3.2 Restrictive lung disease 3.3 Other lung disease with mixed restrictive/obstructive pattern 3.4 Hypoxia without lung disease 3.5 Developmental lung disorders
4	PH Due to Pulmonary Artery Obstruction 4.1 Chronic thromboembolic PH 4.2 Other pulmonary artery obstructions
5	PH With Unclear and/or Multifactorial Mechanisms 5.1 Hematologic disorders 5.2 Systemic disorders 5.3 Others 5.4 Complex congenital heart disease

Considering the true upper limits of normal mPAP, the 6th World Symposium on PH introduced revised hemodynamic definitions, affirming PH in those with mPAP >20 mmHg as opposed to mPAP \geq 25 mmHg as previously defined. A pulmonary vascular resistance (PVR) of at least 3 Wood Units was further mandatory to define pre-capillary pulmonary hypertension. These changes impact identification and management of all groups of PH, including group 5 PH. With updated review and discussion of the literature, the proceedings outlined additional changes specifically impacting group 5 PH, including removal of both splenectomy and thyroid disease as specific subtypes of PH (asserting these conditions as risk factors) and reclassification of lymphangioleiomyomatosis into Group 3 PH (1).

PULMONARY HYPERTENSION IN HEMATOLOGIC DISORDERS

Chronic Hemolytic Anemias

Prevalence

PH is increasingly recognized as a major source of morbidity and mortality in patients with chronic hemolysis, most notably

TABLE 2 | Subclassifications of group 5 pulmonary hypertension: PH with unclear and/or multifactorial mechanisms (1).

5.1 Hematologic disorders	Hemolytic anemias Myeloproliferative disorders
5.2 Systemic and metabolic disorders	Pulmonary Langerhans cell histiocytosis Gaucher disease Glycogen storage disease Neurofibromatosis Sarcoidosis
5.3. Others	Chronic renal failure with or without hemodialysis Fibrosing mediastinitis
5.4 Complex congenital heart diseases	Segmental pulmonary hypertension <ul style="list-style-type: none"> Isolated pulmonary artery of ductal origin Absent pulmonary artery Pulmonary atresia with ventricular septal defect and major aorto-pulmonary collateral arteries Hemitruncus Other Single ventricle <ul style="list-style-type: none"> Unoperated Operated Scimitar syndrome

in the context of sickle cell disease. The prevalence of PH in hemolytic anemia is variable depending upon the diagnostic criteria, methods used to diagnose PH, and population studied. In sickle cell disease (SCD, hemoglobin SS), 30–40% of patients have evidence of elevated pulmonary artery pressures based on tricuspid regurgitant jet velocity (TRV) \geq 2.5 m/s on Doppler echocardiography. However, only 6–10.5% of patients have mPAP of \geq 25 mmHg on invasive hemodynamic evaluation by RHC (2–5). This reflects the fact that estimation of pulmonary artery pressure is often inaccurate by echocardiogram. Additionally, elevated pulmonary artery pressures can be related to high cardiac output (due to anemia) and not always secondary to increased pulmonary vascular resistance (PVR). Using the new definition of PH (mPAP >20 mmHg), the prevalence of PH in SCD is probably higher than that reported in these studies.

PH has also been reported in association with other hemolytic anemias. In a study of 110 patients with β -thalassemia intermedia, 59% of patients had elevated peak systolic tricuspid gradient >30 mmHg on Doppler echocardiography, suggestive of PH. Only 6 patients had RHC and were found to have severe pre-capillary PH. All patients in this study had normal left ventricular function and high cardiac output (6). The prevalence of PH by echocardiography was even higher (75%) in a small study of 35 patients with homozygous β -thalassemia (7). More recently, Derchi et al. evaluated the prevalence of PH in a large multicenter cross-sectional study of 1,309 patients with β -thalassemia in Italy. Nine percent of patients had TRV >3 m/s; patients in the “PH likely” group with TRV \geq 3.2 m/s underwent RHC. The prevalence of RHC-confirmed PH was 2.1%. Increased age and splenectomy were independent risk factors for PH in this study (8).

Notably, the prevalence of PH in patients with hereditary spherocytosis is much lower than reported with SCD and thalassemia. In two retrospective studies, no patients had

a TRV of ≥ 2.8 m/s on echocardiography to suggest PH (9, 10). Nonetheless, PH has been described in hereditary spherocytosis and stomatocytosis, with most cases related to chronic thromboembolic pulmonary hypertension (CTEPH) following splenectomy (11).

Pathogenesis

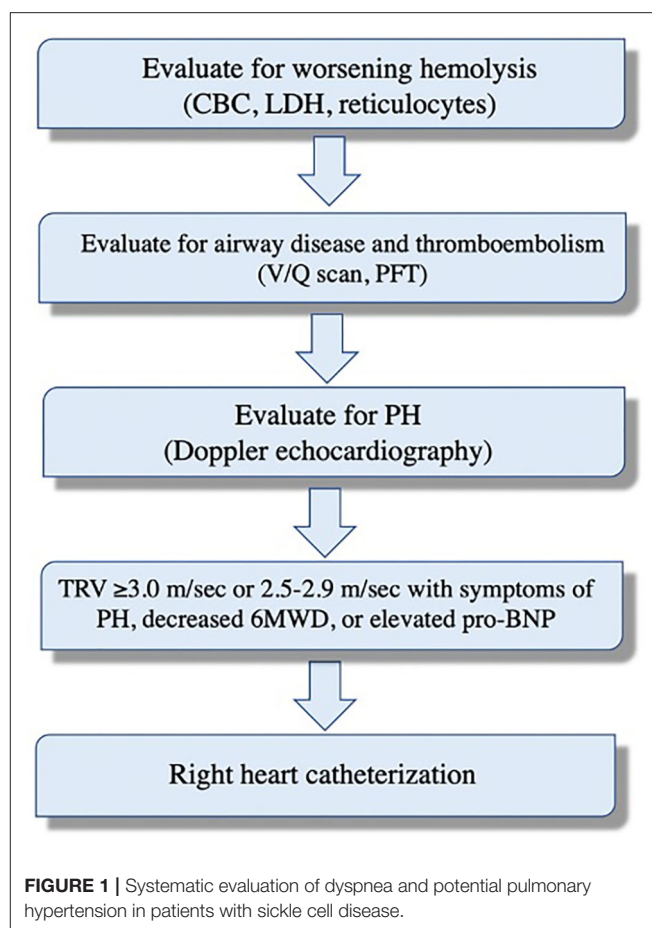
The pathogenesis of PH in chronic hemolysis is multifaceted and is most extensively studied in SCD. The vasomotor tone of the pulmonary circulation is normally regulated by nitric oxide (NO), a powerful vasodilator and modulator of endothelial proliferation. NO is produced by the pulmonary endothelium in response to shear stress on the surface of endothelial cells (12). The enzyme NO synthase catalyzes cleavage of the terminal amino group from the amino acid L-arginine, producing NO. Hemolysis causes the release of arginase-1, which depletes L-arginine eliminating the NO precursor (13). Additionally, studies have shown that free hemoglobin produced by intravascular hemolysis in SCD consumes NO (14, 15). Impaired NO production is also related to hemolysis-induced generation of asymmetric dimethylarginine, an endogenous NO synthase inhibitor (16). Production of heme due to oxidation of free hemoglobin also activates Toll-like receptor 4 promoting vaso-occlusive events (17, 18). Interestingly, a growing body of literature indicates a possible role for free hemoglobin in the pathogenesis of PAH in patients with no known hemolysis. Rafikova et al. found that free hemoglobin levels were significantly higher in patients with known PAH compared to patients with no PAH. Furthermore, the severity of PAH correlated with higher cell-free hemoglobin levels. In an animal model of PAH, inhibition of heme translocation using sulfasalazine prevented the development of PH (19).

Endothelin-1 (ET-1), a potent vasoconstrictor of pulmonary vascular bed, plays a role in the imbalance between vasodilators and vasoconstrictors in SCD. Plasma and urine levels of ET-1 are elevated in patients with SCD compared with matched controls (20). Furthermore, Phelan et al. showed that exposure of cultured human endothelial cells to previously sickled erythrocytes results in enhanced gene expression of ET-1 (21).

Hypercoagulability is another important factor that adds to pulmonary endothelial dysfunction. A hypercoagulable state results from reduced NO, asplenia (functional or surgical), and activation of thrombotic factors (tissue factor, platelets, and thrombin) in SCD (22, 23). This is supported by findings of pulmonary thromboemboli in 38–80% of postmortem lung examinations of deceased SCD patients (24, 25).

Proliferative arteriopathy seen in SCD-PH implies possible contribution of angiogenic factors to the pathogenesis. A recent case-control study of children with β -thalassemia major demonstrated a significantly higher serum level of vascular endothelial growth factor in patients with PH than those without PH or matched healthy children (26).

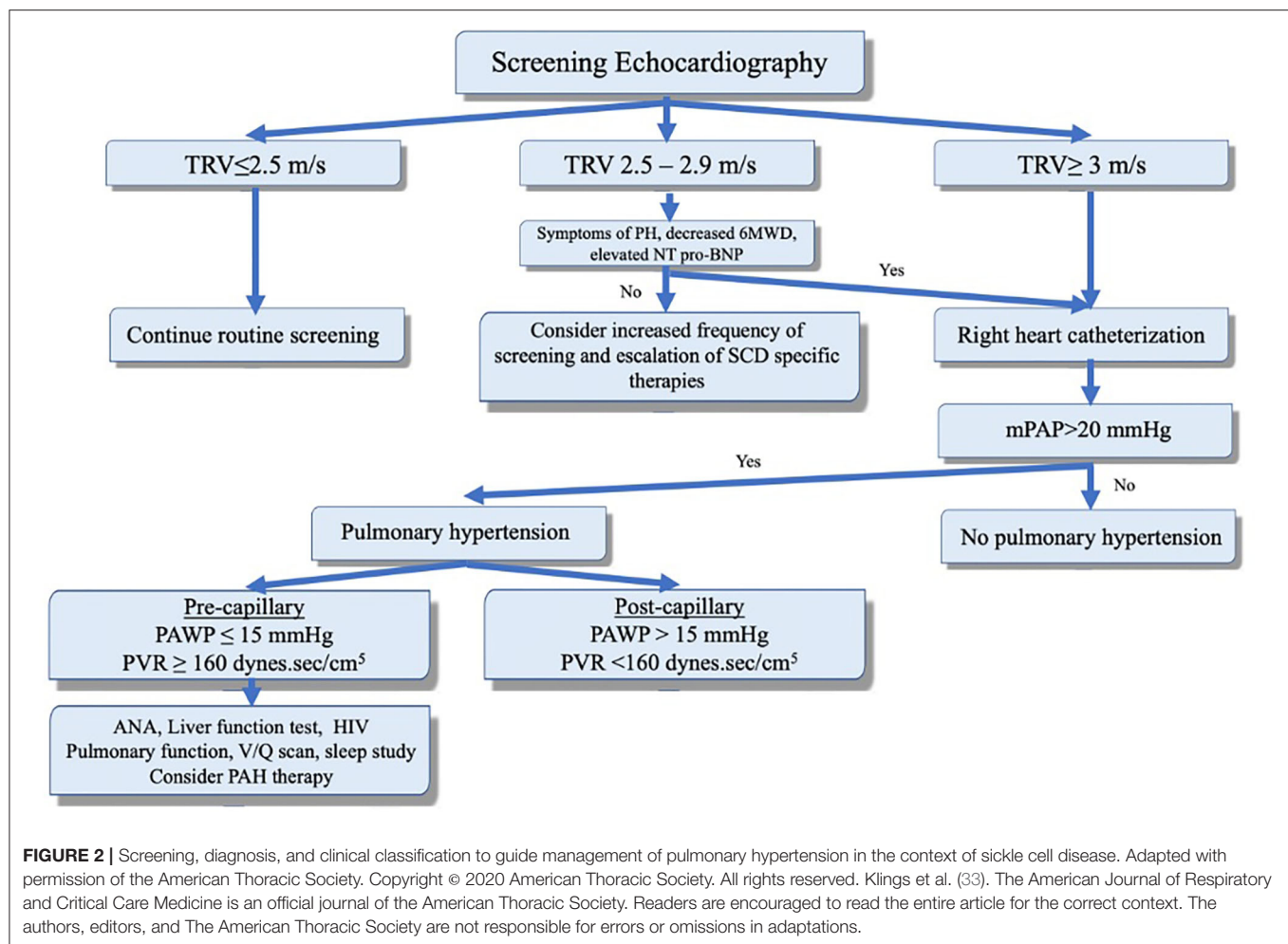
Finally, a significant number of patients with SCD have been found to have restrictive cardiomyopathy and diastolic dysfunction (27). In fact, 50% of patients with SCD and PH have pulmonary artery wedge pressure (PAWP) of >15 mmHg, consistent with pulmonary venous hypertension (i.e.,



post-capillary PH) (3, 5, 28). Vaso-occlusive events, systemic hypertension, iron overload (due to transfusions) and end-organ damage (e.g., renal failure) in patients with chronic hemolysis contribute to diastolic left ventricular dysfunction as well (29). Collectively, all these factors increase the risk of developing PH in patients with SCD and other hemolytic anemias.

