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**Survivina e Smac/DIABLO: papel da apoptose na  
HAP experimental**

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experimental PAH**





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Dissertação apresentada à Universidade de Aveiro para cumprimento dos requisitos necessários à obtenção do grau de Mestre em Bioquímica Clínica, realizada sob a orientação científica do Professor Doutor Tiago Alexandre Henriques-Coelho, Professor Auxiliar Convidado do Departamento de Fisiologia e Cirurgia Cardiorácica da Faculdade de Medicina da Universidade do Porto e da Professora Doutora Rita Maria Pinho Ferreira Professora Auxiliar Convidada do Departamento de Química da Universidade de Aveiro.



À minha família, à Alice



**o júri**  
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## palavras-chave

Hipertensão pulmonar, apoptose, survivina, Smac/DIABLO, remodelação vascular

## Resumo

A hipertensão pulmonar (HP), é uma doença multifactorial e progressiva, caracterizada pela vasoconstrição, remodelação vascular pulmonar e trombose *in situ*, com conseqüente aumento da pressão arterial pulmonar e da resistência vascular pulmonar, frequentemente culminando na insuficiência cardíaca direita e morte. A remodelação vascular pulmonar, um dos principais contribuintes para o desenvolvimento e progressão da HP, é caracterizada por uma proliferação celular excessiva e uma reduzida apoptose. No entanto, pouco se sabe sobre os mecanismos moleculares subjacentes a este desequilíbrio. Assim, o presente trabalho teve como objetivo principal a avaliação da contribuição da apoptose na patogénese da HAP, através da análise da expressão cardíaca e pulmonar da survivina e do Smac/DIABLO e a sua associação às alterações hemodinâmicas e morfométricas. Para o efeito utilizou-se um modelo animal de HAP induzida por administração de monocrotalina (MCT). Os resultados demonstraram que os cardiomiócitos se encontravam hipertrofiados 7 dias após a injeção de MCT o que precedeu as alterações hemodinâmicas que se verificaram apenas ao dia 21. Sete dias após a administração de MCT observou-se sobreexpressão de survivina nos ventrículos, tendo esta aumentado progressivamente com a doença. A nível pulmonar, embora sobreexpressa ao dia 7, a expressão de survivin diminuiu com o desenvolvimento da doença. Apesar de sobreexpressa nos ventrículos e no pulmão 7 dias após injeção de MCT, a expressão do Smac/DIABLO diminuiu progressivamente com a HAP. Em conclusão, os resultados sugerem que a desregulação das vias de sinalização nas quais intervêm a survivina e o Smac/DIABLO está relacionada com a remodelação vascular pulmonar e a hipertrofia dos cardiomiócitos em resposta a estímulos apoptóticos.

**Keywords**

Pulmonary hypertension, apoptosis, survivin, Smac/DIABLO, vascular remodelling

**Abstract**

Pulmonary hypertension (PH) is a multifactorial, progressive disease, characterized by vasoconstriction, pulmonary vascular remodelling and thrombosis in situ, which lead to augmentation of pulmonary arterial pressure and pulmonary vascular resistance, culminating in right heart failure and ultimately in death. Pulmonary vascular remodelling, which is one of the major contributors to the development and progression of PH, is characterized by excessive cellular proliferation and reduced apoptosis. However, little is known about the molecular mechanisms underlying this imbalance. So, the present work aimed to study the contribution of apoptosis to the pathogenesis of PAH, through the analysis of cardiac and pulmonary expression of survivin and Smac/DIABLO along hemodynamic and morphometric alterations in an animal model of monocrotaline (MCT)-induced PAH. Results showed that cardiomyocytes were hypertrophied 7 days after MCT injection, preceding hemodynamic alterations which were only present at day 21. Seven days after MCT administration, survivin overexpression was notorious in the ventricles and progressively increased throughout the development of MCT-induced PAH. On the other hand, although increased 7 days after MCT injection, lung survivin expression progressively decreased between the two days. The upregulation of Smac/DIABLO observed in lung, right and left ventricles 7 days after MCT administration, progressively decreased with the disease progression. In conclusion, data suggest that a deregulation in the balance between survivin and smac/DIABLO might be related with pulmonary vascular remodelling and cardiomyocytes hypertrophy in response to apoptotic stimuli.

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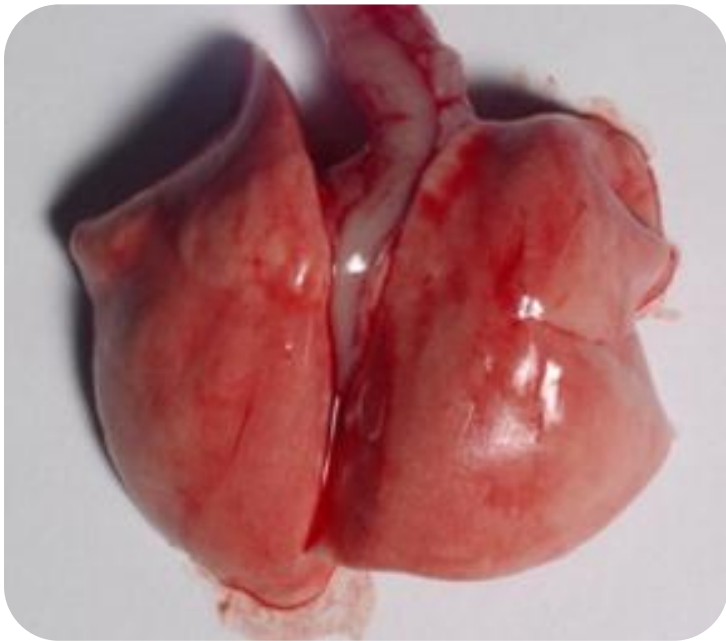
## Abbreviations

