



Idiopathic and connective tissue disease-associated pulmonary arterial hypertension (PAH): Similarities, differences and the role of autoimmunity

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ABSTRACT

Pre-capillary pulmonary arterial hypertension (PAH) is hemodynamically characterized by a mean pulmonary arterial pressure (mPAP) ≥ 20 mmHg, pulmonary capillary wedge pressure (PAWP) ≤ 15 mmHg and pulmonary vascular resistance (PVR) > 2 . PAH is classified in six clinical subgroups, including idiopathic PAH (IPAH) and PAH associated to connective tissue diseases (CTD-PAH), that will be the main object of this review. The aim is to compare these two PAH subgroups in terms of epidemiology, histological and pathogenic findings in an attempt to define disease-specific features, including autoimmunity, that may explain the heterogeneity of response to therapy between IPAH and CTD-PAH.

1. Introduction

Pulmonary hypertension (PH) is a hemodynamic condition defined by an increase in mean pulmonary arterial pressure (mPAP) ≥ 20 mmHg at rest, as assessed by right heart catheterization (RHC) [1]. According to the 2022 guidelines of the European Society of Cardiology (ESC) and European Respiratory Society (ERS), PH is classified in five groups based on hemodynamic profile, clinical presentation, pathology, and therapeutic management [1] (Fig. 1). Group 1 PH is characterized by pulmonary capillary wedge pressure (PAWP) ≤ 15 mmHg and a PVR > 2 Wood units, the opposite of post-capillary PH, which is characterized by a PAWP > 15 mmHg and a PVR ≤ 2 [1].

Group 1 PH, that will hereafter be referred to as pre-capillary pulmonary arterial hypertension (PAH), is also histologically characterized by proliferation and remodeling of the pulmonary small arteries, with

progressive occlusion and increased pulmonary vascular resistance (PVR), leading to right ventricle hypertrophy/remodeling, hence right heart failure.

A hemodynamic profile similar to PAH is also observed in chronic thromboembolic PH (group IV) (Fig. 1) but, in this case, small vessel occlusion is due to chronic multiple thromboembolisms [2].

As outlined in Fig. 1, PAH is classified in 6 subgroups, namely: 1) idiopathic PAH (IPAH); 2) PAH associated to diseases such as connective tissue diseases (CTD), portal hypertension, HIV and other infections; 3) heritable PAH (HPAH); 4) drugs and toxins-induced PAH, 5) PAH with features of pulmonary venous/capillary involvement (PVOD/PCH), and 6) persistent PH of the newborn (Fig. 1).

Although they share common hemodynamic features, these 6 subgroups differ in terms of epidemiology, prognosis, and response to therapy, indicating different genetic backgrounds and pathogenic

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mechanisms underlying the PAH onset. For instance, IPAH and CTD-associated PAH (CTD-PAH) are the most common PAH subtype in the Western world [3]. Patients with CTD-PAH have a poorer prognosis than those with IPAH while, among CTD-PAH, systemic lupus erythematosus (SLE)-associated PAH has a better response to therapy than systemic sclerosis (SSc)-associated PAH [4–7].

Aim of this review is to highlight similarities and differences between IPAH and CTD-PAH in terms of epidemiology, histological features, pathogenesis, and role of autoimmunity to gain a better understanding of heterogeneity in the response to therapy.

2. Epidemiology of PAH

Patient registries with PAH have been used as a tool for characterizing the epidemiology, survival and natural history of the disease, and have permitted comparisons between populations in different geographic areas [8].

Even so, the collection of accurate epidemiology data of all forms of PH and its subsets in registries has been a difficult task in this field mainly due to: a) different ethnicity among the groups studied, b) lack of a uniform disease definition, c) differences in the diagnostic approach applied (echocardiography vs RHC) and d) community-based heterogeneity, and e) inclusion of both incidental (patients who have just received a diagnosis) or “prevalent” cases (patients who had previously received a diagnosis) preceded by data normalization [8,9]. Obviously, an ideal analysis should be based on incident cases only [10], instead of performing statistics on both incidental and prevalent.

For the above reasons, epidemiological data on PAH differ markedly worldwide. In western contemporary registries, the IPAH incidence ranges from 0.9 to 2.6 cases per million inhabitants/ year, and the prevalence from 4.6 to 9 cases per million adults [5,10–18]. In Asian countries only few epidemiological data are available about IPAH, not allowing any substantial comparison with other areas worldwide.

Similarly to CTD-PAH, a female preponderance is commonly observed in IPAH, although a certain variability in the female to male ratio has been reported among registries, ranging from 1.5:1 in Europe [19] to 4:1 in the US [14,20].

Surprisingly, the mean age at diagnosis of IPAH has greatly increased over time, ranging between 45 ± 14 and 65 ± 15 years in current registries, compared to 36 ± 15 years in the first registry created in 1981 (U.S. National Institutes of Health Registry) [19]. The reason for this difference is likely the greater diagnostic accuracy and precision than in the past, in defining PAH clinical settings profiles, thereby reducing

cases of wrongly diagnosed IPAH in younger patients.