Evaluation

Pulmonary symptoms are common in patients with chronic hemolysis and often are multifactorial. Patients with SCD, for example, may have recurrent chest pains, fatigue, and chronic shortness of breath. These symptoms can be related to anemia, acute chest syndrome, obstructive airway disease, thromboembolism, and/or PH (30–32). Therefore, these should be systematically evaluated (Figure 1). Asymptomatic patients also remain at greater risk of developing PH; therefore, screening with Doppler echocardiography is recommended. The American Thoracic Society guidelines recommend screening patients with SCD with Doppler echocardiography every 1–3 years to assess mortality risk. Importantly, echocardiography should not be performed within 4 weeks of acute chest syndrome or within 2 weeks of acute vaso-occlusive crisis as these acutely increase the pulmonary artery systolic pressure (PASP) due to hypoxemia and worsening anemia (33). There are no screening guidelines



for patients with other hemolytic diseases. Nonetheless, it is reasonable to obtain echocardiography on high-risk patients (e.g., severe hemolysis, asplenia, prior venous thromboembolism, and iron overload), especially those with unexplained pulmonary symptoms (34).

PH is suspected if the TRV on echocardiography is >2.5 m/sec. However, $\text{TRV} \geq 2.5$ m/s has a low positive predictive value as only 31% of such patients had $\text{mPAP} \geq 25$ mmHg on RHC (27). A $\text{TRV} \geq 3$ m/s is 3 standard deviations above the population mean and is more specific (PH confirmed in 66–77% of patients on RHC) (4, 5). A TRV between 2.5 and 3 m/s combined with either an NT-proBNP > 164.5 pg/ml or 6-min walk distance of < 333 m increases the likelihood of PH and warrants further evaluation and RHC (Figure 2) (3, 5). PASP is commonly estimated indirectly by measuring the right ventricular systolic pressure which (in the absence of pulmonic stenosis) are equivalent. Using the modified Bernoulli equation ($P = 4V^2$), the PASP is calculated from the TRV. This means that minor variations in measurement of TRV can result in significant differences in estimated pulmonary pressures. Furthermore, even when accurate, elevated PASP can be related to high cardiac output often seen in patients with anemia. On the other hand,

PH can be missed on echocardiography as the TRV is not always measurable (35). Thus, RHC remains the test of choice to confirm the presence of PH and assess the PAWP, cardiac output, and calculated PVR. It is worth mentioning that reduced blood viscosity and high cardiac output (due to anemia) result in a lower baseline PVR than healthy subjects. Thus, PVR of >2 Wood units is considered elevated in these patients (3–5).

Blood biomarkers can also be helpful in the evaluation of PH in chronic hemolytic anemia. Lactate dehydrogenase is a commonly used marker for hemolysis has been shown to correlate with NO resistance and severity of PH (36). NT-proBNP is another useful biomarker for PH and right ventricular dysfunction. In a study by Machado et al., elevated NT-proBNP (>160 pg/mL) in SCD patients correlated with TRV and was predictive of PH. Additionally, elevated NT-proBNP was an independent predictor of mortality (37).

Treatment

Management of PH in hemolytic anemias includes general measures such as supplemental oxygen to reverse hypoxemia and prevent its deleterious effect on the pulmonary vascular tone. Diuretics are used to treat volume overload but should

be used cautiously due to risk of sickling in patients with SCD. Lifelong anticoagulation is indicated in SCD patients with PH secondary to CTEPH. Although surgical pulmonary thromboendarterectomy (PTE) can be curative in CTEPH, it is undoubtedly more challenging in SCD patients due to the increased risk of sickling and occlusive crises during cardiopulmonary bypass. The best perioperative approach in these patients remains controversial. Avoidance of hypoxemia, hypothermia, and acidosis reduces the risk of perioperative complications. Additionally, exchange transfusion to reduce hemoglobin S to <20% prior to surgery is advised (38, 39). This aggressive transfusion regimen, however, has not been shown to reduce perioperative complications in patients with SCD (40). Inoperable patients are considered for balloon pulmonary angioplasty (BPA). In a recent study of PH in SCD, three patients with CTEPH underwent BPA with significant reduction in PVR in 2 patients and normalization of mPAP in the third patient (41).

Optimizing treatment of the underlying hemolytic disease is of paramount importance. Hydroxyurea is a myelosuppressive agent that has been shown to decrease sickle cell hemoglobin polymerization, reducing hemolysis and frequency of acute chest syndrome, and vaso-occlusive crises. Furthermore, hydroxyurea decreases hospitalization and mortality in SCD patients with HbSS and is currently recommended for patients with more than one episode of acute chest syndrome or >3 vaso-occlusive crises per year (33, 42–44). The risk of developing PH is also reduced with hydroxyurea in patients with thalassemia (45). Chronic transfusion is recommended in SCD patients who fail to respond or cannot tolerate hydroxyurea. This strategy is associated with improvement in pulmonary vascular changes in SCD patients (46). Similarly, adherence to transfusion and iron chelation therapy in patients with thalassemia prevented PH (47, 48).

SCD patients who remain symptomatic despite optimization of their SCD therapy (i.e., hydroxyurea, transfusion) are considered for PH-targeted therapy. This is usually reserved for patients with moderate to severe (based on functional class) RHC-confirmed pre-capillary PH with relatively reduced cardiac output. In general, endothelin receptor antagonists (ERAs) are recommended in patients with moderate PH (Functional class 2–3). Bosentan, an endothelin receptor antagonist, was evaluated in two randomized placebo-controlled trials of SCD patients with RHC-proven pre- and post-capillary PH. Both studies were terminated early due to slow enrollment but modest improvement in hemodynamic parameters was observed. Efficacy endpoints were not analyzed to draw definitive conclusions but bosentan was well-tolerated (49). Patients with more severe PH (class IV functional status, evidence of right-sided heart failure, or reduced cardiac output) can be treated with prostanoids. The efficacy of prostacyclins has been evaluated in a small retrospective study of 11 patients who received treprostinil or epoprostenol and showed improvement in right ventricular systolic pressure on echocardiography (50). Patients who need such therapy should be referred to centers with expertise in managing PH in SCD patients due to concerns of complications (e.g., high-cardiac output pulmonary edema, line thrombosis). Phosphodiesterase type 5 inhibitors (PDE-5Is) should generally be avoided in this population due to risk of

increased hospitalizations for painful crises noted in a study of sildenafil (51). Riociguat is a soluble guanylate cyclase stimulator that does not rely on NO and, therefore, is an attractive agent for SCD-PH. Riociguat is currently approved for treatment of Group 1 PAH and Group 4 CTEPH. Weir et al. recently reported in a small case series tolerability of riociguat in four of six patients with sickle related CTEPH. Additionally, riociguat was associated with significant improvement in exercise capacity (mean 6MWD improvement of 56.8 m), functional class, NT-proBNP, and RVSP in 3 patients. Vaso-occlusive crisis developed in only one patient who was not on hydroxyurea (52). A phase 2, multi-multi-center, randomized, double-blind, placebo-controlled, study is now underway to evaluate the effectiveness and safety of riociguat in patients with sickle cell disease.

Myeloproliferative Disorders

Prevalence

Myeloproliferative disorders (MPDs) are characterized by clonal expansion of the multipotent hematopoietic progenitor cell with overproduction of one of the blood elements (i.e., erythrocytes, leukocytes, or platelets). The prevalence of PH in patients with MPDs has been evaluated in multiple small studies. Most of these studies used Doppler echocardiography (PASP > 35 mmHg) to diagnose PH, with an estimated prevalence of 13–48% of PH in MPDs (53–56). Interestingly, in a larger study of 103 patients with MPDs, the prevalence of PH was much lower with only 5 patients found to have PH (57). Among MPDs, patients with chronic myeloid leukemia, polycythemia vera, essential thrombocytosis, and myelofibrosis are particularly at higher risk of developing PH. The presence of PH is associated with poor prognosis in MPDs.

Pathogenesis

PH in MPDs has two main phenotypes: chronic thromboembolic (CTEPH) and proliferative pulmonary arteriopathy. Thrombotic events (both venous and arterial) and microcirculatory disturbances are common in polycythemia vera and essential thrombocytosis patients. In a large study of 1,213 patients with polycythemia vera, 41% developed thrombotic complications (58). This translates into higher risk of CTEPH. In a retrospective small study of 10 patients with MPDs and RHC-confirmed PH, six patients were diagnosed with CTEPH at the time of MPD diagnosis. CTEPH was strongly associated with elevated hematocrit (59). Presence of cardiovascular comorbidities, history of thrombosis or splenectomy and increased age further increase the risk of thrombosis in these patients (60–62).

High hematocrit in polycythemia vera is associated with hypercoagulability as demonstrated by the progressive increase in blood viscosity that parallels higher hematocrit values (63). The major impact of hematocrit on blood viscosity is in the venous (low-shear) circulation as it decreases blood flow and increases the risk of venous thrombosis. In the arterial circulation, under high-shear, platelets are displaced to the periphery by the large red cell mass and come in contact with endothelial cells, causing platelets activation (release of thromboxane A₂, platelet-derived growth factor, vascular endothelial growth factor) and thrombosis (64, 65). The abnormalities of erythrocytes and

platelets in MPDs also contribute to the aggregation of red blood cells and disturbed blood flow. Furthermore, chronically activated leukocytes (especially with JAK2 mutation) interact with platelets and endothelial cells and increase the risk of thrombosis (66–69). Splenectomy is used in the treatment of MPDs and can be associated with a hypercoagulable state as well (61). Notably, obstruction of pulmonary arteries in MPDs is not always related to thrombi but can be secondary to tumor emboli or circulating megakaryocytes with subsequent release of vasoactive substances that will further increase the tone in the pulmonary vascular bed (70, 71).

The second PH phenotype in MPDs is pulmonary arteriopathy which can result from different mechanisms. Portopulmonary hypertension may complicate chronic liver disease which is known to occur in MPDs (72). Medication-induced PH, fitting into Group 1 PH, is another potential cause of PH in these patients. Dasatinib, a tyrosine kinase inhibitor commonly used for treatment of chronic myeloid leukemia, has been associated with development of PH which seems to improve after discontinuation (73). Several TKIs other than dasatinib (e.g., bosutinib, ponatinib, lapatinib) have also been implicated in the development of PH. In a small study of 27 patients who received lapatinib, three patients were found to have PH by RHC with resolution of PH in all three after cessation of therapy (74).

Alkylating agents such as cyclophosphamide, melphalan, and busulfan (used for treatment of myelofibrosis) can cause PH with severely reduced diffusing capacity of the lung for carbon monoxide and abnormal findings on computed tomography of the chest (septal thickening, ground glass opacities, and lymph node enlargement) that are consistent with pulmonary veno-occlusive disease. Unlike dasatinib-induced PH, these patients have a poorer prognosis. Extramedullary hematopoiesis may complicate MPDs and result in infiltrative changes in the portal circulation with subsequent development of portal hypertension and portopulmonary hypertension. In cases of lung involvement, extramedullary hematopoiesis may lead to pulmonary vascular occlusion and PH. Proliferative changes in the pulmonary vessels may also occur due to increased circulating vascular endothelial growth factors and enhanced angiogenesis in MPDs (75). Finally, pre-capillary PH can be related to abnormal JAK2 signaling and depletion of NO. This is supported by improvement in PH and NO with JAK2 inhibitors.

Treatment

Since hematologic abnormalities contribute directly to the pulmonary vascular changes seen in MPDs, treatment of the underlying disease is a potential strategy. Indeed, allogeneic hematopoietic stem cell transplant and JAK inhibitors to treat MPDs have been shown to improve PH (76). For example, treatment with ruxolitinib (JAK1/JAK2 inhibitor) in 15 patients with myelofibrosis and PH resulted in improvement in right ventricular function and PASP by echocardiography in 66% of patients and was associated with significant reduction in NT-proBNP and increase in NO level (77). In cases of drug-induced PH, improvement in PH can be expected with discontinuation of the offending agent (especially in dasatinib-induced PH) although this alone may not be enough to reverse PH. When

appropriate, CTEPH is treated with PTE or BPA. Riociguat, a guanylate cyclase stimulator, is associated with improved functional status and hemodynamics in patients with inoperable disease or suboptimal post-PTE hemodynamic outcomes (78, 79). External beam radiotherapy to the thorax has been used to treat extramedullary hematopoiesis in patients with MPDs and PH with improvement in pulmonary artery pressures (80). PAH-specific therapy has not been studied in MPD-associated PH.