<b>[Ca<sup>2+</sup>]<sub>i</sub></b>	Intracellular calcium concentration
<b>5-HT</b>	Serotonin
<b>ADM</b>	Adrenomedulin
<b>AIF</b>	Apoptosis-inducing factor
<b>ALK</b>	Activin receptor-like
<b>Ang-1</b>	Angiopoietin-1
<b>ANP</b>	Atrial natriuretic peptide
<b>Apaf-1</b>	Apoptotic protease activating factor-1
<b>ATII</b>	Angiotensin II
<b>BMP</b>	Bone morphogenetic protein
<b>BMPR</b>	Bone morphogenetic protein receptor
<b>BNP</b>	Brain natriuretic peptide
<b>BW</b>	Body weight
<b>CaM</b>	Calmodulin
<b>cAMP</b>	Cyclic adenosine monophosphate
<b>CCB</b>	Calcium channel blockers
<b>CD</b>	Cluster of differentiation
<b>cGMP</b>	Cyclic guanosine monophosphate
<b>CPI-17</b>	C-kinase potentiated protein phosphatase-1 inhibitor
<b>CREB</b>	cAMP response element-binding protein
<b>CTEPH</b>	Chronic thromboembolic pulmonary hypertension
<b>cyt c</b>	Cytochrome C
<b>EC</b>	Endothelial cell
<b>ECM</b>	Extracellular matrix
<b>E<sub>m</sub></b>	membrane potential
<b>ERK</b>	Extracellular-signal-regulated kinase
<b>ET-1</b>	Endothelin-1
<b>G</b>	Gastrocnemius
<b>GDF</b>	Growth and differentiation factors
<b>HF</b>	Heart failure
<b>HIV</b>	Human immunodeficiency virus
<b>HR</b>	Heart rate
<b>HW</b>	Heart weight
<b>IAP</b>	Inhibitors of apoptosis proteins
<b>IP<sub>3</sub></b>	Inositol 1,4,5-triphosphate
<b>iPAH</b>	Idiopathic pulmonary arterial hypertension
<b>JNK</b>	c-Jun N-terminal kinase
<b>K<sub>v</sub></b>	Voltage-gated potassium channels
<b>L</b>	Lung
<b>LV</b>	Left ventricle
<b>MAPK</b>	Mitogenic-activated protein kinase

<b>MCT</b>	Monocrotaline
<b>MCTP</b>	Monocrotaline pyrrole or dehydromonocrotaline
<b>MLC</b>	Myosin light chain
<b>MLCK</b>	Myosin light chain kinase
<b>MLCP</b>	Myosin light chain phosphatase
<b>NF-AT</b>	Nuclear factor of activated T cells
<b>NF-<math>\kappa</math>B</b>	Nuclear factor kappa B
<b>NO</b>	Nitric oxide
<b>NYHA</b>	New York Heart Association
<b>PA</b>	Pulmonary artery
<b>PAEC</b>	Pulmonary artery endothelial cell
<b>PAH</b>	Pulmonary arterial hypertension
<b>PAP</b>	Pulmonary arterial pressure
<b>PASMC</b>	Pulmonary artery smooth muscle cell
<b>PCH</b>	Pulmonary capillary hemangiomatosis
<b>PDGF</b>	Platelet-derived growth factor
<b>PGI<sub>2</sub></b>	Prostacyclin
<b>PH</b>	Pulmonary hypertension
<b>PK</b>	Protein kinase
<b>PLC</b>	Phospholipase C
<b>P<sub>max</sub></b>	Peak systolic pressure
<b>PPH</b>	Primary pulmonary hypertension
<b>PVOD</b>	Pulmonary veno-occlusive disease
<b>PVR</b>	Pulmonary vascular resistance
<b>ROC</b>	Receptor-operated calcium channel
<b>ROS</b>	Reactive oxygen species
<b>RV</b>	Right ventricle
<b>RVH</b>	Right ventricle/ventricular hypertrophy
<b>S</b>	Septum
<b>SC</b>	Subcutaneous injection
<b>SERT</b>	Serotonin transporter
<b>SMC</b>	Smooth muscle cell
<b>SOC</b>	Store-operated calcium channel
<b>SPAP</b>	Systolic pulmonary artery pressure
<b>TGF</b>	Transforming growth factor
<b>Tib</b>	Tibia
<b>TNF</b>	Tumor necrosis factor
<b>TRAIL</b>	TNF-related apoptosis-inducing ligand
<b>TRP</b>	Transient receptor potential
<b>TXA<sub>2</sub></b>	Thromboxane A <sub>2</sub>
<b>VDCC</b>	Voltage-dependent calcium channel
<b>VEGF</b>	Vascular endothelial growth factor
<b>VIP</b>	Vasoactive intestinal peptide
<b>WHO</b>	World Health Organization







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## **I – Introduction**

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Pulmonary hypertension (PH) has been defined as a sustained elevation of the pulmonary arterial pressure (PAP) to more than 25 mmHg at rest or greater than 30 mmHg with exercise [1]. This is a multifactorial, progressive disease, with substantial mortality and morbidity. Besides elevated PAP, PH is characterized by: (i) increased pulmonary vascular resistance (PVR); (ii) alterations in the extracellular matrix (ECM); (iii) pulmonary vascular remodelling due to proliferation and migration of endothelial cells (ECs) and vascular smooth cells; (iv) progressive pulmonary vascular obliteration; (v) thrombosis and (vi) fibrosis, frequently leading to right-sided heart failure (HF) and death [2]. Untreated patients have a median survival time of 2.8 years [3]. Although it is found in both sexes at all ages, it has a higher incidence in women than men (2:1) and the average age of presentation is 36 years old [4].

Over the years the cellular and molecular mechanisms underlying the PH have been studied, not only to better understand these alterations, but also to develop therapies that may lead to a better prognosis. The therapies implemented at this moment are not entirely efficient because although they increase patients' survival, they do not lead to a cure. Therefore, the study of these mechanisms remains essential to better understand the development and progression of the disease and to plan a targeted and personalized therapy. For that reason, using a monocrotaline (MCT) animal model we studied the contribution of apoptosis to the pathogenesis of pulmonary arterial hypertension (PAH), through the analysis of the apoptotic proteins survivin and Smac/DIABLO.





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## **II – State of Art**

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## 1. Pulmonary Hypertension: Classification, Risk Factors and Diagnosis

In 1891, during an autopsy, the German physician Ernst von Romberg observed pulmonary vascular lesions without a known cause, which he named “pulmonary vascular sclerosis”, a classical description of PH (reviewed in [5]). The term primary pulmonary hypertension (PPH) was first used in 1951 after Dresdale and his associates reported data on 39 patients with unexplained PH (reviewed in [6]). Due to the rising incidence of PH in Europe, in 1973 at an international conference on PPH, World Health Organization (WHO) proposed a classification for PH that included two groups: PPH, a rare disorder that occurred without a known underlying cause, or secondary PH, a disease that developed in the presence of an underlying cause or in the presence of risk factors [6]. Later in 1998, this classification was reviewed at the Second World Symposium on PAH (Evian, France). In the “Evian Classification” PH was divided into five major groups based on pathobiologic and clinical features as well as therapeutic options [7]. During this Symposium, different risk factors for PAH: (i) drugs, toxins and chemicals (e.g. aminorex, meta-amphetamines, benfluores, cocaine, chemotherapeutic agents), (ii) demographic and medical conditions (e.g. gender, pregnancy, obesity), and (iii) diseases (HIV infection, thyroid disorders, collagen vascular disease) were categorized as definitive, very likely, possible or unlikely according to the strength of their association with PH and their probable casual role [7].