In Western patients, IPAH is overwhelmingly the most frequent PAH subtype, accounting for 30–48% [11,12,14,17,21], while CTD-PAH is the second most common subtype with an estimated prevalence ranging from 15% to 30% [11,12,14,17,21]. By contrast, the most common subtype in Asia is congenital heart disease (CHD)-PAH (43%), followed by IPAH (35%) and CTD-PAH (19%) [8,22]. Among CTD-PAH, SSc-PAH is the most prevalent in the US (62%) [14] and Europe (61%–76%) [1,15,17], while in Asia high rates of SLE-PAH (51%) have been reported, and SSc-PAH accounts for only 9% [8,22].

3. Pathology of vascular lesions leading to PAH

PAH is characterized by complex arterial lesions involving pre- and intra-acinary pulmonary arteries [23]. Early changes in the endothelial layer are responsible for a specific picture defined as pulmonary vascular disease (PVD) [24]. PVD is characterized by a markedly decreased diffusing lung capacity for carbon monoxide (DLCO), despite an almost normal or only slight decrease of forced vital capacity (FVC), while the FVC/DLCO ratio always remains above 1. At this stage, in most patients PAP is still normal and the patient is usually asymptomatic. Even so, when a FVC/DLCO ratio increase >1.6 occurs, the risk of development of PAH is higher, as suggested by the results of the PHAROS study, showing that a FVC/DLCO ratio >1.6 , along with a DLCO $<55\%$, is highly predictive of PAH development [25,26]. Indeed, in the portion of PVD+ patients that will develop PAH, endothelial changes will evolve dramatically into a continuous anarchic remodeling (vascular remodeling) generating the clinical condition of PAH, one of the major causes of death in CTD-PAH patients.

In patients with PAH, like IPAH and CTD-PAH, vascular remodeling involves all layers of the pulmonary arterial vessels. The histopathological key features of vascular remodeling are listed in Table 1. More specifically, constrictive lesions, such as intimal thickening, media hypertrophy and adventitial thickening, are the result of an imbalance between proliferation and apoptosis of the various cell types forming the vascular walls [23]. Intimal thickening, mainly composed of myofibroblasts and smooth muscle cells (SMC) [27,28] may be eccentric, concentric non-laminar or concentric laminar, the latter being specific to SSc [23]. Media hypertrophy is the result of either expansion of elastic fibers of muscular arteries or SMC hyperplasia and hypertrophy [23]. Adventitial thickening is due to the deposition of collagen by adventitial fibroblasts. Plexogenic lesions consist of slit-like channels of endothelial cells that have undergone proliferation, surrounded by myofibroblasts,

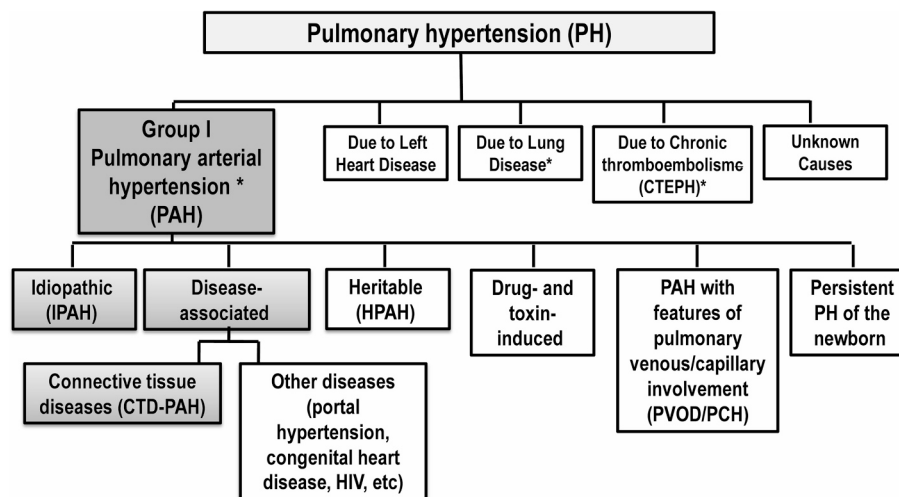


Fig. 1. Clinical classification of pulmonary hypertension according to the 2022 guidelines of the European Society of Cardiology (ESC) and European Respiratory Society (ERS).

PAH subtypes object of this review are in grey boxes. *Pre-capillary PH.

Table 1

Pre-capillary arteries histopathological changes in idiopathic pulmonary arterial hypertension (IPAH) as compared to connective tissue disease (CTD)-associated PAH (CTD-PAH).

Histology findings	PAH subgroup				
	IPAH	CTD-PAH			
		SSc	MCTD	SLE	RA
Intimal thickening	Yes	Yes	Yes	Yes	Yes
Intimal fibrosis	RO	Yes	Yes	Yes	Yes
Plexogenic lesion	Yes	RO	RO	Yes	NR
Media hypertrophy	Yes	Yes	Yes	Yes	Yes
Adventitial thickening	Yes	NR	NR	NR	NR
Inflammatory infiltrates	Yes	Yes	Yes	Yes	Yes
Fibrinoid vasculitis	Yes	RO	RO	Yes	NR
PVOD-like pattern	No	Yes	SF	SF	SF

MCTD, mixed connective tissue disease; NR, not reported; PAH, precapillary pulmonary arterial hypertension; PVOD, pulmonary veno-occlusive disease; RA, rheumatoid arthritis; RO, rarely observed; SF, severe form; SLE, systemic lupus erythematosus; SSc, systemic sclerosis.