PULMONARY HYPERTENSION IN SARCROIDOSIS

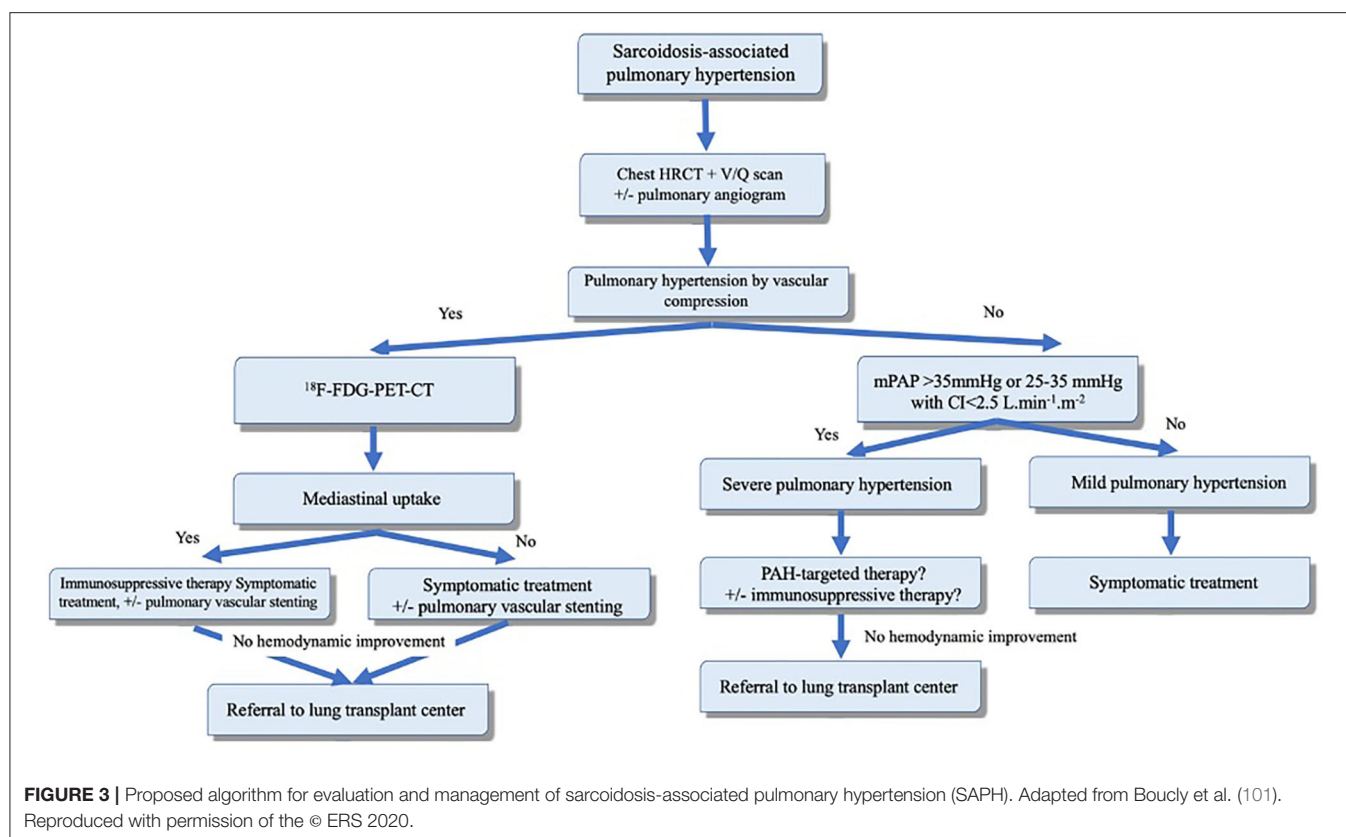
Prevalence

Sarcoidosis is characterized by granulomatous inflammation to an unknown trigger that primarily affects the lungs. The prevalence of sarcoidosis-associated PH (SAPH) is variable. This variation comes from inconsistency in methods used to diagnose SAPH. Autopsy studies have shown pulmonary vascular involvement in ~5% of patients with significant parenchymal lung disease (81). Similar prevalence has been found in studies that evaluated SAPH based on clinical evidence of right ventricular failure (i.e., cor pulmonale) (82). However, studies that used hemodynamic assessment using RHC to diagnose SAPH reported higher prevalence up to 23% (83, 84). The prevalence is substantially higher in patients with pulmonary symptoms and advanced lung disease due to sarcoidosis. In a study of 15 patients with advanced fibrotic sarcoidosis, Emirgil et al. found evidence of PH on RHC in 10 patients (67%) (85). Another study of 363 patients with advanced sarcoidosis (listed for lung transplantation) found evidence of PH on RHC in 74% of these patients (86). Racial differences may significantly impact the course of disease and risk of developing PH. For example, in a recent large prospective study of 512 patients, predominantly Caucasian, with sarcoidosis, SAPH was found in 2.9% of those with intermediate or high probability for PH (by echocardiography) (87). It is important to note that some patients with sarcoidosis may have normal pulmonary artery pressure at rest with evidence of exercise-induced PH (88).

Pathogenesis

There are several mechanisms of PH in sarcoidosis. Each one of these may have a different response and outcome to a specific therapy. Pre-capillary PH may result from hypoxic pulmonary vasoconstriction and destruction of the distal vascular bed with advanced interstitial lung disease. This is supported by the finding that the majority of SAPH patients have advanced fibrotic lung disease (89–91). This usually leads to mild to moderate PH (mPAP <35 mmHg). Nevertheless, at least one third of sarcoidosis patients have no significant fibrosis which indicates that hypoxemia and fibrosis are not the only explanation for SAPH.

Granulomatous vasculopathy and angiitis with obliteration of arterioles or venules has been reported in autopsies of patients with pulmonary sarcoidosis (92–94). Occlusion of pulmonary vessels by granulomas causes PH and sometimes can be mistaken with thromboembolic disease. This pathology is associated with more severe PH than that related to lung fibrosis. Granulomatous



vasculitis can also affect the post-capillary vessels causing focal stenosis and a clinical pattern consistent with pulmonary veno-occlusive disease (95, 96).

Moreover, sarcoidosis is associated with increased risk of venous thromboembolism, which carries associated risk of developing CTEPH in some patients (97, 98). Sarcoidosis can be associated with mediastinal and hilar lymphadenopathy that can cause extrinsic compression of pulmonary arteries (and venous stenosis) or fibrosing mediastinitis, resulting in SAPH.

Cardiac involvement occurs in 20–25% of patients with sarcoidosis (99). Thus, post-capillary PH can develop in sarcoidosis due systolic or diastolic dysfunction with left ventricular involvement. In a retrospective study of 130 sarcoidosis patients with persistent dyspnea, 29% of patients with SAPH had an elevated PAWP >15 mmHg (89). Because cardiac involvement can be subtle, screening for cardiac sarcoidosis using magnetic resonance imaging and 18 F-fluorodeoxyglucose positron emission tomography is recommended (100). Finally, post-capillary SAPH can be due to extrinsic compression of the pulmonary veins.

Treatment

As outlined earlier, SAPH can be related to multiple mechanisms (Figure 3). However, some patients may have a prominent mechanism. For example, in the presence of advanced parenchymal (fibrotic) lung disease, PH with mPAP of <35 mmHg is likely related to hypoxemia and destruction of the pulmonary vascular bed associated with

fibrotic changes. Treatment in these patients should focus on treatment of parenchymal lung disease and correction of hypoxemia. PAH-specific therapy carries the risk of worsening ventilation-perfusion mismatch and, therefore, should be avoided.

PAH-specific therapy can be considered in carefully selected patients with moderate to severe SAPH that is “out-of-proportion” to the degree of parenchymal lung disease and hypoxia (i.e., mPAP >35 mmHg). Such degree of PH suggests a component of vasculopathy that may respond to pulmonary vasodilators. Vasoreactivity has been demonstrated in a small prospective observational study of 8 patients with moderate-to-severe SAPH with improvement in mPAP and PVR to inhaled NO (102). This highlights the potential role of pulmonary vasodilators in these patients. Such therapy has been evaluated in multiple studies and shown to improve the hemodynamic parameters with no significant change in functional status or 6-min walk distance (Table 3). Similarly, in a double-blind placebo-controlled trial, bosentan improved mPAP and PVR, but failed to demonstrate benefit in 6-min walk distance (110).

Immunosuppressive therapy alone may result in hemodynamic improvement in patients with granulomatous vasculopathy or compressive lymphadenopathy (101). SAPH due to extrinsic compression can be treated with pulmonary artery or vein angioplasty and stenting (114). Overall, patients with SAPH have poor prognosis and should be evaluated early for lung transplantation.

TABLE 3 | Existing published studies evaluating treatment of sarcoid-associated pulmonary hypertension.

Publication	Study design (number of subjects)	Treatment (# of patients)	Outcome
Preston et al. (102)	Prospective observational (8)	Inh NO (5), inh NO with IV epo (1), CCB (2)	Short term 20% decreased PVR and mPAP; long term increased 6MWT
Culver et al. (103)	Retrospective chart review (7)	Bosentan (3), bosentan and IV epo (4)	Decreased mPAP at 6–18 mo in about 50% patients
Fisher et al. (104)	Retrospective case series (7)	IV epo (6), subcut trep (1)	Improved functional class
Milman et al. (105)	Retrospective chart review (12)	Sildenafil (12)	Decreased mPAP and PVR, increased CO, no change 6MWT
Barnett et al. (106)	Retrospective case series (22)	IV epo (1), bosentan (12), sildenafil (9)	Increased 6MWT and functional class, decreased mPAP and PVR
Baughman et al. (107)	Prospective open label 16 weeks (15)	Inh iloprost (15)	Decreased mPAP and PVR in 6 of 15 and increased 6MWT in 3 of 15 patients
Baughman et al. (89)	Retrospective chart review (5)	Bosentan (5)	Decreased mPAP in 3 of 5 patients at 4 mo
Judson et al. (108)	Prospective open label 12 weeks (25)	Ambrisentan (21)	No change 6MWT; 11 patients discontinued drug at 12 weeks
Dobarro et al. (109)	Retrospective chart review (8)	Sildenafil (9), bosentan (2); only 8 followed up with repeat RHC	Increased 6MWT and decreased NT-proBNP; non-statistically significant increase in CO/CI and decreased PVR
Baughman et al. (110)	Prospective placebo-controlled 16 weeks (35)	Bosentan (23), placebo (12)	Decreased mPAP and PVR; no change in 6MWT
Keir et al. (111)	Retrospective (33)	Sildenafil (29), bosentan (3)	Increased 6MWT, decreased NT-proBNP, improved TAPSE
Bonham et al. (112)	Retrospective case series (26)	Parenteral prostacyclin with epo (7) and trep (6), ERAs (12), PDE-5i (20), CCB (1)	Increased CI/CO, decreased PVR, improved functional class, decreased NT-proBNP
Ford et al. (113)	Prospective open label 24 weeks (12)	Tadalafil (12)	No change 6MWT at 24 weeks

Inh, inhaled; NO, nitric oxide; IV, intravenous; epo, epoprostenol; CCB, calcium channel blocker; mPAP, mean pulmonary arterial pressure; PVR, pulmonary vascular resistance; 6MWT, 6 min walk test; CO, cardiac output; CI, cardiac index; mo, months; NT-proBNP, N terminal pro brain natriuretic peptide; TAPSE, tricuspid annular plane systolic excursion.

PULMONARY HYPERTENSION IN CHRONIC KIDNEY DISEASE AND END STAGE RENAL DISEASE

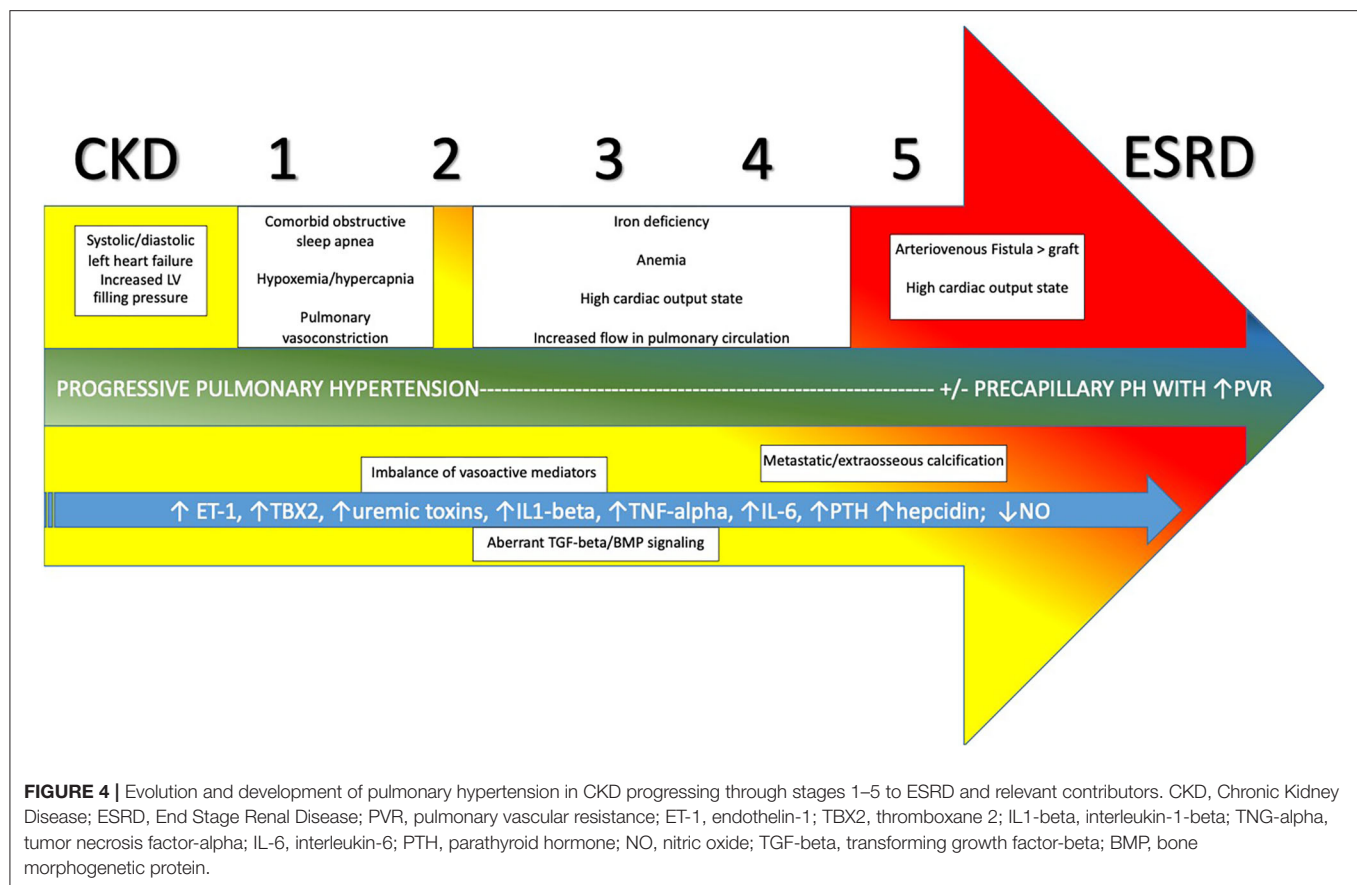
The evaluation and management of PH in the context of chronic kidney disease (CKD), particularly end stage renal disease (ESRD), presents unique challenges to the clinician in terms of elucidating the main drivers of PH in this patient population. The clinical presentation of such patients is one in which PH is often multifactorial, hence the placement of this clinical entity into Group 5 of the PH classification schemata. As such, management relies in large part on attempts to correct and optimize multiple physiologic derangements often seen in CKD and ESRD.