The classification was modified once more in 2003 at the Third World Symposium (Venice, Italy). The nomenclature “idiopathic pulmonary arterial hypertension” (iPAH) was adopted instead of PPH. In 2008 at the Fourth World Symposium on PH (Dana Point, California) (Table 1) the classification of PH was updated, and the term “non-PAH” was approved for other categories than PAH [8]. Additionally, left-heart disease PH was subdivided in systolic HF, diastolic HF and valvular heart disease and schistosomiasis was included as a new class of disease-associated PAH [8]. More recently, in March of 2013 the Fifth World Symposium was realized in Nice (France), however the new guidelines have not been published yet.

So, nowadays PH is classified in: (i) PAH; (ii) pulmonary veno-occlusive disease (PVOD) and/or pulmonary capillary hemangiomatosis (PCH); (iii) PH associated with left heart disease; (iv) PH associated with lung disease and/or hypoxia; (v) chronic

thromboembolic PH (CTEPH) and (vi) PH with unclear or multifactorial mechanisms (Table 1).

**Table 1 | Clinical Classification of Pulmonary Hypertension, from the 4th World Symposium on Pulmonary Hypertension (Dana Point, 2008)** (adapted from [8]).

Pulmonary Arterial Hypertension
<b>Group 1: Pulmonary Arterial Hypertension</b>
1.1 Idiopathic
1.2 Familial/Heritable (BMP2, ALK1, endoglin, unknown)
1.3 Drugs and toxin-induced
1.4 Associated with:
1.4.1 Connective tissue disease
1.4.2 HIV infection
1.4.3 Portal hypertension
1.4.4 Schistosomiasis
1.4.5 Congenital heart disease
1.4.6 Chronic haemolytic anaemia
1.4.7 Others
1.5 Persistent pulmonary hypertension of the newborn
<b>Group 1': Sub-class of Pulmonary Arterial Hypertension</b>
1'. Pulmonary veno-occlusive disease and/or pulmonary capillary hemangiomatosis
Non-Pulmonary Arterial Hypertension
<b>Group 2: Pulmonary Hypertension associated with left heart disease</b>
2.1 Systolic dysfunction
2.2 Diastolic dysfunction
2.3 Valvular disease
<b>Group 3: Pulmonary Hypertension associated with lung disease and/or hypoxia</b>
3.1 Chronic obstructive pulmonary disease
3.2 Interstitial lung disease
3.3 Sleep-disordered breathing
3.4 Chronic exposure to high altitude
3.5 Alveolar hypoventilation disorders
3.6 Developmental abnormalities
<b>Group 4: Chronic Thromboembolic Pulmonary Hypertension</b>
<b>Group 5: Pulmonary Hypertension with unclear or multifactorial mechanisms</b>
5.1 Hematologic disorders (e.g. Myeloproliferative disorders)
5.2 Systemic disorders (e.g. vasculitis, sarcoidosis, neurofibromatosis)
5.3 Metabolic disorders (e.g. glycogen storage disease, thyroid disorders)
5.4 Congenital heart disease (other than systemic-pulmonary shunt)
5.5 Others
Legend: BMP2 – bone morphogenetic protein receptor-II; ALK1 – activin receptor-like kinase 1; HIV – human immunodeficiency virus

Besides the classification based on pathobiologic and clinic features, patients with PH can also be classified according to their functional capacity (Table 2). This classification has been adopted from the New York Heart Association (NYHA) for left heart disease [9]. Both clinical and functional classifications are useful for diagnosis, prognosis and therapy guidance of PH.

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**Table 2 | Functional classification of Pulmonary Arterial Hypertension** (adapted from [9]).

Class I:	Patients with pulmonary hypertension without resulting in limitation of physical activity.
	Normal physical activity does not cause undue dyspnoea or fatigue, chest pain or near syncope.
Class II:	Patients with pulmonary hypertension resulting in slight limitation of physical activity.
	They are comfortable at rest, but normal physical activity causes undue dyspnoea or fatigue, chest pain or near syncope.
Class III:	Patients with pulmonary hypertension resulting in marked limitation of physical activity.
	The patient is comfortable at rest, but less than normal physical activity causes undue dyspnoea or fatigue, chest pain or near syncope.
Class IV:	Patients with pulmonary hypertension with inability to carry out any physical activity without symptoms.
	These patient manifest signs of right heart failure. Dyspnoea and/or fatigue may even be present at rest. Discomfort is increased by any physical activity.

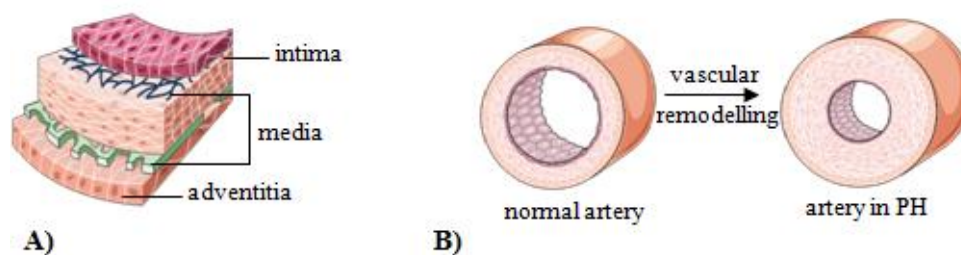
In advanced disease the patient may present non specific symptoms such as: chest pain, breathlessness, fatigue, weakness, palpitations, oedema, ascites and syncope [10, 11]. After careful examination, the clinician can also found: (i) hepatomegaly, (ii) right ventricular hypertrophy (RVH) and right atrial dilation on the electrocardiogram and (iii) increased systolic pulmonary artery pressure (SPAP) on the echocardiography [10-12]. For a differential diagnosis and assessment of disease severity it may be required: (i) antinuclear antibodies, (ii) autoimmune disease markers, (iii) HIV and viral hepatitis screening, (iv) coagulation disorder markers (e.g. protein S and C, von Willebrand factor) and (v) cardiac deterioration markers (e.g. cardiac troponin T, B-type natriuretic peptide (BNP)) [11, 12].