SMC and extracellular matrix. Finally, the PVOD-like pattern is characterized by capillary congestion in the alveolar parenchyma, and obliterative fibrosis of venules.

As depicted in Table 1, cellular intimal thickening, media hypertrophy and inflammatory infiltrates [29–31] are all common to both IPAH and CTD-PAH. By contrast, plexogenic lesions and fibrinoid vasculitis are often observed in IPAH and SLE-PAH, but rarely seen in SSc-PAH and MCTD-PAH [29,32,33].

Adventitial thickening occurs more commonly in IPAH than in CTD-PAH [27,34]. On the other hand, arterial intimal fibrosis rarely occurs in IPAH, while it is a common feature of PAH associated to SSc, SLE, MCTD, and rheumatoid arthritis (RA) [29,32].

Similarly, the PVOD-like pattern is rarely observed in IPAH, being more often detected in SSc-PAH and in the most severe forms of CTD-PAH, including SLE, MCTD and RA [29,32] (Table 1).

Overall, the evidence suggests that while PVOD-like pattern and concentric laminar intimal thickening are associated with poor response to therapy, plexogenic lesions and fibrinoid vasculitis, which are more frequently found in IPAH and SLE-PAH, are associated with a more favorable response to therapy, and hence more favorable prognosis.

4. Pathogenesis

4.1. The role of genetics

In the last decades more insight into the pathogenesis of PAH has revealed similarities in genetic background and pathogenic mechanisms between IPAH and CTDs-PAH.

The first evidence for the genetic role in PAH was reported in heritable PAH [35,36], in which a high frequency of bone morphogenic protein (BMP) 2 receptor (BMP2) gene mutations were found. BMP2 belongs to the TGF- β superfamily. Upon BMP2 binding to its ligand BMP, an inhibition of endothelial and SMC proliferation and migration occurs, preventing neointimal formation [37,38]. Some BMP2 gene mutations are therefore associated to the reduction of BMP2 expression and of signaling, together with an increased SMC proliferation and migration. These mutations associated to BMP2 defects of expression (and/or function) have been detected in about 80% of patients with heritable PAH, 10 to 40% of patients with IPAH [39,40], but rarely in CTD-PAH, including RA, and Sjögren's syndrome [41,42]. Even so, PAH is observed in only 20% of BMP2 mutations carriers [43], suggesting that additional, still unknown genetic, epigenetic and/or environmental factors contribute to the development of the disease. In addition, the finding of a BMP2 protein expression reduction in BMP2 wild-type patients with IPAH [44] and SSc-PAH [45,46] suggests that additional mechanisms, besides BMP2 mutations, might decrease BMP2

expression, thus contributing to the PAH onset. This is the case of the loss of SRY-Box Transcription Factor 17 (SOX17) in endothelial cells shown, by an *in vitro* study, to be associated to the reduction in BMP2 expression, and impairment of BMP signaling [47], predisposing to PAH.

Genome- and exome-sequencing studies have revealed additional genes, other than BMP2, carrying rare pathogenic mutations, predisposing to both IPAH and CTD-PAH (mainly SSc- and SLE-PAH) (Table 2) [39,42,48–52]. These include other TGF- β /BMP signaling-related genes (GDF2, ACVRL1, ENG, SMAD9), potassium channel genes, namely KCNK3, KCNA5, ABCC8, and transcription factors such as T-Box factor 4 (TBX4) and SOX17, the latter two involved in lung and vascular development, respectively (Table 2). Even so, some of these genes have not yet been validated (Table 2).

4.2. Molecules contributing to endothelial dysfunction

Endothelial dysfunction plays a pivotal role in the pathogenesis of PAH. The imbalance between vasodilator Vs vasoconstrictor and/or anticoagulant Vs coagulant factors promotes vascular remodeling. More specifically, overproduction of endothelin-1 (ET-1) leading to vasoconstriction, impaired synthesis of the vasodilator nitric oxide (NO) and the prostacyclin (PGI₂)/thromboxane A₂ (TXA₂) imbalance, favoring thrombosis, have been recognized as the main contributors to vascular remodeling in the PAH pathogenesis (Fig. 2).

4.2.1. ET-1

ET-1 is produced by endothelial cells and acts as a potent vasoconstrictor and inducer of smooth muscle cell (SMC) proliferation. These effects occur through ET-1 binding to two endothelin receptor isoforms, namely type A (ETAR) and type B (ETBR). By contrast, the binding of ET-1 to endothelial cell-associated ETBR (the only one to be expressed on these cells) induces the release of NO and PGI₂ [53], hence vasodilation and inhibition of platelet aggregation/activation. Increased levels of ET-1 have been reported in the plasma [54] and lungs [53] of patients with IPAH. Elevated plasma ET-1 levels have also been found in both SSc [55] and SSc-PAH patients [56] and are correlated with impaired right ventricular function [55].