Definition, Epidemiology, and Scope of the Problem

CKD is defined as abnormal renal function present for at least 3 months and is staged in severity based on the degree of reduction of glomerular filtration rate and the degree of albuminuria. ESRD is defined as renal failure requiring regular interval, long-term dialysis, or renal transplant to maintain survival. The true prevalence of PH amongst patients with CKD and ESRD is difficult to discern, due to the lack of robust data. Most of what is known is drawn from relatively small, retrospective, and almost exclusively echocardiography based studies. Amongst patients

with CKD and not yet on dialysis, the prevalence of PH was observed to be upwards of 30% or more. Prevalence was highest among those with most advanced CKD (CKD 5) (115, 116). When considering those patients on hemodialysis (HD), a meta-analysis of known studies by Tang and colleagues (116) showed the prevalence of PH (based largely on echocardiography) between 19 and 56%. There is less data available with regard to the prevalence of PH in the setting of peritoneal dialysis (PD), but it appears to be lower than that of HD (117).

O'Leary et al. (118) recently published a retrospective analysis of a large cohort of patients with varying degrees of CKD that underwent RHC. PH was present in 1,267/1,873 (68%) of patients, using the accepted definition of PH at that time of mean PAP \geq 25 mmHg. Of these, 76% had post-capillary PH and 24% had pre-capillary disease, using elevated PAWP (\geq 15 mmHg) as the differentiating factor. The authors also noted that mortality was higher among patients with advanced stages of CKD, and further compounded by the presence of PH. The distribution of hemodynamic profiles seen underscores the significant role of Group 2 PH phenomenon related to elevated left ventricular filling pressures in driving PH in an overwhelming majority of patient with CKD. Comorbid left ventricular systolic and/or diastolic dysfunction, as well as a propensity to volume overload at times are the main drivers of this. However, many of those with CKD/ESRD with and without elevated PAWP have other factors driving PH as well. Among these are alterations in circulating



inflammatory and vasoactive mediators, endothelial dysfunction, alterations in cardiopulmonary flow related to anemia and arteriovenous fistula (AVF), factors related to dialysis itself, and comorbid Group 3 PH processes (see **Figure 4**).

Altered Inflammatory and Vasoactive Mediators and Related Endothelial Dysfunction

The presence of uremic toxins in patients with advanced CKD or ESRD can contribute to pulmonary vascular endothelial dysfunction. Specifically, mono-methyl L- Arginine, symmetric dimethyl arginine, and asymmetric dimethyl arginine are present in increased levels and inhibit NO synthase, impairing the intrinsic production and bioavailability of NO, the vasodilatory effects of which serve to maintain normal pulmonary vascular tone. There is also evidence that increased levels of uremic toxins are associated with increased inflammation, endothelial dysfunction, and oxidative stress (119). There is decreased availability of circulating NO in patients on HD with PH vs. those without PH. In addition, patients on HD with PH showed decreased ability to recover NO levels after HD, suggesting impairment in NO synthesis (120). As noted previously, ET-1 contributes to pulmonary vascular remodeling, is present in excess in patients with Group 1 PAH, and is also known to be elevated in patients with CKD and those on HD. Further, HD

does not facilitate clearance of ET-1 and levels remain elevated after HD is performed (120). Angiopoietin 1 and 2 are additional mediators that modulate vascular smooth muscle growth via action on endothelial cells. There is evidence that both are important in the pathogenesis of CKD and PAH (121).

Cellular inflammation is a key driver of disease in both PH and CKD. In addition to the noted effects of uremic toxins, levels of circulating pro-inflammatory cytokines are increased in patients on HD with PH relative to those without, particularly IL-1beta, TNF-alpha, and IL-6. High-sensitivity c-reactive protein was also noted to be higher in the group with PH (122). The transforming growth factor beta (TGF-beta) and bone morphogenetic protein (BMP) signaling pathways modulate downstream gene expression that controls properties of vascular structure and tone. It has been demonstrated that this pathway is aberrant in CKD and PAH, leaning in a pro-inflammatory direction that favors vascular remodeling. Thus, there is a strong likelihood that the TGF-beta/BMP axis is relevant in patients with both PH and CKD/ESRD (123, 124).

Alterations in Cardiopulmonary Flow

Intentional arteriovenous shunting caused by the surgical creation of AVF or placement of a synthetic arteriovenous graft (AVG) contributes to a high cardiac output state. This leads initially to increased loading on the left ventricle. When this phenomenon is coupled with the increased output state, a

syndrome of high-output heart failure may manifest. With time, persistence of this high flow is thought to lead to increased shear forces on the pulmonary vascular endothelium that can cause pulmonary vascular remodeling and an intrinsic arteriopathy in some patients (125). The propensity to develop PH is higher in patients with AVF relative to AVG, as AVF can enlarge with time and develop progressively higher flows, which in turn increases flow through the pulmonary circulation to increasing degrees. It has been noted when flow through the fistula exceeds 2 L/min or more than 20% of cardiac output, the risk of pulmonary hypertension is higher. Furthermore, presence of the AVF on the upper arm presents increased likelihood of PH as compared to AVF of the lower arm (126).

Practices regarding the placement and maintenance of AVF for HD have come under a bit more scrutiny in recent years as a result of noted effects on circulatory and pulmonary vascular physiology. There has been deliberate advocacy on the part of the Centers for Medicare and Medicaid Services to utilize AVF over catheter-based HD in most patients, driven largely by legitimate concern for the risk of catheter related bloodstream infection and thrombosis (127). Since the inception of this initiative, it has been increasingly recognized that some patients with CKD/ESRD may have subclinical or mild PH before arterio-venous access for dialysis is placed. Some advocate for more systematic echocardiographic screening for PH prior to placement of AVF. If evidence of PH is present, then PD and/or expedited/prioritized renal transplant evaluation may be more appropriate (128). Further study is needed in this regard.

Anemia due to reduced renal production of erythropoietin, as well as iron deficiency and anemia of chronic disease is very common in patients with CKD and ESRD. This contributes to compensatory increase in cardiac output (much like that seen in SCD) and thus exposes the pulmonary circulation to higher than usual flow and the risks of vascular remodeling noted above. Iron deficiency in both PAH and CKD appears to be modulated largely by increased levels of hepcidin, which inhibits enteric iron transport and uptake (129, 130). Iron deficiency thus may ultimately affect pulmonary vascular regulation through downstream effects on hypoxia inducible factors. Risk for development of PH in the context of CKD/ESRD and anemia is most significant when hemoglobin is <10 g/dl (131).

Factors Related to Dialysis

As noted, HD presents a higher likelihood of developing PH, due in part to the use of AVF or AVG in many patients. In addition, it appears that the type of dialysis membrane used may contribute to presence of PH as well. Thromboxane B₂, a pro-proliferative, pro-thrombotic, and pro-inflammatory molecule, is known to be elevated in patients with PAH compared to controls. In addition, it has been shown that thromboxane B₂ levels are higher in patients on HD and/or with PH as compared to those not on HD and/or with no evidence of PH (132). The reason for elevated thromboxane B₂ in some patients on HD appears to be related to increased release from circulating monocytes when cellulose dialysis filter membranes are utilized as compared to polysulfone membranes (133, 134).

The likelihood of PH is also directly proportional to older age and number of years on HD (131). The reason for this is likely simply related to the duration of exposure of the pulmonary circulation to the aforementioned pathophysiologic effects of HD. An additional phenomenon that may contribute to pulmonary vascular changes and PH is the showering of micro-bubbles from the dialysis machine with each HD session. These micro-bubbles have the potential to obstruct small pulmonary capillaries, leading to local inflammatory changes and microthrombosis, setting the stage for progressive pre-capillary arteriopathy to progress as HD continues over time (135, 136).

Comorbid Group 3 PH Processes

As noted previously, Group 3 PH is often driven by abnormal oxygenation and ventilation, commonly seen during sleep in patients with obstructive sleep apnea (OSA). There is a high prevalence of OSA amongst patients with CKD, independent of the need for HD (137, 138). The associated hypoxemia, and in some cases hypercapnia, contributes to pulmonary vasoconstriction and some degree of PH in many patients with CKD. Asymmetric dimethyl arginine is also noted to be elevated in patients with sleep disordered breathing, implicating impairment of NO production as at least one plausible molecular mechanism for the development of PH attributable to OSA (139). This pathway may be of particular relevance to patients with CKD. As such, screening for nocturnal hypoxemia and OSA with formal polysomnography is of paramount importance. Those found to have sleep disordered breathing should be titrated for and treated with nocturnal non-invasive positive airway pressure therapy.

In addition to OSA, it has been noted that secondary hyperparathyroidism seen in CKD/ESRD can lead to extraosseous calcium deposition in the interalveolar septae and pulmonary arterial system. Alveolar septal deposition can contribute to restrictive physiology on pulmonary function testing, as well as reduction in diffusing capacity (140). The overall clinical impact of these phenomenon with respect to the development of PH is not as clear compared to that of OSA. Additionally, it has been shown that parathyroid hormone levels do not differ between patients with and without PH in the context of CKD/ESRD (141).

PH Diagnosis and Invasive Hemodynamic Evaluation in Patients With CKD/ESRD

As with all forms of pulmonary hypertension, the initial preferred tool for screening is transthoracic echocardiography. This is especially important in CKD/ESRD patients with unexplained dyspnea. Timing of echocardiography is most optimal soon after the completion of a session of dialysis, with the patient at or very near their established dry weight. The presence of estimated PASP ≥ 50 mmHg and or significant right ventricular dilatation or dysfunction should prompt invasive hemodynamic measurement by RHC. As with echocardiography, RHC should be performed in the relative immediate time period after a session of dialysis, such that the left and right sided circulation are as volume and pressure off-loaded as possible, ensuring that hemodynamic

measurements are reflective largely of alterations in flow and/or PVR that may be present.

The PEPPER study (142) was an important endeavor that helped to understand the frequency of PH in patients with advanced CKD (stage 4 or 5) and unexplained dyspnea, as well as to understand the effect of HD on pulmonary hemodynamics. In the study, 31 CKD patients not on dialysis and 31 patients on HD with dyspnea underwent RHC. In the group on HD, 78% were found to have PH (13% pre-capillary PH, 65% post-capillary PH). Among those not on HD, 77% had PH (6% pre-capillary PH, 71% post-capillary PH). Further, in the group on HD, 25/31 underwent RHC pre and post dialysis. It was noted that mean PAP and PAWP were both significantly lower post-dialysis, underscoring the importance of timing of RHC relative to HD.

In patients that are on HD and have an AVF, it may be useful to study the effects of fistula occlusion during RHC, however there is scant literature to guide this practice. This must be done cautiously and for a relatively short time, as there is some risk of associated thrombosis of the AVF. Published studies looking at the hemodynamic effects of fistula compression are echocardiography based. Yigla et al. found that after compression of AVF for 1 min in four patients on HD with echocardiographic evidence of PH resulted in a 10% reduction in mean estimated PA systolic pressure and a 15% reduction in estimated cardiac output (reduction from supranormal values) (143).

Treatment of PH in the Context of CKD and ESRD

Given that the majority of PH in the context of ESRD/HD is pulmonary venous hypertension, the mainstay of treatment is optimizing volume status with diuretics or dialysis, controlling systemic blood pressure, and medical therapies that optimize systolic and diastolic function of the left ventricle. The importance of correcting hypoxemia and sleep disordered breathing when relevant was previously discussed. If clinical history or risk factors suggest that CTEPH may be a possibility, then consideration must be given to screening ventilation-perfusion lung scan. If there is evidence of CTEPH, then additional imaging and hemodynamic workup is indicated to determine if the patient is a candidate for PTE or BPA. For patients in whom there is exceedingly high AVF flow and elevation in cardiac output driving PH, consideration should be given to revision of the AVF via banding or ligation (144, 145).

In general, trials of off label use of PAH-specific therapies in group 5 PH related to CKD/ESRD should be reserved for a very select few patients for whom all other contributing factors to PH have been optimized but have persistent and significant RHC-proven PVR elevation, particularly in the context of reduced cardiac output reflective of RV failure. The presence of clear Group 1 PAH comorbidities such as connective tissue disease, HIV, or portal hypertension, may also make a case for treatment with PAH therapies more compelling in the setting of a hemodynamic profile consistent with clear pre-capillary disease. As with all group 5 PH, there is no randomized, controlled data regarding the use of PAH-specific therapy in CKD/ESRD

associated PH. Though not absolutely contraindicated in renal failure, PDE-5 inhibitors and riociguat should be used with caution in patients with advanced CKD/ESRD due to potential alterations in pharmacokinetics. As such, dose adjustments may be needed in this context. (146, 147).