## 2. Pulmonary Vascular Morphology in Pulmonary Hypertension

In order to maintain its main function – gas exchange – pulmonary circulation is a low-pressure, high-flow and low-resistance system. Therefore the walls of the pulmonary arteries (PAs) are relatively thin [13]. The pulmonary vascular wall has 3 layers: (i) an outer adventitia containing fibroblasts, (ii) a media, composed of smooth muscle cells (SMCs) and an elastic lamina (the internal and external, separating the three layers), and (iii) an intima, with a single layer of ECs and a basement of connective tissue (Figure 1) [13, 14].



Pathological changes underlying PH involves vascular remodelling, which is characterized by the thickening of all three layers of pulmonary vessel wall due to the hypertrophy or hyperplasia of one or more cell types as well as increased deposition of ECM components (example: collagens) (Figure 1) [13, 15, 16]. The thickening of media (proliferation and hypertrophy of medial smooth muscle) and/or intima (EC hypertrophy and hyperplasia, oedema and thickening of the endothelial basement membrane) leads to narrowing and occlusion of pulmonary artery, increasing PVR and PAP [13, 15, 16]. In the adventitia layer is also seen the proliferation of fibroblasts and the deposition of collagen [16].



**Figure 1 | Pulmonary Vascular Morphology.** A) The pulmonary vascular wall has 3 layers: an outer adventitia, a media and an intima; B) In PH, vascular remodelling is characterized the thickening of all three layers of pulmonary vessel wall.

In severe stages of PH occurs the formation of a layer of ECM and cells (SMCs or myofibroblasts embedded in mucopolysaccharide) between the internal elastic lamina and the endothelium – neointima – which also contribute to the elevation of PVR [13, 17]. These myofibroblasts express markers such as  $\alpha$ -smooth muscle actin and vimentin but there is no expression of smooth muscle myosin (characteristic marker of highly differentiated SMCs), cluster of differentiation 31 (CD31), CD34 or factor VIII (ECs markers) [17]. Plexiform lesions, another form of vascular remodelling, also predominates in severe stages of PH. They occur due to the disorganised proliferation of ECs supported by a myofibroblast stroma, contributing to the elevation of PVR [13, 18]. These ECs express angiogenic markers such as vascular endothelial growth factor (VEGF) and its receptors [19, 20]. In some forms of PH the release of reactive oxygen species (ROS), growth factors, cytokines and angiogenic factors from the abundant mast cells, T and B lymphocytes and macrophages that surround PAs and arterioles may play a role in vascular remodelling [18]. With time, all of these vascular remodelling events lead to the reduction of PAs' lumen due to its obliteration, resulting in PVR rise and PAP elevation. The augmentation of the PVR results in an overwork of the heart especially of the right ventricle (RV) inducing RVH [21]. Muscle mass accumulates in the RV (through the enhancement of protein synthesis and myocyte hypertrophy)

increasing wall thickness, and it assumes a more rounded shape compressing the left ventricle (LV) [21, 22]. Although in an initial phase the overwork of the RV can maintain the cardiac output, the persistent elevated resistance leads to a progressive dysfunction that frequently culminate in right heart failure [21].

### 3. Pathobiology of Pulmonary Hypertension

#### 3.1. Genetic Associations

Mutations in bone morphogenetic protein (BMP) receptor 2 (BMPRII), and single nucleotide polymorphisms of  $K_V$  (voltage-gated  $K^+$  channels) 1.5, transient receptor potential (TRP) channels and serotonin transporters (SERT) have been implicated in the development of PH.

Bone morphogenetic proteins (BMPs) are multi-functional growth factors that belong to the transforming growth factor (TGF) superfamily. They have multiple important roles in vascular remodelling, including inhibition of apoptosis and SMC proliferation, regulation of vasoactive peptides and growth factors, and modulation of immune cell function [23]. Signalling pathways involving BMPs involve the formation of hetero-complexes between one of type-I receptors and one of type-II receptors [24]. The TGF- $\beta$  superfamily type II receptors are constitutively active serine/tyrosine kinases that initiate intracellular signalling in response to specific ligands [24]. After BMP binding, type-II receptor phosphorylates type-I receptor, activating its kinase domain [24, 25]. In turn, activated type-I receptor phosphorylates the cytoplasmic signalling proteins responsible for TGF- $\beta$  superfamily signal transduction – Smad – that translocates to the nucleus where interacts with various transcription factors, regulating gene transcription (e.g. Bcl-2 and  $K^+$ ) [24, 25]. There is evidence that MAPKs (mitogenic-activated protein kinase), including ERKs (extracellular-signal-regulated kinase), p38, and JNKs (c-Jun N-terminal kinases) might also be activated via TGF- $\beta$  and BMPs [24-26]. Mutations in the gene coding for BMPRII (locus in chromosome 2q31-32) have been found in more than 70% of familiar PAH, and 10% to 40% in iPAH [27]. In spite of the autosomal dominant inheritance of BMPRII gene mutations, this disease has a low penetrance since only approximately 20% of individuals carrying the mutation will develop PAH [27]. One hundred and forty four mutations in BMPRII have been

reported until now, particularly missense mutations that lead to substitution of cysteine residues in the ligand-binding and/or kinase domain of BMPRII, which culminate in the traffic disruption of the mutant BMPRII to the cell surface retaining the receptor in the cytoplasm [28, 29]. On the other hand, in non-cysteine missense mutations in the kinase domain, BMPRII mutants reach the cell surface but fail to activate Smad either by disruption of its serine-threonine kinase activity or the inability to complex with type I receptors [28]. In addition, missense mutations in the cytoplasmic domain attained the cell surface and retained the ability to activate Smad signalling [28]. All these mutations will result in a common feature – gain of function ligand-independent activation of p38 pro-proliferative pathways, inhibiting Smad-dependent signalling with the consequent vascular remodelling observed in familial PAH [28, 30, 31]. Indeed, the decline of BMPRII function in the endothelium might lead to increased apoptosis favouring the development of apoptosis-resistant clones of ECs resulting in the formation of plexiform lesion in PH patients [32]. In addition, mice with BMPRII mutations showed that loss of BMPRII signalling might induce and/or predispose PH with elevation of PAP, wall thickness, pulmonary arterial muscularization and RVH [33, 34].