4.2.2. NO, and related molecules

NO is a potent pulmonary vasodilator, and acts as an inhibitor of platelet activation and SMC proliferation [57]. It is produced by the vascular endothelium following NO synthase activation. There are three NO synthase isoforms (NOS), namely neuronal (nNOS), endothelial (eNOS) and inducible NOS (iNOS), the latter being activated during inflammation. NO is rapidly converted to its catabolite (NO_x), which is the most representative marker of NO production. NO_x levels are directly correlated to NO production. In fact, a low NO_x plasma level reflects low NO production and this is the reason why NO_x is lower in IPAH patients than in controls and inversely correlated with mPAP, PVR and survival [58]. The activity of all three NOS isoforms is negatively regulated by asymmetric dimethylarginine (ADMA), an endogenous NOS inhibitor. Plasma levels of ADMA are increased in IPAH patients and associated with an unfavorable pulmonary hemodynamic outcome [59]. Similarly, increased ADMA levels are associated with the presence of PAH, directly correlated with the severity of PAH [60,61] and inversely correlated with 6-min walking test results in CTD-PAH including SSc-PAH [62].

4.2.3. PGI₂/TXA₂

PGI₂ is synthesized and released by endothelial cells. It is the catabolic product of arachidonic acid following the action of two well-known enzymes, namely cyclooxygenase-2 (COX-2) and prostacyclin synthase, the former being up-regulated by inflammation [63] and hypoxia [64] (Fig. 2).

Once released by endothelial cells, PGI₂ acts as a powerful inhibitor of platelet aggregation and also promotes vasodilation. Its activity is

Table 2
Gene variants involved in PAH pathogenesis.

Gene	Encoded protein	Protein function	Validation of alleles variants (AV) predicted to be deleterious		References
			Validation	Comments	
GDF2	BMP-9	Ligand of BMPR2 and ALK1, promoting vascular stability	Not validated	NA	[42,49,146]
ACVRL1	ALK1	Endothelial receptor mediating TGFβ/BMP signaling	Validated by clinical and instrumental data	AV are directly associated to high mPaP and PVR and low 6Mwt in FPAH	[39,42,52]
ENG	Endoglin	TGFβ coreceptor, involved in TGFβ/BMP signaling	Not validated	NA	[42,52]
SMAD9	SMAD8	Downstream effectors of BMP signaling	Validated	Assessed by hereditary analyses in HPAH	[42,52]
KCNK3	TASK1	Potassium channel	Validated	Assessed by hereditary analyses in HPAH	[42,52]
KCNK5	KCNA5	Potassium voltage-gated channel	Validated	AV are directly associated to high mPaP and PVR and low 6Mwt	[39,42,50]
ABCC8 ^a	SUR1	Subunit of the ATP-sensitive potassium channel	Validated	Loss of function (in IPAH e HPAH)	[42,48,50]
TBX4	TBX4	Regulation of embryonic lung development	Not validated	NA	[42,146]
SOX17	SOX17	Regulation of vascular differentiation	Validated	Assessed by hereditary analyses in HPAH	[42,52]
EIF2AK4	GCN2	Downregulation of protein synthesis	Not validated	NA	[42,146]
ATP13A3	ATPase 13A3	Transport of cations across membranes	Not validated	NA	[42]
CAV1	Caveolin-1	Regulation of eNOS activity	Not validated	NA	[42]
KDR ^a	VEGFR2	Regulation of angiogenesis	Validated	Assessed by hereditary analyses in HPAH	[51]
FBLN2 ^a	Fibulin-2	Protection against vascular injury, regulation of blood pressure	Validated	AV are directly associated to high mPaP and PVR and low 6Mwt	[51]
PDGFD ^a	PDGF-D	Regulation of cell proliferation, migration and survival	Validated in PDGF-Knock in mouse overexpressing PDGF in the heart	Vascular smooth muscle cell proliferation and vascular remodeling with wall thickening	[51]

AV, alleles variants; CTD-PAH, connective tissue disease-associated pulmonary arterial hypertension; HPAH, hereditary pulmonary arterial hypertension; IPAH, idiopathic pulmonary arterial hypertension; mPaP, mean pulmonary arterial pressure; NA, not available; PAH, pulmonary arterial hypertension; PVR, pulmonary vascular resistance; 6MWT, six minutes walking test.

^a IPAH only.