Renal transplant surgery is generally regarded as not presenting prohibitive levels of operative risk to patients with mild to moderate degrees of PH and RV dysfunction, particularly considering the procedure is not usually associated with large intravascular volume and hemodynamic shifts as seen for example in liver transplantation. Patients with significant PH, and particularly severe right ventricular dysfunction, are considered with an abundance of caution for renal transplantation, especially if interventions to attempt to improve pulmonary hemodynamics have failed. There is no clear evidence based data available to systematically risk stratify patients with ESRD and PH for renal transplant surgery (148). Independent of immediate perioperative surgical risk however, it is noted that pre-transplant PH (based on echocardiographic studies) is associated with a significant increased risk of graft failure (149). Furthermore, post-transplant survival is decreased in patients with elevated estimated PASP on echocardiogram > 50 mmHg and in those with elevated PVR > 3 Wood units (150, 151). In general, approach to renal transplantation candidacy in the setting of PH should be done in an individualized, multidisciplinary fashion, especially given the lack of robust studies to reliably predict outcomes.

PULMONARY HYPERTENSION ASSOCIATED WITH LESS COMMON COMORBIDITIES

Pulmonary Langerhans Cell Histiocytosis

Pulmonary Langerhans cell histiocytosis (PLCH) is an uncommon cystic lung disease characterized by pathologic dendritic cell infiltration of the walls of the distal bronchiole. The involved dendritic cell subset, the Langerhans cell, localizes within the airway epithelium and processes encountered inhaled antigens, becoming activated when additional molecular signals indicate a state of danger (152). Activated dendritic cells migrate to lymphoid tissue to induce the adaptive immune response. PLCH occurs almost exclusively in smokers (153), and smokers with or without pulmonary diseases have significantly greater presence of Langerhans cells in the lung (154, 155). Cigarette smoking further induces production of molecules such as TNF-alpha, TGF-beta, GM-CSF, and osteopontin; these factors promote development, activation, and survival of Langerhans cells, though migration out of the airway tissue may be impaired (156–158). The bronchiolocentric nodules characteristic of PLCH contain a mixed population of Langerhans and other inflammatory cell types (159). Factors predisposing the small population of smokers developing PLCH remain incompletely understood.

Relative to other forms of chronic lung disease, PLCH is associated with higher prevalence of PH, and the vascular component of disease can be great (160, 161). PH has been

found in 41–100% of PLCH patients, with prevalence particularly high (>90%) in those undergoing lung transplant evaluation (160, 162–164). Presence of PH has been consistently associated with increased mortality in PLCH (163, 165). While inverse correlation between pulmonary artery pressure elevation and spirometry has been noted in some studies (163, 166), others have not demonstrated this finding (160, 161). Examinations of tissue specimens have demonstrated typical vascular changes of PAH, including intimal fibrosis and medial hypertrophy of small and medium-sized pulmonary arteries (160). Observation of Langerhans cells with vessels is rare, and vascular involvement can also be noted in regions without parenchymal infiltration of Langerhans cells (160, 161). In contrast to most forms of PAH, capillary and venular pathology appears to be common in PLCH-PH (160, 165–168).

Management of PLCH, for those with or without associated PH, should most importantly center around smoking cessation. Stabilization and even substantial disease regression has been observed in those successful at quitting (169–172). While most case reports and series have focused on improvements in radiology or pulmonary function following smoking cessation, benefits on the pulmonary vascular component of disease should also be presumed. One case report of a patient with PLCH and associated PH has described PH resolution following smoking cessation (173). For patients with symptomatic PLCH refractory to smoking cessation, steroids, and chemotherapeutic agents have been used with mixed results (174–178). Use of PAH-specific therapy in those with PLCH-PH has been best described in a series by Le Pavec et al., where 14 patients were treated with an ERA, PDE-5I, and/or inhaled iloprost. Use of PAH-specific therapy was associated with improvements in mPAP and PVR on follow-up hemodynamic reassessments; a trend toward improved survival was also observed (167). With histologic observation of pulmonary venous involvement in PLCH-PH, the potential for development of pulmonary edema in response to PAH-specific therapy must be acknowledged, and cases consistent with this outcome have been documented after initiation of intravenous epoprostenol (160). Encouragingly, oxygenation of treated patients did not worsen in the series by Le Pavec (167). Given vascular complexity, lung transplant should be considered in those with associated PH and in others with severe PLCH. PLCH patients make up <1 percent of lung transplant recipients in the United States, with overall similar post-transplant outcomes to other patient groups (162, 164).

Neurofibromatosis Type 1

Neurofibromatosis type 1 (NF1) is an autosomal dominant genetic disorder caused by mutation in NF1, the tumor suppressor gene encoding neurofibromin. Incidence is 1 in 3,000 births, with near complete penetrance; skin manifestations are usually prominent and early in onset. Major manifestations leading to diagnosis include café au lait macules, cutaneous and subcutaneous neurofibromas, axillary and inguinal freckling, and Lisch nodules of the iris (179). NF1-related parenchymal lung disease affects 10–20% of adults and is characterized by cysts, micronodules, and interstitial abnormalities (179–181). Arterial and venous vascular lesions, including arterio-venous malformations, arterial aneurysms,

coarctation of the abdominal aorta, and renal artery stenosis, are important but under recognized sources of morbidity in NF1 (182, 183). Neurofibromin is expressed in endothelial and vascular smooth muscle cells, and it is thus suspected that absence of functional protein leads to dysregulation of growth and proliferation of intimal and medial layer cells, in systemic and pulmonary circulations (182, 184). Patients with NF1 also have increased malignancy risk, and cancer is the leading cause of death with life expectancy reduced by 10–15 years (185, 186).

Pulmonary hypertension in NF1 patients has been described only in case reports and series, thus prevalence of NF1-PH is difficult to estimate but thought to be rare. The largest series of PH in NF1 recently characterized 49 patients from the French PH National Referral Center. Despite NF1 affecting males and female equally, 80% of NF1-PH patients were female, and median age at diagnosis of PH was 62. No specific NF1 mutations were associated with PH, and others with NF1-PH were not observed in the families of most patients. Hemodynamics at diagnosis were consistent with severe pre-capillary PH (mean mPAP 45 with PVR 10.7 WU). The majority of patients had normal or mildly abnormal spirometry, with decreased diffusion capacity for carbon monoxide (median 30% predicted) and resting hypoxemia. Imaging was most frequently notable for diffuse lung cysts, ground glass opacities, and emphysema (187). Importantly, this series and another literature review have identified normal imaging or single radiologic abnormalities in nearly 30% of NF1-PH patients (184, 187). Lung tissue has been infrequently available; the few histologic studies have demonstrated arterial remodeling with muscularization of small arterioles as well as particular increase in intimal layer thickness (187, 188). Findings of thrombosis have not been noted, and presence of plexiform lesions has been variable (187, 189). Three patients undergoing transplant all had pulmonary venous wall thickening (187), and a patient with pulmonary capillary hemangiomatosis associated with NF1 has also been reported (190).

Prognosis of NF1-PH is poor, with transplant free survival of 87, 54, and 42% at 1, 3, and 5 years, respectively (187). Of patients in the large series by Jutant et al., 45/49 were started on PAH-specific therapy; ERAs, PDE-5Is, and prostacyclin derivatives were all used alone or in combination. Treated patients demonstrated improvement in cardiac index and PVR; however, oxygenation worsened slightly and clinical improvements were modest and not sustained (187). Lung transplantation has been successfully performed for NF1 patients (187, 188); however, concern for malignancy risk following transplant and immunosuppression can be a barrier (191). The tyrosine kinase inhibitor sorafenib, which is capable of suppressing kinases operating in pathways also shared by targets of neurofibromin, was used in a single patient with refractory NF1-PH. Hemodynamic and clinical improvements resulted after 3 months of use (192). Given neurofibromin's role in containment of cell proliferation, investigation of drugs impacting cell cycle regulation are particularly of interest in NF1-PH.

Gaucher Disease

Gaucher disease (GD) is an autosomal recessive lysosomal storage disease with an incidence of 1 in 50,000 births. GD

results from inadequate function of the lysosomal enzyme glucocerebrosidase, leading to accumulation of its substrate in macrophages, described as Gaucher cells. Accumulation of macrophages is particularly prominent in the bone marrow, liver, and spleen, with resulting hepatosplenomegaly, thrombocytopenia, anemia, osteopenia, and painful bone crises. GD is classified into three subtypes (GD 1, 2, and 3), with GD type 1, the non-neuronopathic form of GD, being the most common. Pulmonary involvement can be seen in all GD types, though those with GD1 are at most risk for developing pulmonary vascular complications of the disease (193). Enzyme replacement therapy for GD was first approved by the United States Food and Drug Administration in 1991 and has provided significant improvements in hematologic counts, spleen and liver volumes, and other clinical parameters for GD type 1 (194, 195). Importantly, use of enzyme replacement therapy has dramatically decreased need for splenectomy, previously performed for GD due to lack of other management options (195–197).

The prevalence of PH in adult GD type 1 patients, as estimated by echocardiography, ranges from 7 to 30% (198, 199). Mild PASP elevation (35–50 mmHg) is noted in the majority, with 0.76% documented to have PASP >50 (198). Treatment with enzyme replacement therapy is associated with lower prevalence of PH (7.4% in treated patients vs. 30% in untreated patients) (198). Though enzyme replacement therapy has been questioned to be causative of PH (200, 201), most current evidence suggests protection, and multiple examples of PH-related clinical improvements after initiation exist (198). In contrast, splenectomy has strong association with development of GD-PH. Severe PH complicating GD occurs predominantly in females who have undergone splenectomy as part of GD management (198, 199). The largest published series of GD-PH patients included 14 adult patients from a single referral center. Females made up 71% of the population, with a mean age of 36 at PH diagnosis, and all patients had previously undergone splenectomy at mean age 12. Other pertinent findings from this series includes lack of correlation between development of PH and GD severity by other metrics. Heteroallelic mutations (with one copy of the most common mutation N370S) were more prevalent than N370S homozygotes (199). Of interest, this report and others provide examples of patients also meeting criteria for hepatopulmonary syndrome; in these cases PH was either coexisting with hepatopulmonary syndrome or was unmasked after initiation of enzyme replacement therapy and resulting resolution of hepatopulmonary syndrome (199, 202). When tissue has been available, observed arterial changes have resembled those of PAH, including medial hypertrophy and intimal fibrosis, as well as plexiform lesions (203–205). In

addition, intravascular Gaucher cells have been observed, though are not uniformly present (205–207).

If not previously initiated, most GD-PH patients will start on enzyme replacement therapy, with case reports of this alone leading to significant improvement (198, 199). ERAs, PDE-5Is, and prostanoids have all been used in individual patients with GD-PH, generally with positive outcomes (198, 199, 208). Given shared risk factor (splenectomy) with CTEPH, use of anticoagulation should be discussed particularly in splenectomized patients, though data about its role is lacking. Coumadin has been used in several patients in combination with enzyme replacement therapy and PAH-specific therapy (198). In cases of refractory GD-PH, imatinib has been used successfully in a single patient (209). Finally, though other comorbidities can pose a challenge to candidacy, lung transplant has been used as a treatment modality for a few cases of severe GD-PH (203, 205).

DISCUSSION

Group 5 pulmonary hypertension remains a varied and challenging group of clinical entities. Understanding of the pathogenetic factors underlying many of the clinical conditions in the group has advanced considerably, however the relative rarity and/or variable clinical presentation of many of these disease processes has made it challenging to identify unifying treatment principles that can be applied broadly. Greater understanding may be gained by enhanced prospective registries of the more common conditions associated with PH, particularly sarcoidosis, sickle cell disease, myeloproliferative disorders, and chronic kidney disease/ESRD. More robust randomized trials of PAH therapies will be needed to understand if any of the ever-increasing number of PAH treatments may be of benefit in any of these populations.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Novel *TNIP2* and *TRAF2* Variants Are Implicated in the Pathogenesis of Pulmonary Arterial Hypertension

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Background: Pulmonary arterial hypertension (PAH) is a rare disease characterized by pulmonary vascular remodeling and right heart failure. Specific genetic variants increase the incidence of PAH in carriers with a family history of PAH, those who suffer from certain medical conditions, and even those with no apparent risk factors. Inflammation and immune dysregulation are related to vascular remodeling in PAH, but whether genetic susceptibility modifies the PAH immune response is unclear. *TNIP2* and *TRAF2* encode for immunomodulatory proteins that regulate NF- κ B activation, a transcription factor complex associated with inflammation and vascular remodeling in PAH.

Methods: Two unrelated families with PAH cases underwent whole-exome sequencing (WES). A custom pipeline for variant prioritization was carried out to obtain candidate variants. To determine the impact of *TNIP2* and *TRAF2* in cell proliferation, we performed an MTS [3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium] assay on healthy lung pericytes transfected with siRNA specific for each gene. To measure the effect of loss of *TNIP2* and *TRAF2* on NF-kappa-beta (NF- κ B) activity, we measured levels of Phospho-p65-NF- κ B in siRNA-transfected pericytes using western immunoblotting.

Results: We discovered a novel missense variant in the *TNIP2* gene in two affected individuals from the same family. The two patients had a complex form of PAH with interatrial communication and scleroderma. In the second family, WES of the proband with PAH and primary biliary cirrhosis revealed a *de novo* protein-truncating variant in the *TRAF2*. The knockdown of *TNIP2* and *TRAF2* increased NF- κ B activity in healthy lung pericytes, which correlated with a significant increase in proliferation over 24 h.

Conclusions: We have identified two rare novel variants in *TNIP2* and *TRAF2* using WES. We speculate that loss of function in these genes promotes pulmonary vascular

remodeling by allowing overactivation of the NF- κ B signaling activity. Our findings support a role for WES in helping identify novel genetic variants associated with dysfunctional immune response in PAH.