In addition, serotonin (5-HT) pathway, which has been related with PSMCs hyperplasia, also plays an important role in the development and progression of PH. Serotonin transporter is highly expressed in the PSMCs and have been implicated in vascular remodelling, since patients with iPAH have increased expression of SERT in lung tissues especially in the media of thickened PAs [35]. Furthermore, mice with SERT gene knockout developed less severe hypoxia PH than wild-type control and selective SERT inhibitors attenuate hypoxia- and MCT-induced PH [36-38]. On the other hand, elevated expression of SERT is related with increased severity of PH and transgenic mice selectively overexpressing SERT gene in SMCs developed spontaneously PH in absence of hypoxia, and overstated PH after exposure to hypoxia [39, 40]. Finally, SERT is encoded by a single gene on chromosome 17q11.2 and variants in upstream promoter region have been described. The polymorphism consists in the insertion or deletion of 44pb, designated the long (L) and the short (S) allele, respectively, affecting SERT expression and function. The L-allele induces 2- to 3-fold greater level of SERT gene transcription than S-allele [35, 41]. The L/L genotype might confer genetic susceptibility to iPAH because this homozygous form is much more frequent in these patients (65%) compared with healthy individuals (27%) [35].

Intracellular  $\text{Ca}^{2+}$  concentration ( $[\text{Ca}^{2+}]_i$ ) also has been related with pulmonary vasoconstriction as well as pulmonary vascular wall thickening and remodelling underlying PH. Calcium influx through membrane is controlled by multiple ion channels including: i) voltage-dependent  $\text{Ca}^{2+}$  channel (VDCC) that is regulated through changes in membrane potential ( $E_m$ ); ii) receptor-operated  $\text{Ca}^{2+}$  channels (ROC) which is activated by the interaction agonist-membrane receptor; and iii) store-operated  $\text{Ca}^{2+}$  channels (SOC) that is activated by depletion of  $\text{Ca}^{2+}$  from intracellular stores [42]. In turn, SOC and ROC function is dependent on the expression of TRP channel genes. TRP genes, such as TRPC1, TRPC4, TRPC5, TRPC6 and TRPC7 are expressed in human PSMCs, and TRPC1, TRPC3 and TRPC4 are present in PAECs [43, 44]. Additionally, TRPCs seems to play a role in PH, since PSMCs of iPAH overexpressed TRPC3 and TRPC6 mRNA and protein compared to normal individuals, enhancing cell growth and proliferation [43]. Furthermore, it has been reported that a single-nucleotide polymorphism in the TRPC6 gene promoter is associated with elevation of  $[\text{Ca}^{2+}]_i$  and proliferation of PSMCs in 12% of iPAH patients', predisposing to the disease [45].

### 3.2. Neurohumoral Mechanisms

The primary function of the pulmonary endothelium is to maintain a low PVR through the balance between: (i) vasodilators and vasoconstrictors, (ii) mitogens (proliferation) and growth inhibitors (apoptosis) and (iii) antithrombotic and prothrombotic determinants [46, 47]. In patients with PH, pulmonary endothelial dysfunction appears to play an important role in the changes of pulmonary vasculature [48]. Besides disorganized proliferation of ECs that lead to the formation of plexiform lesions, these cells have an altered production of vasoactive mediators (such as prostacyclin ( $\text{PGI}_2$ ), nitric oxide (NO), endothelin-1 (ET-1), serotonin (5-HT) and thromboxane ( $\text{TXA}_2$ )) affecting vascular tone and SMCs growth which may facilitate the pulmonary vascular hypertrophy and structural remodelling (Table 3) [47, 48]. In addition, endothelial dysfunction might affect the production of anticoagulant factors altering the balance between prothrombotic and antithrombotic determinants, impairing vascular homeostasis that can lead to thrombosis and fibrosis [48].

**Table 3 | Neurohumoral alterations described in the pathobiology of PH**

Mediators	Physiological Effects				Levels in PH		References
	Vascular Dilatation	Cell Proliferation	Platelet Aggregation	Apoptosis	Serum	Lungs	
<b>PGI<sub>2</sub></b>	✓	✗	✗			↓	[49-51]
<b>TXA<sub>2</sub></b>	✗	✓	✓				[49, 50, 52]
<b>NO</b>	✓	✗	✗	✓		↓↑	[46, 53, 54]
<b>ET-1</b>	✗	✓	✓	✓	↑	↑	[55-59]
<b>5-HT</b>	✗	✓			↑	↑	[35, 39, 60-62]
<b>VIP</b>	✓	✗	✗	✗	↓	↓	[63-65]
<b>ADM</b>	✓	✗		✗	↑		[66, 67]
<b>ATII</b>	✗	✓				↑	[68-70]
<b>Ang-1</b>				✗		↑	[71, 72]
<b>VEGF</b>		✓		✗		↑↓	[73-77]
<b>PDGF</b>		✓		✗		↑↓	[78, 79]
<b>ANP, BNP</b>						↑	[80-83]
<b>Apelin</b>	✓	✗		✓		↓	[84, 85]
<b>Ghrelin</b>	✓						[86-88]

Legend: **5-HT**, Serotonin; **ADM**, Adrenomedulin; **Ang-1**, Angiopoiein-1; **ANP**, Atrial Natriuretic Peptide; **ATII**, Angiotensin II; **BNP**, Brain Natriuretic Peptide **ET-1**, Endothelin-1; **NO**, Nitric Oxide; **PDGF**, Platelet-Derived Growth Factor; **PGI<sub>2</sub>**, Prostacyclin; **TXA<sub>2</sub>**, Thromboxane A<sub>2</sub>; **VEGF**, Vascular Endothelial Growth Factor; **VIP**, Vasoactive Intestinal Peptide; ✗, no; ✓, yes; ↓, decrease; ↑, increase