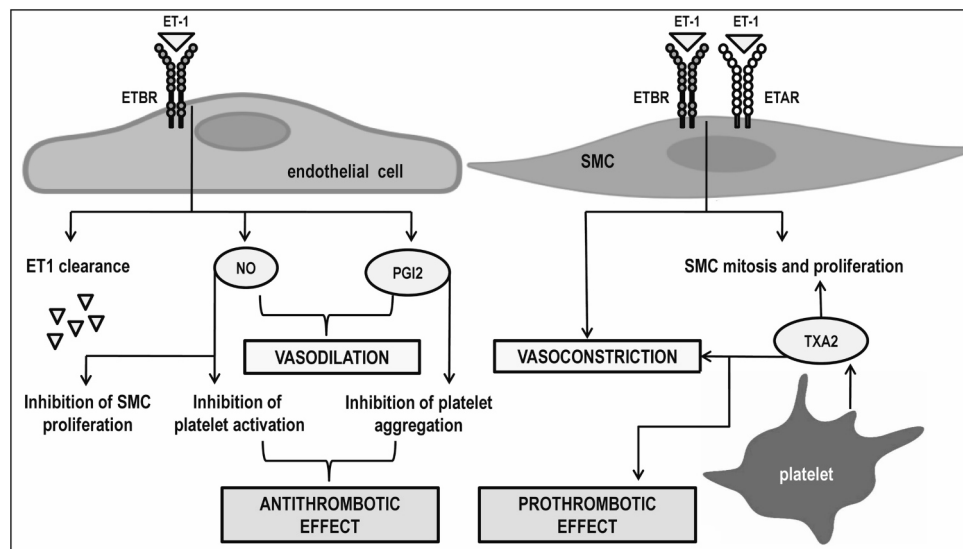


Fig. 2. Vascular tone is maintained by a fine-tuning balance between vasodilator Vs vasoconstrictor and/or anticoagulant Vs coagulant factors. The binding of endothelin-1 (ET-1) to the endothelin receptor type B (ETBR) on endothelial cells mediates ET-1 clearance and stimulates the release of nitric oxide (NO) and prostacyclin (PGI₂). NO inhibits smooth-muscle cells (SMC) proliferation and, along with PGI₂, induces vasodilatation. In addition, both NO and PGI₂ produce antithrombotic effects by inhibiting platelet activation and aggregation, respectively. All these protective effects against PAH are counterbalanced by the action of ET-1 itself as well as of the prothrombotic platelet-derived thromboxane A₂ (TXA₂) on SMC. Indeed, ET-1 binding to its receptors ETBR and ETAR, along with TXA₂ signaling, induces vasoconstriction and SMC mitosis and proliferation.

counterbalanced by the prothrombotic and vasoconstrictor effect of TXA₂, produced by platelets. An imbalance between PGI₂ (decreased production/release) and TXA₂ (increased production/release) [65] occurs in IPAH and CTD-PAH, resulting in vasoconstriction and vascular SCM proliferation [65]. In fact, in IPAH patients, a decreased PGI₂ expression has been detected in small and medium-sized pulmonary arteries walls, while no expression has been found in plexiform and

concentric lesions [66].

4.2.4. Vascular endothelial growth factor (VEGF)

In the intricate network of cytokines involved in the pathobiology of PAH, VEGF, a potent angiogenic cytokine which enhances vascular permeability, occupies a pivotal position.

The VEGF family comprises five members, namely VEGF-A, VEGF-B,

VEGF-C, VEGF-D, and Placental Growth Factor. For each member, several VEGF splice-variant isoforms are generated, which signal through specific distinct receptors known as VEGFR1, VEGFR2, and VEGFR3.

VEGF-A is the most powerful angiogenic cytokine with pro-survival and vascular permeability-enhancing effects, exerted primarily by binding to VEGFR2, and to a lesser extent to VEGFR1 [67].

There is considerable evidence supporting the involvement of VEGF-A in PAH, although also some controversy. Elevated plasma levels of VEGF-A have been documented in IPAH [68–71] and SSc-PAH [70,72,73]. In both diseases, plasma [69] or serum [74] VEGF-A levels are directly correlated with PVR. Adachi et al. [74] reported higher levels of serum VEGF-A165b, an inhibitory splice variant of VEGF-A, in CTD-PAH as compared with control but not in IPAH. Conversely, another study by Suzuki S et al. demonstrated higher plasma levels of VEGF-A165b in IPAH than in controls, but not in CTD-PAH [75]. These discrepancies may be explained by the different biological samples used for VEGF-A165b assay determination, as the former employed serum samples [74] and the latter plasma samples [75].

Placental Growth Factor, a VEGF family member, is a pro-survival and pro-angiogenic factor which exerts these effects through binding to VEGFR-1. Placental Growth Factor is now recognized as a marker for PAH in SSc [76]. In the study by Adachi et al. [74], Placental Growth Factor serum levels were higher in CTD-PAH than IPAH or healthy controls and correlated with 6MWD and BNP levels. In the same study, the Placental Growth Factor serum titer was significantly lower in IPAH than in healthy controls [74]. These results are at variance with those reported by Tiede SL, who found that Placental Growth Factor levels were significantly elevated in both CTD-PAH and IPAH as compared with the non-PH control group [77].

High levels of soluble VEGFR1 (sVEGFR1) have been detected in CTD-PAH and IPAH [70,78]. In SSc patients, sVEGFR-1 was higher in patients with PH than in those without PH [76], as well as in patients that would later develop PAH than in those who would not [70]. On the other hand, VEGFR-2 has been found to be overexpressed in plexiform lesions from IPAH patients [79,80], while low levels of soluble VEGFR-2 have been detected in IPAH and SSc-PAH as compared to controls [71].

4.2.5. CXCL12

CXCL12, also known as stromal cell-derived factor (SDF-1), is a chemokine belonging to CXC family. It is expressed and secreted by various tissues and cell types and regulates several biological processes, including embryogenesis, organogenesis, angiogenesis and vasculogenesis, and tissue repair. CXCL12 acts through its binding to CXCR4 and CXCR7 receptors. The axis CXCL12/CXCR4 is critical for chemotactic endothelial cell migration [81,82]. There is also evidence that links this axis to enhancement of VEGF signaling. Indeed, *in vitro* studies have suggested that CXCL12 promotes VEGF release [83–86] which, in turn, increases CXCR4 expression on endothelial cells [83], sensitizing these cells to CXCL12 signaling. On the other hand, the axis CXCL12/CXCR7 has been suggested to regulate endothelial proliferation and vascular regeneration and repair [82].