Keywords: pulmonary arterial hypertension, NF- κ B, inflammation, massive paralleled sequencing, *TNIP2*, *TRAF2*

INTRODUCTION

Pulmonary arterial hypertension (PAH) is a rare disorder associated with progressive elevation of pulmonary pressures that, if untreated, leads to right heart failure and death (1, 2). The major pathological features of PAH are obliterative vasculopathy and progressive loss of distal pulmonary microvessels that are unresponsive to available therapies (3, 4). While the cause of PAH remains incompletely understood, pathogenic variants in the bone morphogenetic protein receptor 2 (*BMPR2*) have been associated with both familial (60%) and sporadic PAH (20–25%) (5, 6). Studies have now shown that *BMPR2* is responsible for ensuring appropriate vascular repair in response to injury and loss of function variants are associated with impaired angiogenesis, as well as endothelial and smooth muscle cell proliferation within the vessel wall (7–9). Given the low penetrance of *BMPR2* variants, it is likely that additional genetic modifiers are required for the development of PAH. Besides genetic predisposition, immune dysregulation and inflammation have been established as risk factors for PAH development (10, 11). Recent work in this area has led to identifying immune subtypes within PAH, which may explain phenotypic differences among patients (12). Understanding the interaction between genetics and immunity could provide an avenue for revealing the basis of PAH susceptibility and might offer opportunities for novel therapeutics that could change the natural history of this disease.

Whole-exome sequencing (WES) is a valuable tool for discovering novel pathogenic variants in the protein-coding regions and the genome's intron-exon boundaries. The use of WES to analyze probands and family members (i.e., trios and quartets) has led to the identification of novel disease genes associated with multiple genetic disorders, including PAH (13–16). In this study, we employed WES to detect novel pathogenic variants in a patient trio and a quartet that had previously tested negative for known PAH pathogenic variants using our customized parallel sequencing PAH gene platform (17). We report the discovery of potentially pathogenic variants in *TNIP2* and *TRAF2*, two genes associated with inflammation and immunity whose role in PAH has not been previously documented. Through the use of patient-derived cells and *in vitro* gene knockdown studies, we show that *TNIP2* and *TRAF2* act by suppressing the activation of NF- κ B, a transcription complex that controls the expression of inflammatory cytokines and genes associated with cell proliferation in PAH (18–21). We conclude that *TNIP2* and *TRAF2* pathogenic variants could increase PAH susceptibility through their capacity to alter cellular immune responses and drive abnormal cellular proliferation in the pulmonary vasculature.

MATERIALS AND METHODS

Patient Studies

All patients involved in this study gave their informed consent to participate. Samples were obtained from the Spanish PAH registries (REHAP and REHIPED), and the ethical committee approved the project of scientific research of the Hospital Universitario La Paz, Madrid (PI-1210, PI-3996). PAH was confirmed in all patients by right heart catheterization (RHC) following the recommendations of the 6th World Symposium and the European Society of Cardiology (1, 2, 22). Screening of family members was carried out by performing a family pedigree of all individual index cases of PAH. In those with a clinical suspicion of heritable PAH, evaluation of medical history, physical examination, laboratory tests, 12-lead electrocardiogram, and transthoracic echocardiography of all family members was done to select possible affected members and RHC candidates.

Whole Exome Sequencing

The workflow used for WES analysis is summarized in **Supplementary Table 1**. Library preparation was carried by Agilent SureSelect™ (v. 6.0), all exons kit followed by sequencing in a NovaSeq Sequencer (Illumina, USA). The exomes were analyzed by Saphetor Varsome software (23) and VarSeq (Golden Helix, USA) to prioritize variants in the probands with segregation patterns consistent with potential disease pathogenesis. Copy number variant (CNV) analysis was carried out through VarSeq to detect large rearrangements (deletions, insertions, and Indels of more than 1 Kb). Segregation analysis and validation of the variants detected in the WES was performed by classical Sanger sequencing. Screening for pathogenic variants in well-known PAH genes was carried out through a custom gene panel described previously (17).

Variant prioritization was carried out by a custom algorithm (see **Supplementary Table 1**). The first step was to filter out variants not meeting the standard quality requirements for depth, genotype quality, and variant allele frequency, followed by segregation analysis based on the variant's suggested pattern of inheritance. All variants with a population frequency above 1% were filtered out, and the pathogenicity of the remaining variants was assessed using several bioinformatic tools included in the dbNSFP database (24) plus the computation of the CADD score, which predict the deleteriousness of variants throughout the human genome (25). The variant analysis was prioritized according to predicted pathogenicity. Predicted impact of variants on candidate gene function and role in PAH pathogenesis was carried out using *in silico* analysis via STRING and literature review, respectively.

Pericyte Isolation and Culture

Pericytes were obtained from human donor lung tissue samples and lineage was confirmed by FACS, and western immunoblotting as previously described (26–28). 3G5 (ATCC CRL-1814) hybridoma cells were cultured in 1X DMEM with 10% FBS. Culture medium was collected and concentrated by centrifugal filter Amicon Ultra-15 (Millipore). 3G5 IgM was isolated using the IgM Purification Kit (Pierce), and concentration was determined using the Lowry Method. The day before the extraction, 40 μ l of rat anti-mouse IgM magnetic beads (Invitrogen 11039D) were incubated with 10 μ g of 3G5 IgM overnight. The next day, fresh lung tissue was washed with 1X HBSS twice and digested in a 10 ml 1X HBSS solution containing 10 mg Collagenase (Sigma-Aldrich, USA), 1 mg dispase, and 75 μ g DNase for 15–30 min at 37 degrees on an orbital shaker at vigorous speed (150 rpm). An equivalent amount of ice-cold pericyte medium (ScienCell, USA) containing 2% FBS was added immediately after digestion. The suspension was passed through a 100 μ m mesh filter (BD Falcon, USA) to remove debris and undigested fibrous tissue and washed once with 1X PBS. The cell pellet was resuspended in 1 ml 1X PBS containing 0.2% FBS and 40 μ l of antibody-coated magnetic beads and gently rotated in 4 degrees for 45 min. Cells were used between Passage 8–10.

Western Immunoblotting

Cells were washed three times with ice-cold 1x PBS, and lysates were prepared by adding lysis buffer (1X radio-immunoprecipitation assay buffer, 1 mMol/L phenyl-methylsulfonyl fluoride, and 1x phosphatase inhibitor) and vortexed for 10s before centrifugation. Supernatants were transferred to fresh microcentrifuge tubes and stored at -20°C . The protein concentration was determined by Pierce BCA assay (Thermo Scientific Product #23227, Rockford, Illinois). Equal amounts of protein were loaded onto each lane of a 4–12% Bis-Tris gel (Life Technologies) and subjected to SDS-PAGE electrophoresis under reducing conditions. After blotting, polyvinylidene difluoride (PVDF) membranes were blocked for 1 h in blocking buffer (5% milk powder in 0.1% PBS/Tween 20) and incubated with primary antibodies overnight at 4°C , followed the next day by incubation with horseradish peroxidase-conjugated secondary antibodies. Bands were visualized using the ECL kit (Thermo Fisher).

RNA Interference

To achieve gene knockdown, 2 μ mol/L silencing RNA (siRNA) of ABIN2 (encoded by TNIP2) (L-014328-01, Dharmacon, Lafayette, CO), TRAF2 (L-005198-00, Dharmacon, Lafayette, CO), or non-targeting siRNA control transfected into healthy pulmonary pericytes. Transfection was performed using a Nucleofector II (Program U-025; Lonza) with the Basic Smooth Muscle Cell Nucleofection kit (Lonza). All experiments were performed 48 h after electroporation.

MTS Proliferation Assay

Cells were seeded in eight replicates on collagen-coated 96-well plates (5.0×10^3 cells per well) (Nunc, Rochester, NY) and cultured in complete medium overnight at 37°C and 5% CO_2 . The next day, reduced serum media and MTS reagent were

added to each well for 1 h in accordance with the manufacturer's instructions. MTS absorbance was measured at 490 nm.

Statistical Analysis

The number of samples studied per experiment is indicated in the figure legends. Values from multiple experiments are expressed as mean \pm SEM. Statistical significance was determined using the Mann-Whitney test. A value of $P < 0.05$ was considered significant.

RESULTS

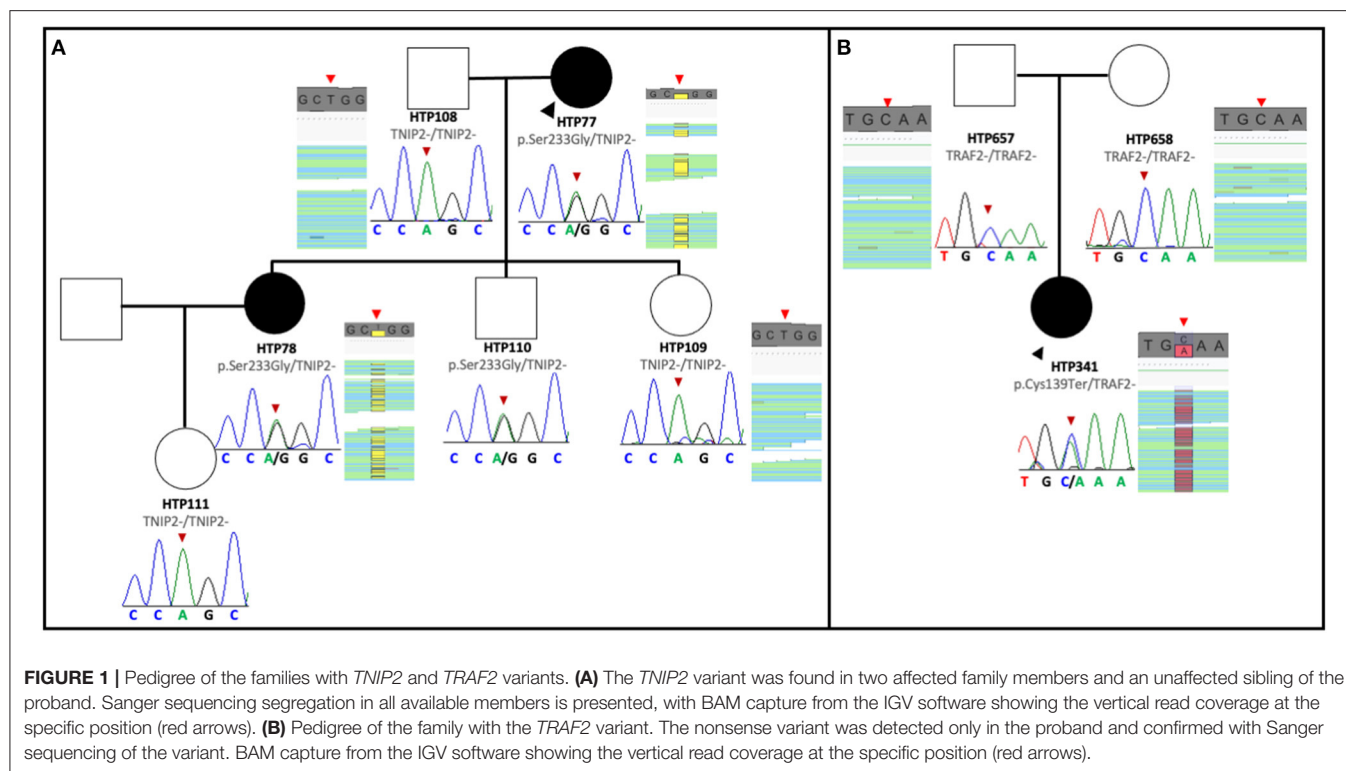
Discovery of the TNIP2 Variant

The pedigree of the first family is shown in **Figure 1A**. The index case was a 63-year-old female with a history of Ostium Secundum atrial septal defect (ASD), which was surgically corrected in 1994, and severe restrictive lung disease caused by thoracic deformities and diaphragmatic paralysis. The patient was diagnosed with PAH when she presented with exertional dyspnea and chest pain. Despite treatment with sildenafil and ambrisentan, the patient died 4 years after diagnosis from progressive right heart failure. Her 45-year-old daughter was diagnosed in 2015 with PAH associated with limited scleroderma and remains clinically stable with triple therapy of tadalafil, bosentan, and intravenous epoprostenol. The clinical characteristics of the two affected patients are summarized in **Table 1**. The remaining family members were screened and demonstrated no signs and symptoms of PAH to date. Given the familial predisposition, we conducted a genetic screen of the patients and family members using an in-house custom panel (HAPv2.1) (17), which was negative for mutations in 21 genes associated with PAH. As a next step, we conducted WES on the two patients and four unaffected family members, followed by segregation analysis (**Figure 1A**).

Analysis of WES data confirmed the absence of mutations in PAH associated genes. However, we discovered that both PAH patients were carriers of a novel missense variant in TNFAIP3 Interacting Protein 2 (*TNIP2*). This gene encodes for the protein ABIN2, which acts as an inhibitor of the NF- κ B signaling pathway (29). The variant is predicted to change an adenine to a cytosine base at position 697 (NM_024309.4:c.697A>C), which results in a switch from serine to glycine at position 233 [p.(Ser233Gly)] of the peptide chain, located between the two ubiquitin-binding domains of the protein. This novel *TNIP2* variant was predicted to be damaging based on the estimates obtained via dbSNFP and would likely result in loss of protein function (**Table 2**). Additionally, this variant is absent from several control population databases (gnomAD exomes, gnomAD genomes, Kaviar, Beacon, Bravo, ESP, 1000G phase III). Interestingly, one of the unaffected family members was also found to be a carrier of the same *TNIP2* variant and is currently undergoing clinical surveillance in our clinic.