Contraction and relaxation as well as SMCs proliferation and apoptosis are also controlled by multiple protein kinases (PKs) (e.g. PKC, Rho kinase) and cyclic nucleotides (e.g. 3'-5'-cyclic adenosine monophosphate (cAMP) and 3'-5'-cyclic guanosine monophosphate (cGMP)). Indeed, PKA, that can be trigger by PGI<sub>2</sub>, activates K<sub>V</sub> channels in vascular SMCs promoting vasodilatation [89]. On the contrary, PKC, which can be trigger by ATII, ET-1 and TXA<sub>2</sub>, is able to adjust [Ca<sup>2+</sup>]<sub>i</sub> mediating contraction through activation of Ca<sup>2+</sup> channels and inhibition of K<sup>+</sup> channels [90]. PKC is also capable to intervene in contraction through phosphorylation of CPI-17 (C-kinase potentiated Protein phosphatase-1 Inhibitor) which in turn inhibits myosin light chain (MLC) phosphatase (MLCP) increasing MLC phosphorylation [90, 91]. Inhibition of MLCP and CPI-17 phosphorylation is also achieved by Rho kinase, also promoting smooth muscle contraction [92]. Regarding cAMP role, it is hypothesize that vascular relaxation is accomplish by (i) lowering [Ca<sup>2+</sup>] through inhibition of inositol 1,4,5-triphosphate (IP<sub>3</sub>) formation due to phospholipase C (PLC) inhibition, (ii) inhibition of Ca<sup>2+</sup> release from sarcoplasmic reticulum, (iii) stimulation of Ca<sup>2+</sup> uptake and/or extrusion, and (iv) inhibition of Ca<sup>2+</sup> entry [90]. In addition, cAMP through increasing MLCP activity and reducing MLC kinase (MLCK), p42/44, MAPK and Rho kinase activities can regulate vascular tone [91].

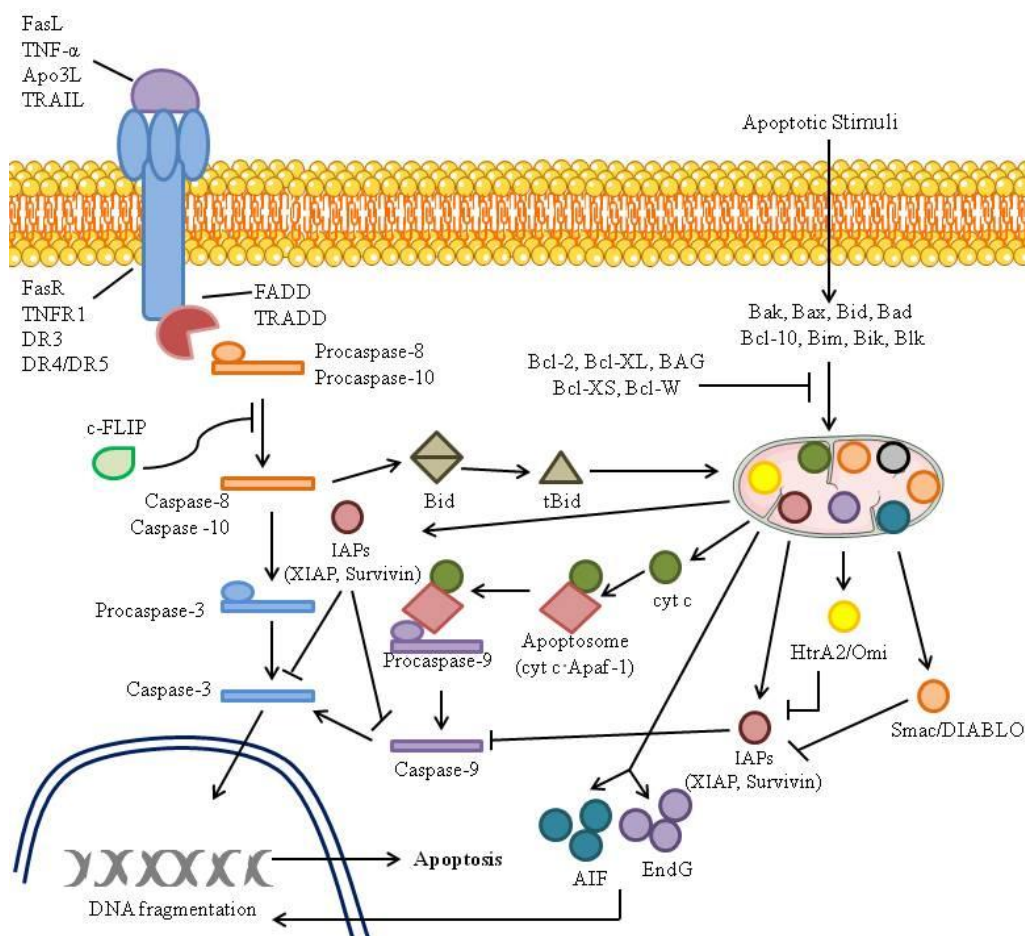
Furthermore, [Ca<sup>2+</sup>]<sub>i</sub> determines smooth muscle contraction, and its rise is the main trigger of pulmonary vasoconstriction as well as SMC migration and proliferation, leading to pulmonary vascular wall thickening and remodelling. Indeed, [Ca<sup>2+</sup>]<sub>i</sub> is an important second messenger for PASMC migration and proliferation, once some signal transduction proteins involved in cellular proliferation are Ca<sup>2+</sup> dependent (e.g. mitogen-activated protein II kinase) [93, 94]. The fact that resting [Ca<sup>2+</sup>]<sub>i</sub> in proliferating PASMC is greater than in growth-arrested cells supports the role of enhanced Ca<sup>2+</sup> influx into the cytoplasmic space is also essential for cell growth and proliferation [95]. Moreover, resting [Ca<sup>2+</sup>]<sub>i</sub> is higher in PASMCs of iPAH patients than in cells of normal individuals, demonstrating once more the role of [Ca<sup>2+</sup>]<sub>i</sub> in PH [96]. In addition, augmentation in [Ca<sup>2+</sup>]<sub>i</sub> leads to elevation of nuclear Ca<sup>2+</sup> concentration which is essential for cell cycle progression, since at least four steps are sensitive to Ca<sup>2+</sup>/CaM (calmodulin) complex activation [97-99]. Additionally, Ca<sup>2+</sup> influx via VDCC (voltage-dependent Ca<sup>2+</sup> channels) activates several transcription factors, such as CREB (cAMP response element-binding protein), NF-AT (nuclear factor of activated T

cells) and NF- $\kappa$ B (nuclear factor kappa B) that are involved in cell proliferation, protein synthesis and inflammation [42].