Several evidences have suggested that CXCL12 is involved in the pathogenesis of PAH, including IPAH and CTD-PAH. Indeed, increased plasma levels of CXCL12 have been found in both IPAH and CTD-PAH when compared to controls [82,87], with CXCL12 being associated to a poor prognosis [87]. In addition, a marked increase in the expression of CXCL12 and its receptors CXCR4 and CXCR7 has been detected in lung tissue and pulmonary infiltrates from IPAH patients [82,88]. In SSc, high levels of CXCL12 have been detected in microvascular endothelial cells in early stages of the disease [89], and a CXCL12 polymorphism has been associated with PAH [90].

4.2.6. Caveolin-1

Caveolin-1, the major protein constituent of caveolae plasma membrane, interacts (and forms complexes) with a wide range of proteins,

including eNOS, VEGFR [91], and the reactive oxygen species (ROS)-producing enzyme dihydropyridine-adenine dinucleotide phosphate (NADPH) oxidases [92] (Fig. 3A). Under physiological conditions, the binding of caveolin-1 to the aforementioned proteins negatively regulates their activity, and maintains ROS and NO in a steady state, favoring vasodilation through Protein Kinase G (PKG) (Fig. 3A).

Defective caveolin-1 expression causes the persistent production of NO and ROS (mainly superoxide) which rapidly inter-react to form peroxynitrite (Fig. 3B). The latter, in turn, induces the nitration of PKG, its functional impairment and hence vasoconstriction [93].

As caveolin-1 is a down regulator of vascular endothelial growth factor (VEGF) receptor, reduced caveolin-1 expression can mediate an up-regulation of VEGF signaling, favoring PAH (Fig. 3B). Low expression of caveolin-1 has been detected in endothelial cells of lung tissues from IPAH patients [93]. Furthermore, caveolin-1 serum levels were significantly lower in IPAH as compared to chronic obstructive pulmonary disease-PAH and non-PAH subjects [94]. Finally, supporting the role of caveolin 1 in PHA, is that its expression in bone marrow mesenchymal stem cells from SSc patients is significantly lower than in cells from healthy subjects [95].

4.2.7. The Warburg effect

Growing evidence has demonstrated that mitochondrial and intracellular metabolism dysfunction are involved in the pathogenesis of PAH [96]. In this context, a major key role is played by the shift from mitochondrial oxidative phosphorylation to glycolysis in the cytosol, known as the Warburg effect, which leads to the inhibition of apoptosis and promotion of proliferation (Fig. 4). The Warburg effect is mediated by the pyruvate dehydrogenase complex (PDC), the gate-keeping enzyme of glucose oxidation. PDC is a multi-enzyme complex composed of pyruvate dehydrogenase (E1 component), dihydrolipoamide acetyltransferase (E2 component), and dihydrolipoamide dehydrogenase (E3 component), whose activity is regulated by the reversible phosphorylation of the E1 component catalyzed by pyruvate dehydrogenase kinase (PDK). There are four PDK isoforms (PDK1–4) that regulate the activity of PDC by modulating its phosphorylation state. PDK-1 and PDK-2 are strongly up-regulated in IPAH-pulmonary SMC and, to a lesser extent, in endothelial cells, fibroblasts, and inflammatory cells [97], while PDK4 expression has been found in pericytes of PAH patients lungs and correlated with pericyte proliferation and survival [98].

Besides PDK, the activity of PDC can be enhanced by Sirtuin-3 (SIRT-3), the main mitochondrial deacetylase. A down-regulation of SIRT-3 has been found in IPAH-pulmonary arterial SMC compared to normal cells [99]. Additionally, a *SIRT-3* loss-of-function polymorphism has been associated with IPAH but not with CTD-PAH [99].

Another molecule which appears to be strictly related to PAH is transcription factor hypoxia inducible factor 1 α (HIF-1 α), whose activity is regulated by the oxygen state (Fig. 4). Under normoxic conditions, HIF-1 α undergoes a proteasomal-dependent degradation while in hypoxic conditions it accumulates, inducing the expression of target genes, including VEGF, CXCR4 and PDK [100,101] (Fig. 4).

HIF-1 α is expressed in plexiform lesions [102] and a high HIF-1 α expression and transcriptional activity has been found in IPAH-endothelial cells as compared to controls [103].

4.3. The role of autoimmunity

The detection of autoantibodies in IPAH patients sera [104–107] has suggested that autoimmunity might play a role in PAH pathogenesis, similarly to what was observed earlier in CTD-PAH [108–111]. Some autoantibodies (Ab) can be found in both IPAH and CTD-PAH, and some Ab found only in CTD-PAH.