Discovery of the TRAF2 Variant

The pedigree of the second family is shown in **Figure 1B**. The index case was a 27-year-old female that presented to the clinic in 2015 complaining of dizziness and was subsequently diagnosed with idiopathic PAH. Following initiation of tadalafil and ambrisentan, the patient developed elevated liver function



enzymes, which prompted ambrisentan cessation. During the workup for elevation in liver enzymes, the patient was diagnosed with primary biliary cirrhosis, and the PAH subtype was revised to portopulmonary hypertension. Intravenous epoprostenol was temporarily initiated to stabilize the patient, and treatment was later switched to triple oral therapy with tadalafil, macitentan, and selexipag (Table 1). At the time of writing, the patient remains clinically stable.

As part of the clinical evaluation, we applied the custom gene panel HAPv.2.1 to screen the patient and her unaffected parents but found no evidence of pathogenic variants. However, WES analysis detected a novel pathogenic variant in the Tumor Necrosis Factor Receptor-Associated Factor (*TRAF2*). This gene encodes for a protein involved in the TNF-dependent activation of NF- κ B (30). The variant is predicted to change cysteine to adenosine in position 417 (NM_021138.4:c.417C>A), introducing a premature stop codon and resulting in a truncated *TRAF2* transcript. Analysis of the parents' exomes showed that the variant is *de novo* in the proband. This variant is also absent from the control population databases (gnomAD exomes, gnomAD genomes, Kaviar, Beacon, Bravo, ESP, 1000G phase III), and the majority of the pathogenic *in silico* tools from the dbSNP suggested a damaging effect for this variant (Table 2).

TNIP2 and TRAF2 Interact With Pathways Associated With PAH

To date, no studies have shown evidence of *TNIP2* and *TRAF2* involvement in the pathogenesis of PAH. To determine whether these genes could potentially be interacting with components

of known PAH-associated pathways, we carried out an *in silico* protein-protein screen using STRING (Figure 2A). This tool illustrates protein-protein interactions based on function and previously described associations (31). Our STRING analysis showed that *TNIP2* and *TRAF2* indirectly interacted with the transforming growth factor- β (TGF- β) family genes via *CAV1*, a known risk gene for hereditary PAH (13). Notably, *BMPR2*, the gene most commonly implicated in hereditary and idiopathic PAH, is part of the TGF- β family identified within the STRING cluster. As expected, we found strong associations between *TNIP2* and *TRAF2* with proteins involved in the NF- κ B pathway (Figures 2B,C). Of the proteins associated with familial PAH, only *CAV1* directly interacts with *TRAF2* based on *in vitro* studies demonstrating that these proteins form a cytoplasmic protein-protein complex that modulates NF- κ B activation in response to TNF (32, 33).

Loss of TNIP2 and TRAF2 Result in Increased Pericyte Proliferation and Activation of NF- κ B

TNIP2 and *TRAF2* are part of a gene network responsible for regulating the NF- κ B pathway's signaling activity. The NF- κ B proteins exist in the cytoplasm in an inactive state and can be rapidly activated in response to harmful stimuli such as oxidative stress, bacterial lipopolysaccharides, environmental toxins, and inflammation. Once activated, the NF- κ B protein complex translocates to the nucleus and binds to target genes involved in cell proliferation, survival, and the production of cytokines that regulate innate and adaptive immune

TABLE 1 | Clinical features of patients.

Patient ID	HTP 77	HTP 78	HTP 341
Etiology	Group 1: PAH + ASD	Group 1: PAH + scleroderma	Group 1: PAH + primary biliary cirrhosis
Date of birth	22/09/1946	15/10/1975	24/08/1988
Age at diagnosis (years)	63	27	27
Current age (years)	Dead with 67 years	45	32
WHO functional class	III	I	I
Last known PAH therapies	Sildenafil, ambrisentan	Bosentan, tadalafil, epoprostenol	Selexipag, macitentan, tadalafil
6-min walk test (meters)	Not performed	595	454
CPET (date)	Not performed	May of 2018	March of 2018
RER		1.41	1.15
Maximum V _{O2} —mL/Kg/min (% of predicted value)		11.0 (33)	17.0 (53)
VE/VCO ₂ slope		33	38
Mean pulmonary artery pressure (mmHg)	42	51	55
Pulmonary capillary wedge pressure (mmHg)	14	6	8
Pulmonary vascular resistance (Wood units)	6.0	9.1	17.0
Cardiac output (lpm)	4.6 lpm	4.6 lpm	2.7 lpm
Cardiac index (lpm/m ²)	3.0	3.1	1.8
Mean arterial pressure or blood pressure (mmHg)	96	83	74
Echocardiographic abnormalities	Repaired Ostium Secundum ASD. Mild dilatation and hypertrophy of the RV and mild RV systolic dysfunction. Mild RA enlargement.	Eccentricity index of 1.7. RV hypertrophy and dilatation. RV systolic dysfunction. Moderate RA enlargement.	Eccentricity index of 1.1. RV hypertrophy and dilatation. Subtle RV systolic dysfunction. Mild RA enlargement.
Chest computed tomography abnormalities	Thoracic deformities and diaphragmatic paralysis	Bilateral ground glass opacities related with PH.	No abnormalities noted in chest CT
Forced expiratory volume in 1 s (mL) and predicted value by age and sex (%)	660 (43%)	2,790 (86%)	2,00 (89%)
Forced vital capacity (mL) and predicted value by age and sex (%)	1,060 (56%)	3,240 (61%)	3,185 (91%)
Diffusing capacity of the lungs for carbon monoxide (DLCO)	Not available	45%	56%
Partial pressure of oxygen (PaO ₂)	75 mmHg	57 mmHg	55 mmHg
V/Q scintigraphy and/or CT pulmonary angiography	Global matched perfusion and ventilation defects	Normal V/Q scan. Main pulmonary artery enlargement. RA and RV enlargement	Matched perfusion and ventilation defects in the upper right lung segment in V/Q scan.

Summary of the most relevant clinical characteristics of the patients in which a genetic variant was detected. ASD, atrial septal defect; CPET, cardiopulmonary exercise test; CT, Computed tomography; PH, pulmonary hypertension; RA, right atrium; RER, respiratory exchange ratio; RV, right ventricle; V/Q, Ventilation-Perfusion scan; WHO, World Health Organization.

TABLE 2 | Candidate genes and variant *in silico* analysis.

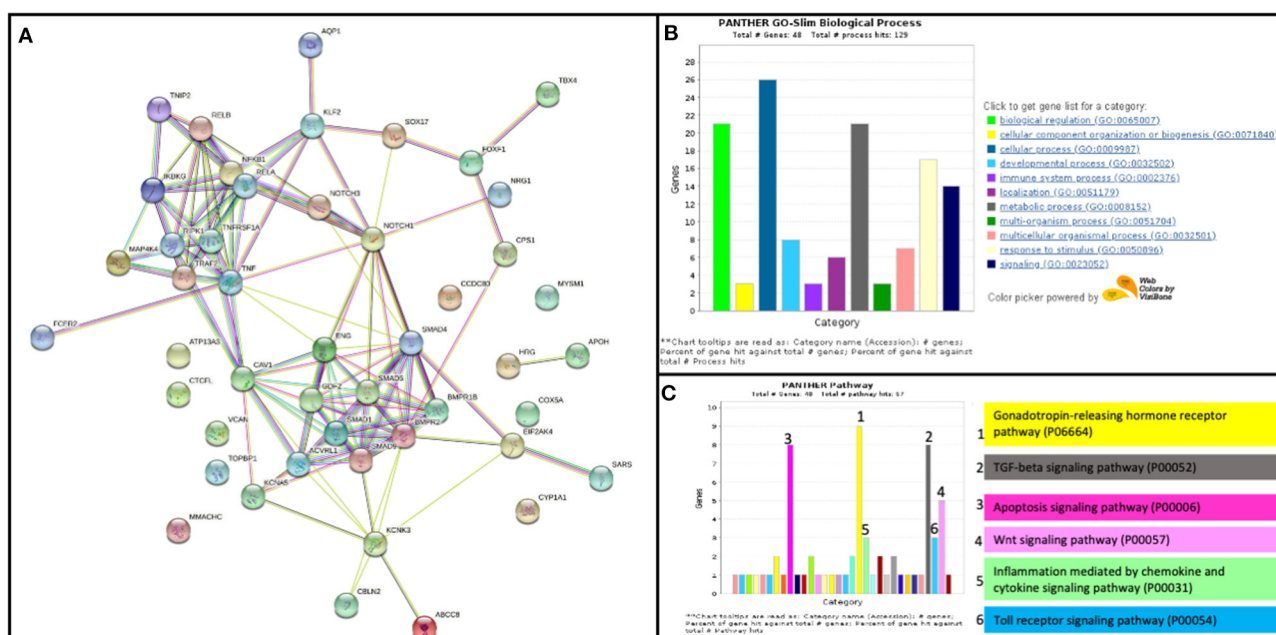
Gene	Variant information (hg19)				Pathogenicity		Population frequencies
	cDNA position	Protein position	Effect	Zygosity	dbSNFP	CADD	AF
TNIP2	NM_024309.3:c.697A>G	NP_077285.3:p.Ser233Gly	Missense	Heterozygous	12/19	25.8	0
TRAF2	NM_021138.3:c.417C>A	NP_066961.2:p.Cys139Ter	Non-sense	Heterozygous	5/9	35	0

Variants were annotated with the human reference genome hg19. Pathogenicity *in silico* prediction was obtained from the aggregation database dbSNFP and CADD (Table 1 for more info). AF, Allele frequency was estimated from several population pseudo-control databases: gnomAD genomes (v2.1.1), gnomAD exomes (v2.1.1), Kaviar (version 160204-Public), Beacon (v2.0), 1000G, Phase III and Bravo.

responses (34). Inappropriate NF- κ B activation is associated with a wide range of diseases, including cancer, autoimmune conditions such as scleroderma, and PAH (35). Given that

TNIP2 and TRAF2 serve to regulate NF- κ B activity, we hypothesized that the loss of these proteins would lead to NF- κ B overactivation.

STRING



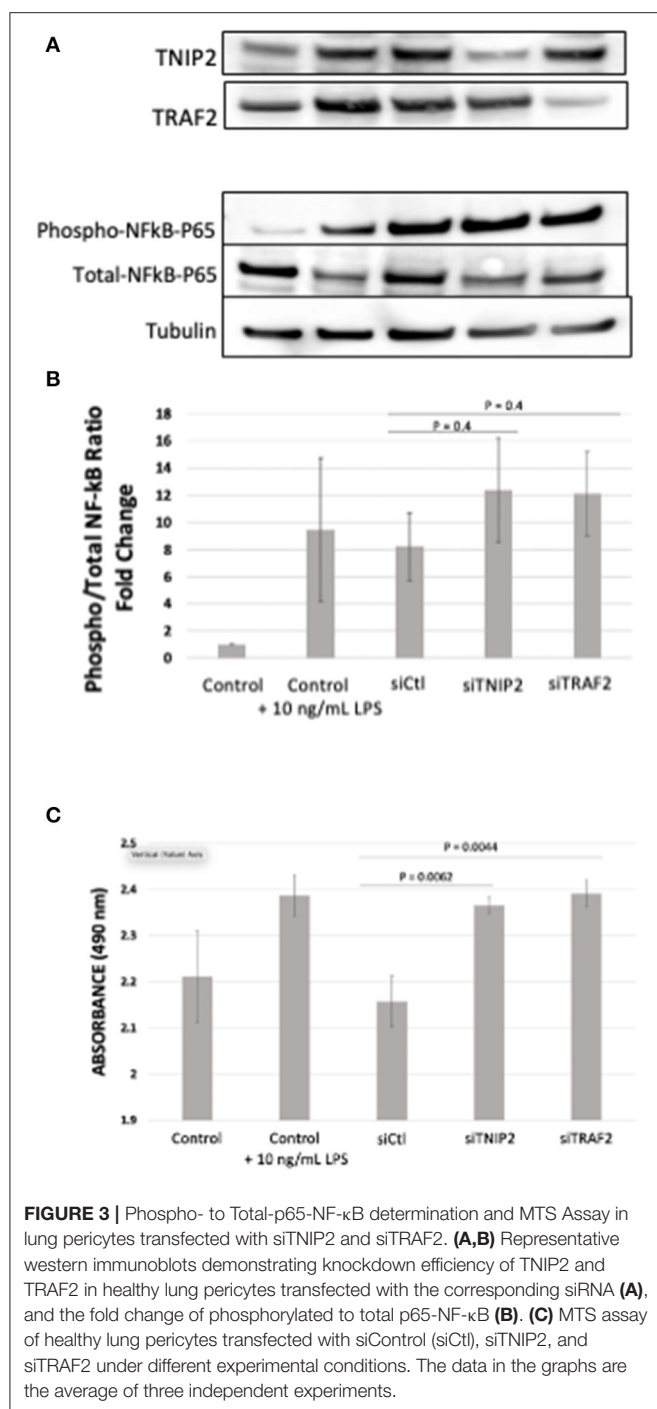
To evaluate the biological consequences of deleterious variants affecting *TNIP2* and *TRAF2*, gene silencing via siRNA transfection in human-derived pulmonary pericytes was performed. We chose to study pericytes because these cells have been shown to contribute to progressive small vessel loss and muscularization through hyperproliferation and inflammatory cytokine production (26, 27, 36). A query of our recently published pericyte RNA-seq analysis confirmed that both *TNIP2* and *TRAF2* are expressed in healthy and PAH pericytes, although no significant differences in expression were observed (26). Transfection of siTNIP2 and siTRAF2 resulted in >50% reduction in corresponding protein levels (**Figure 3A**). To determine whether *TNIP2* and *TRAF2* knockdown led to inappropriate activation of NF- κ B, we used western immunoblotting to probe whole cell lysates for phospho-p65-NF- κ B, the active form of p-65-NF- κ B that becomes activated to trigger transcription of NF- κ B target genes (34, 35). Stimulation with lipopolysaccharide (LPS, a known activator of NF- κ B) triggered an increase in phospho-p65-NF- κ B and was used as a positive control (37). Compared to baseline controls and cells transfected with a non-specific siRNA (siCtl), both siTNIP2 and siTRAF2 pericytes demonstrated a significantly increased phospho-p65-NF- κ B of similar magnitude as that observed with LPS stimulation (**Figure 3B**). To assess the effect of *TNIP2* and *TRAF2* knockdown on pericyte proliferation, we used an MTS assay to quantify cell numbers under different

experimental conditions over 24 h. Compared to the controls, both siTNIP2 and siTRAF2 pericytes demonstrated significantly increased cell proliferation (**Figure 3C**). Interestingly, the degree of proliferation observed in both siTNIP2 and siTRAF2 pericytes was similar to that of healthy pericytes stimulated with LPS.