### 3.3. Role of Apoptosis

Although recent therapies target specific cellular and molecular pathways involved in the pathogenesis of PAH, they are essentially vasodilators, contributing to symptoms relief and improvement but not leading to a better prognosis. So, in the last years investigations have been redirected towards a better understanding of the cellular and molecular mechanisms underlying vascular remodelling, including the role of apoptosis in PH pathogenesis. Indeed, it has been described an imbalance between mitogens and growth inhibitors in PH that along with plexiform lesions and neointima formation highlight a deregulated apoptosis and cell proliferation in the disease.

Apoptosis, or programmed cell death, is important for maintenance of tissue homeostasis through the elimination of harmful cells. In the vasculature, apoptosis modulators are multiple and complex, and might include ROS, NO and ATII systems [100-102]. Apoptosis is characterized by cell shrinkage, extensive protein cross-linking and chromatin condensation (pyknosis) and migration to nuclear membrane [103-105]. In a later stage, although cell membrane remains intact, there is DNA fragmentation (karyorrhexis) into 180 to 200 base pairs promoted by  $\text{Ca}^{2+}$  - and  $\text{Mg}^{2+}$ -dependent endonucleases and formation of apoptotic bodies that are quickly removed via phagocytosis by surrounding cells [105, 106]. Neither the process of apoptosis nor removal of apoptotic bodies is associated with an inflammatory reaction [107, 108]. Apoptosis can occur mainly via two main pathways: intrinsic (or mitochondrial) and extrinsic (or death receptor) ones (Figure 2). Both pathways converge to the same execution pathway that is initiated by the cleavage and activation of execution/effector caspases (cysteinyll aspartic acid-proteases) – caspase-3, caspase-6 and caspase-7 – resulting in the characteristic alterations previously mentioned [109].



**Figure 2 | Apoptosis signalling pathway.** Apoptosis can occur mainly via two pathways: intrinsic and extrinsic ones. In the extrinsic pathway activation of death receptors activates initiator caspase 8 initiating the caspase cascade culminating in apoptosis. The intrinsic pathway is determined by mitochondrial permeability which is regulated through the balance between pro-apoptotic and anti-apoptotic mediators. Activation of pro-apoptotic ones lead to the release of cytochrome c (cyt c), Smac and Omi promoting apoptosis through apoptosome formation and neutralization of inhibitors of apoptosis proteins (IAPs). A crosstalk between these two pathways has been described involving, for instance, Bid cleavage by caspase 8, releasing cyt c from the mitochondria.

The extrinsic pathway involves the activation of transmembrane death receptors which belongs to the TNF (Tumor Necrosis Factor) receptor gene superfamily. These family of receptors is characterized for having a similar cysteine-rich extracellular domain and a cytoplasmic domain called “death domain” [110]. The best-characterized death receptors and corresponding ligands include: i) FasL/FasR, ii) TNF- $\alpha$ /TNFR1, iii) Apo3L/DR3, iv) Apo2L (or TRAIL)/DR4 and v) Apo2L (or TRAIL)/DR5 [111]. After the ligand bind the death receptor, an adaptor protein (FADD for FasL and TRADD for TNF- $\alpha$ ) binds through its own death domain to the clustered receptor death domain



[112, 113]. This adaptor protein also has a “death effector domain” that will associate with an initiator caspase, such as caspase-8 and caspase-10, through dimerization of the death effector domain, activating it by self-cleavage [114, 115]. Then, the initiator caspase (caspase-8) trigger execution pathway through activation of downstream effector caspase, including caspase-3, culminating in apoptosis [115]. This extrinsic pathway can be inhibited through a protein that binds both initiator caspase and/or adaptor protein, such as c-FLIP which can associate with both FADD and caspase-8 inhibiting apoptosis [116, 117].

The intrinsic pathway is determined by mitochondrial permeability which is regulated through the balance between pro-apoptotic (Bcl-10, Bak, Bax, Bid, Bad, Bim, Bik and Blk) and anti-apoptotic (Bcl-2, Bcl-XL, Bcl-XS, Bcl-W and BAG) mediators [109]. When the cell is exposed to some stimuli, such as deprivation of growth factors, exposition to agents that damage DNA or accumulation of unacceptable amounts of misfolded proteins, there is the activation of pro-apoptotic mediators (Bax and Bak), which translocate into the mitochondria and form channels through which cytochrome c (cyt c), Smac/DIABLO, serine protease HtrA2/Omi and other proteins escape to the cytosol [118-120]. Released cyt c activates Apaf-1 (apoptotic protease activating factor-1) and procaspase-9, forming the apoptosome (cyt c/Apaf-1) which is responsible for the activation of initiator caspase 9, culminating in the activation of execution pathway, via activation of caspase-3 [109, 115]. In addition, it was recently demonstrated that besides activating caspases, cyt c can also activate  $K^+$  channels [121]. Smac/DIABLO and HtrA2/Omi have been described to promote apoptosis through the inhibition of IAP (inhibitors of apoptosis proteins) activity, such as XIAP [119, 120, 122]. In addition, alterations in mitochondrial permeability during apoptosis also lead to the release of AIF (apoptosis-inducing factor) and endonuclease G that induce chromatin condensation and DNA fragmentation through a caspase-independent cell death [120].