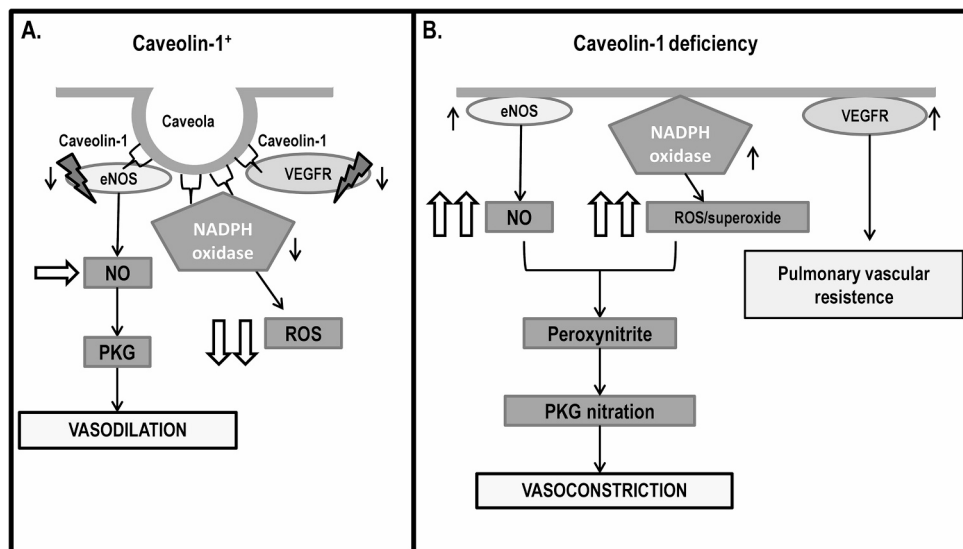


Fig. 3. The role of caveolin-1 in pulmonary arterial hypertension. eNOS, endothelial nitric oxide synthase; NADPH, dihydronicotinamide-adenine dinucleotide phosphate; NO, nitric oxide; PKG, Protein Kinase G; ROS, reactive oxygen species; VEGFR, vascular endothelial growth factor receptor.

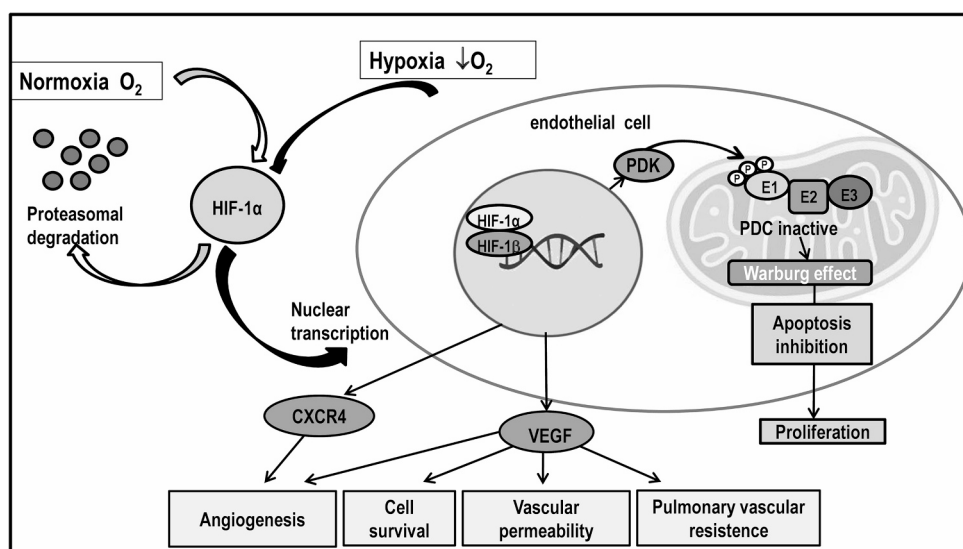


Fig. 4. Hypoxia-inducible factor (HIF) signaling in pulmonary arterial hypertension. Under normoxia, HIF-1α is degraded by the ubiquitin-proteasome system. Under hypoxic conditions, HIF-1α migrates to the nucleus and activates the transcription of genes, including the pro-angiogenic receptor CXCR4, vascular endothelial growth factor (VEGF) and pyruvate dehydrogenase kinase (PDK). VEGF contributes to pulmonary vascular resistance, besides promoting angiogenesis, cell survival and vascular permeability. On the other hand, PDK inactivates the mitochondrial pyruvate dehydrogenase complex (PDC) through the phosphorylation of the E1 component, and induces the Warburg effect, which results in apoptosis inhibition and vascular cell proliferation.

4.3.1. Autoantibodies common to both IPAH and CTD-PAH

4.3.1.1. Anti-endothelial antibodies (AECA). AECA have been detected in 77.8% of CTD-PAH and in 62.1% of IPAH [105]; their more significant antigenic targets were found to be Laminin A/C and tubulin β-chain [112].

Although expressing distinct reactivity profiles [113], IgG AECA from CTD-PAH and IPAH patients have been demonstrated to have a functional role in enhancing pro-adhesive and pro-inflammatory molecules *in vitro* [114].

In IPAH patients only the AECA IgG/IgM ratio is correlated with the severity of the disease [105]. In SSc and MCTD, the prevalence of AECA is higher in patients with PAH than in those without PAH

[105,115,116].