DISCUSSION

In this study, we utilized a WES custom gene prioritization method optimized to identify potentially pathogenic variants in PAH patients and family members to elucidate possible genetic mechanisms influencing disease development. This approach enabled the discovery of novel variants in *TRAF2* and *TNIP2*, two genes involved in regulating the NF- κ B pathway. We demonstrated that loss of *TNIP2* and *TRAF2* results in inappropriate NF- κ B activation and increased proliferation of healthy lung pericytes through the use of siRNA-based knockdown. A graphical abstract is included in **Figure 4**. To our knowledge, ours is the first report to document a potential link between *TNIP2*/*TRAF2* loss of function and PAH in humans.

Since the description of *BMPR2*, the increasing prevalence of genetic screening has shown that pathogenic variants in this gene are present in ~60% of hereditary PAH cases and 20–25% of idiopathic cases. *BMPR2*, as well as other genes (i.e., *KDR*, *SOX17*, *TBX4*), may also be involved in the development of associated forms of PAH (APAH) (38–41). Multiple groups have



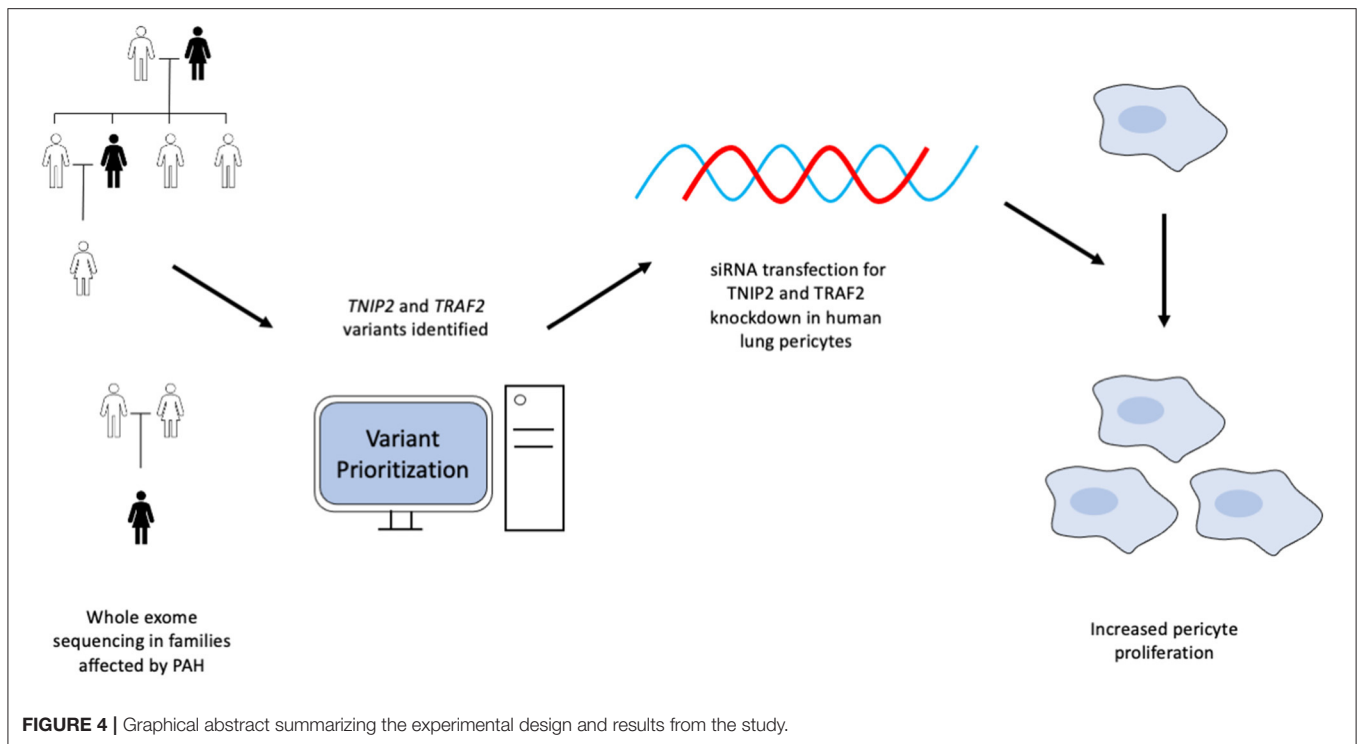
applied custom panels, whole-exome sequencing (WES), or even whole-genome sequencing to elucidate these mechanisms (42), which include variation in gene regulatory elements in addition to the protein-coding sequences themselves (43). Whole-genome sequencing in more extensive and more inclusive cohorts will be vital for uncovering intronic and regulatory genetic sequences that may impact PAH development in patients with unrevealing results upon initial gene panel screening. In many pulmonary hypertension centers, however, genetic analysis is not frequently

offered in patients with PAH associated with other medical conditions (e.g., connective tissue diseases, congenital heart diseases), limiting the data available in this population. This practice will be essential to address future clinical guidelines to deepen our understanding of genetic variation in APAH.

As part of the clinical evaluation, our group routinely performs genetic screening for patients diagnosed with idiopathic, heritable, and associated PAH forms with suspected genetic components. We have developed a custom NGS panel of genes that is updated yearly to reflect published risk genes in PAH and recently reported our experience applying it in 300 patients, some of whom suffered from APAH (17). In all patients with inconclusive results on the gene panel, we move to WES, preferably including parents and other proband relatives, to screen for new risk genes. In recent years, multiple publications have demonstrated the role of genetic variants in the development of many PH subtypes, some of which were previously not known to have a genetic cause. For example, variants in the endoglin gene are linked with PAH associated with connective tissue disease (44), polymorphisms in estrogen-related genes may impact the risk of portopulmonary hypertension (45), and *SOX17* have been implicated in PAH associated with congenital heart disease (41). Also, variants in *CAV1*, *TRPM8*, and *BNP* have been linked with PH secondary to chronic obstructive pulmonary disease (46–48). These findings underscore the importance of expanding our genetic sequencing application to PH populations not traditionally thought to have a genetic predisposition.

In the present study, our WES data suggest an association between *TNIP2* and *TRAF2* variants with the development of APAH. By analyzing exome data of individuals affected with PAH compared to healthy family members, we focused on genes that correlate with phenotypic differences. After our prioritization analysis, we selected the variants in *TNIP2* and *TRAF2* based on the criteria described in **Supplementary Table 1**. These two variants were absent in control population databases, and the majority of the *in silico* pathogenic tools suggest a damaging effect for these variants. Both variants are highly conserved through evolution, highlighting these residues' importance in the affected protein's function. Importantly, these two genes have plausible mechanistic significance as regulators of the NF-κB pathway, previously associated with PAH (49–53). As is the case with the discovery of all novel variants, consideration of pseudocontrol human population gene frequency, conservation in orthologs through evolution, pathogenicity prediction, and biological context was necessary to establish how *TNIP2* and *TRAF2* were involved in PAH.

The family genetic analysis showed that the variant in *TNIP2* is present in both affected individuals and one unaffected member. We suspect two possible explanations for this observation. First, the variant may have incomplete penetrance or cause variable age of disease onset, allowing some carriers to be unaffected for some or all of their lives. Secondly, it may be necessary for an additional disease process to trigger the development of PAH in those who carry this variant. The high-impact nonsense *TRAF2* variant identified in a separate family was only seen in the affected proband, suggesting a *de novo* event



or germinal mosaicism of the parents. Because there was no previous evidence linking pathogenic variants in *TNIP2* and *TRAF2* with PAH, we performed *in silico* analysis to search for relationships between these genes and others implicated in PAH pathogenesis. A study of the protein network by STRING showed that TRAF2 interacts with *CAV1*, a gene linked to hereditary PAH (54). In addition to STRING, there are numerous methods to identify critical gene-gene interactions and visualize pathways affected by genes of interest. Available tools include GSEA from the Broad Institute for gene list analysis and EnrichmentMap to visualize relationships between distinct pathways (55). We believe that investigators should implement these algorithms as part of ongoing genetic studies in PAH.

Inflammation has garnered increasing interest in PAH in recent years (56). Immune phenotypes characterized by differing signatures of inflammatory cytokines offer prognostic and potentially therapeutic significance (12). Antiinflammatory agents such as anakinra and tocilizumab are currently being investigated in PAH clinical trials (57, 58). NF- κ B is known to regulate the expression of a wide array of inflammatory genes, and both human and animal studies indicate that NF- κ B activity is increased in PAH. Core processes in PAH pathophysiology, such as aberrant pulmonary artery wall cell proliferation, apoptosis, and arterial obliteration, are attenuated with NF- κ B inhibition (50). The *TNIP2* gene codes for ABIN2 (A20 binding inhibitor of NF- κ B activation-2), a cytosolic protein that interacts with several membrane receptors (CD40, TNF receptor 1, and toll-like receptor 4) to suppress downstream activation of the NF- κ B pathway (29, 59–61). TRAF2 belongs to a family of adaptor proteins associated with TNF receptors 1 and 2 to modulate TNF stimulation response in immune

cells. Prior work supports TRAF2 as an essential mediator of cell survival and apoptosis through its capacity to activate or inhibit NF- κ B, depending on the cell type (62, 63). Also, TRAF2 is involved in a membrane complex that can recruit caspases to promote apoptosis independent of the NF- κ B pathway (64). Lastly, some authors have noted common hyperproliferative mechanisms driving PAH pathogenesis and cancer development (65), and pathogenic variants in TRAF proteins have been previously described in cancer (66). Specifically, *TRAF2* has suggested roles as an oncogene in epithelial cancer, osteotropic breast cancer and colon cancer (67–69), mainly through the NF- κ B pathway. As evidenced by our *in vitro* studies, decreased expression of *TNIP2* and *TRAF2* can increase NF- κ B activation in human pericytes. More importantly, pericytes demonstrated a change in proliferative activity under these conditions. These data support our hypothesis that pathogenic variants in *TNIP2* and *TRAF2* result in dysregulated NF- κ B activation and could drive abnormal cell proliferation in PAH, as illustrated by the response of lung pericytes transfected with the corresponding siRNAs (Figure 3). However, a manifestation of clinically detectable PAH may require the presence of predisposing conditions (a “second-hit”), such as primary biliary cirrhosis or scleroderma, as seen in our subjects.

Our study has several limitations. First, whole-exome data was only available for a limited number of probands and their relatives, therefore limiting our power to detect novel variants potentially associated with PAH. Secondly, the technology used for our genetic analysis variants does not allow identification of variants outside of exons or exon-intron boundaries, which may hold essential regulatory elements. Additionally, our cell studies used pericytes from human donors, not from the subjects

analyzed in our study. These donors do not have predisposing inflammatory or cardiac conditions, which we believe in having been essential contributors to PAH development in the probands we analyzed. However, our data support the ongoing investigation of NF- κ B as a potential therapeutic target in PAH and show that our genomics-to-bench approach can be useful for expanding our view of PAH genetics.

CONCLUSION

Identifying genetic biomarkers related to PAH holds the potential for a tremendous impact on patient care and the development of personalized treatments. We suggest that WES be considered for patients with associated forms of PAH that do not have causative variants identified with targeted gene panels, especially in cases with family history supportive of inherited disease. Genetic sequencing of family members with and without the disease, when feasible, will provide additional power to link genotype, phenotype, and endotype in PAH. Understanding the genetic underpinnings that predispose individuals to immune dysregulation in the pulmonary vascular wall will be paramount for advancing targeted therapies for this deadly disease. Variants in *TRAF2* and *TNIP2* may contribute to hyperproliferation and increased NF- κ B activation in some individuals with PAH, however further studies are needed to describe the possible relationship of these genes with aberrant immune activity.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: NCBI BioProject, accession no: PRJNA702898 (ACCESSION PRIVATE).

ETHICS STATEMENT

All patients involved in this study gave their informed consent to participate. Samples were obtained from the

Spanish PAH registries (REHAP and REHIPED), and the Ethical Committee approved the project of scientific research of the Hospital Universitario La Paz, Madrid (PI-1210, PI-3996).

AUTHOR CONTRIBUTIONS

JT and VDJP designed and supervised the work. NG, PA, and JT performed all genetic studied and NGS analysis. SP, DC, SA, HP, and AC performed all the functional and *in vitro* studies. ACU, NO, PE, and PL reviewed all the clinical features of the patients. JN performed the microarray analysis. JT, VDJP, SP, NG, and DC wrote the manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmed.2021.625763/full#supplementary-material>

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