More recently, Ab targeting BMPR2 have been detected in a subset of patients with IPAH but not in CTD-PAH [107]. Ab directed against the type I BMP receptors BMPR1A and Activin Receptor-Like Kinase type 1 (ALK1) have been found in SLE-PAH patients at higher titers than in SLE patients without PAH [117].

Ab to angiotensin receptor type 1 (AT1R) mediate pleiotropic vasoconstrictor and pro-inflammatory effects of angiotensin II. These Ab, together with Ab to endothelin-1 type A receptor (ETAR), have been detected in SSc-PAH at a higher frequency and at higher levels than in IPAH [109]. Moreover, high titers of both Ab have been associated with a worse prognosis in SSc-PAH [109]. In SLE-PAH, anti-ETAR Ab occur more frequently in patients with PAH than in those without PAH, and

their levels appear to be correlated with sPAP, as determined by echocardiography and RHC [110]. Finally, *in vitro* and *in vivo* studies have shown that anti-ETAR and anti-AT1R Ab have also been found to be involved and contribute to the SSc pathogenesis [109,118–120]. More recently, anti-ETBR Ab have been detected in SSc-PAH and IPAH, and higher titers have been found in SSc-PAH patients than in healthy donors [121]. However, the potential pathogenic role of anti-ETBR Ab has not yet been investigated.

4.3.1.2. Anti-fibroblast antibodies (AFA). AFA have been described in SSc [122,123] and have been detected in 30% of patients with SSc-PAH and 40% of those with IPAH [104]. Common AFA target antigens in SSc-PAH and IPAH have been identified by two-dimensional immunoblotting, and include proteins involved in the regulation of cytoskeletal function, cell contraction, oxidative stress, cell energy metabolism, namely vimentin, calumenin, tropomyosin 1, heat shock proteins 27 and 70, glucose-6-phosphate-dehydrogenase [124]. In SSc, a pathogenic role of AFA has been suggested by their ability to promote fibroblast activation and induce a pro-inflammatory and pro-adhesive phenotype [125,126]. Whether AFA can exert similar functions in IPAH remains to be established.

4.3.2. CTD-PAH associated autoantibodies

4.3.2.1. Anti-centromeric proteins (CENP) antibodies. Anti-CENP Ab (ACA), whose main targets are CENP-B and CENP-A, have been detected in 20–30% of patients with SSc and in up to 80% of patients with limited cutaneous involvement [26,123,127,128]. Patients positive for ACA are at higher risk of developing PAH than ACA negative patients [129]. Anti-CENP-B Ab have been suggested to contribute to the pathogenesis of SSc [130–132] by interfering with the normal wound healing mediated by CENP-B in pulmonary arterial SMC, and thus promoting vascular remodeling [130,132,133]. To date, there is no evidence of a potential pathogenic role of anti-CENP-A Ab in SSc. Our group has recently identified anti-CENP-A Ab subgroups whose levels are associated with clinical features of PVD in ACA positive patients [26,128].

More recently, we demonstrated that anti-CENP-A Ab subsets can cross-react with the E2 component of the mitochondrial PDC, the major autoantigen in primary biliary cirrhosis (PBC) [123]. These findings, along with the increased risk of PBC in SSc ACA+ AMA+ patients [134], suggest a pathogenic significance of anti-CENP-A Ab cross-reacting with PDC-E2. Whether these Ab may interfere with the PDC activity, which seems to play a role in PAH, can only be speculated as yet.

4.3.2.2. Anti-U1 ribonucleoprotein (RNP) antibodies. Anti-U1 RNP Ab serve as marker of MCTD but these Ab are detectable in other CTD, such as SSc (6.6%–11.3% of patients) [135,136], and SLE (13–30% of patients) [137]. Several studies have demonstrated a strong association between anti-U1 RNP Ab and PAH in SLE [138–140], as well as in SSc and MCTD [141,142]. Moreover, a recent meta-analysis study has identified anti-U1 RNP Ab as a risk factor for PAH in CTD-patients [143]. Okawa-Takatsuji et al. demonstrated that anti-U1 RNP Ab purified from CTD patients are able to induce the expression of adhesion molecules and MHC class II molecules on pulmonary artery endothelial cells [144,145], suggesting their potential role in triggering vasculopathy.

5. Conclusions

The increasing understanding of the PAH pathogenesis has highlighted the central role of endothelial dysfunction in both IPAH and CTD-PAH diseases. Interestingly, growing evidence is accumulating, demonstrating that, like CTD-PAH, IPAH is characterized by an autoimmune signature, with Ab targeting endothelial cells and fibroblasts. These findings may pave the way to novel therapeutic approaches for IPAH. Despite the many pathogenic factors shared by IPAH and SSc-

PAH, both diseases are characterized by distinctive features, namely the Warburg effect in the former and the high prevalence of Ab directed against ETAR, AT1R and the presence of ACA in the latter. The evidence that these autoantibodies are pathogenic in SSc might explain, at least in part, the poor response to therapy in this disease as compared to IPAH. Therapies targeting these Ab may offer a valuable strategy to counteract PAH development and/or progress in SSc-PAH.

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Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Data availability

No data was used for the research described in the article.

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