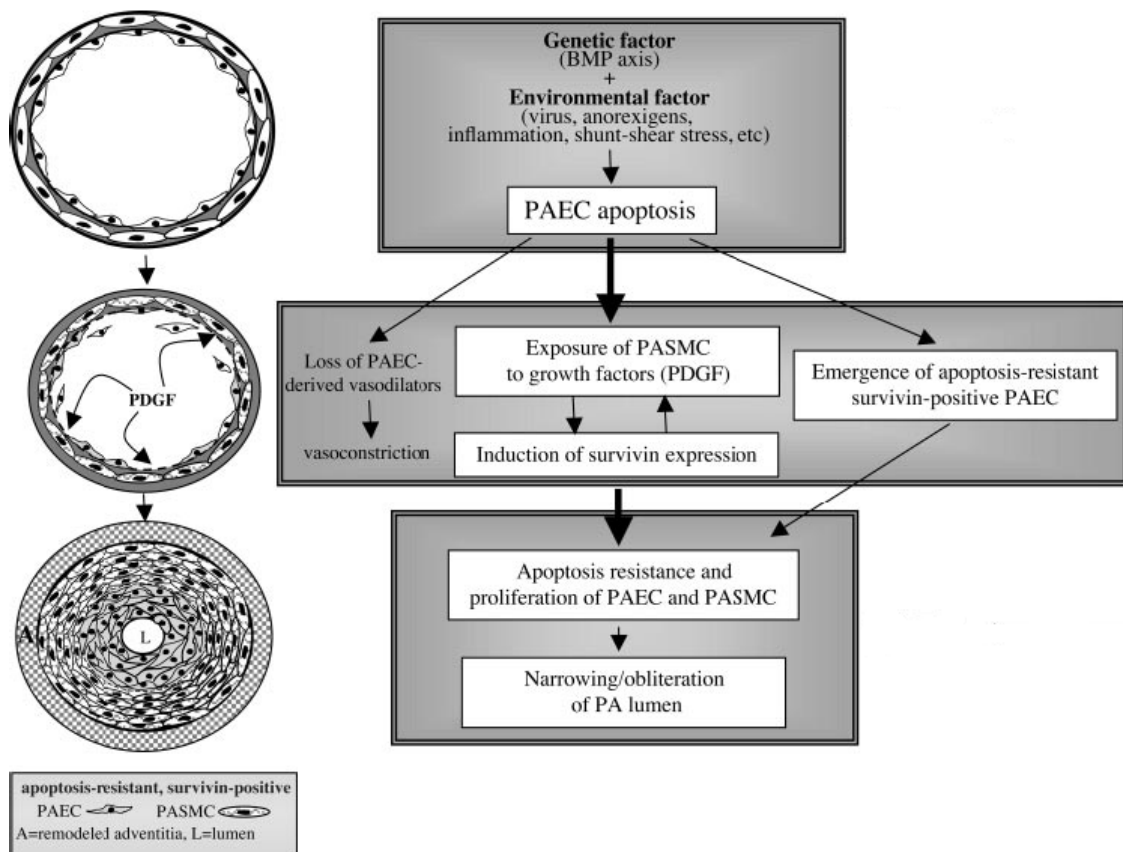
A crosstalk between these two pathways was already described involving, for instance, Bid cleavage through caspase 8 (Figure 2). Truncated Bid translocates to the mitochondria and causes cyt c release through the homo-oligomerization of Bak and/or Bax [123-125]. Recently, it was suggested that apoptosis induced by TRAIL (TNF-related apoptosis-inducing ligand) appears to require the release of mitochondrial Smac/DIABLO [126]. TRAIL induces apoptosis through activation of its death receptors, DR4 and DR5; only, TRAIL-induced apoptosis requires Bax translocation

from the cytosol to the mitochondria. In addition, this translocation of Bax depends on the caspase 8 activation and in turn activated caspase 8 cleave Bid, which translocate to mitochondria releasing Smac/DIABLO [126]. Considering recent works [127, 128] describing TRAIL as a potential therapeutic target in PH, it would be interesting to investigate TRAIL in the development of PH and if its administration can also attenuate or even reverse vascular remodelling seen in PH.

Additionally, maintenance of high concentrations of intracellular  $K^+$  ( $[K^+]_i$ ) is important to the regulation of normal cell volume and suppression of caspases and nucleases [129, 130]. Therefore,  $[K^+]_i$  and  $K^+$  channels seems to be important in apoptosis modulation. Indeed, elevated  $K^+$  loss or  $K^+$  efflux through  $K^+$  channels (via their activation) results in reduction of  $[K^+]_i$  inducing apoptotic volume decrease and activation of caspases and endonucleases that will trigger apoptosis [130, 131]. On the contrary, inhibition of  $K^+$  channels maintain sufficient  $[K^+]_i$  attenuating apoptotic volume decrease and inhibiting apoptosis [131]. Moreover, overexpression of Bcl-2 in rat PASMCs leads to reduction of  $K_V$  channels subunits inhibiting apoptosis [94]. Overexpression of Bcl-2 was also found in the lungs of iPAH patients, which might contribute to the downregulation of  $K_V$  channels in these patients [132].

In PH, the resistance to apoptosis observed in PASMCs seems to result, in part, from the diminished expression of  $K_V$  channels, since PASMCs from iPAH patients' present reduced mRNA expression of  $K_V$  channels subunits, probably due to defects in their gene transcription [133, 134]. Remillard and colleagues [135] also demonstrated that  $K_{V1.5}$  (an  $\alpha$  subunit of  $K_V$  channels) encoding gene of iPAH have multiple single nucleotide polymorphisms which probably contributes to altered function and/or expression of this channels in PASMCs of patients. Additionally, it is believed that certain appetite suppressants (such as aminorex, fenfluramine and dexfenfluramine) are associated with an increased risk of PAH, and they appear to contribute to  $K_V$  channels downregulation, leading to PH [136, 137]. In SMCs, BMP inhibits proliferation and induces apoptosis, on the other hand in ECs it is responsible for cell survival (proliferation) [138, 139]. However, studies had suggested that alterations in BMP axis induce PAECs apoptosis [139]. Besides the genetic susceptibility associated to PH due to mutation in BMP signalling, initial environmental injuries, such as hypoxia, increase flow, inflammation, infections, drugs or toxins, might lead to ECs apoptosis [140-142]. Apoptosis of ECs impairs endothelial function providing conditions for an increased

proliferation of the remaining endothelium and favouring the emergence of apoptosis-resistance and hyperproliferative ECs, that are characteristics of PH later stage [140, 141, 143]. Beyond the release of growth factors and other mediators, the initial apoptosis of ECs leaves SMCs more susceptible to these factors that stimulates proliferation and inhibits apoptosis of SMCs (Figure 3) [144].



**Figure 3 | Apoptotic-based theory for the development of PH.** Although the mechanisms underlying PH need to be better characterized, at the moment an apoptotic-based theory for the development of PH is one of the most accepted in the scientific community. In addition to the genetic susceptibility associated to PH due to mutation in BMP signalling initial environmental factors, such as hypoxia, inflammation or drugs, might lead to PAECs apoptosis. The apoptosis of ECs leads to an endothelial dysfunction providing conditions that increase proliferation of the remaining endothelium and favouring the emergence of apoptosis-resistance, hyperproliferative ECs, expressing survivin, and vasoconstriction, that are characteristic of later stage of PH. In addition, beyond induces the release of growth factors and other mediators the initial apoptosis of ECs leaves SMCs more susceptible to those factors and all this stimulates proliferation and inhibits apoptosis in SMCs. Consequently, the apoptosis resistance and proliferation of both PAECs and PASMCs leads to narrowing and occlusion of pulmonary artery increasing PVR and PAP, frequently culminating in right-sided heart failure and death. (adapted from [142])

### 3.3.1. Survivin and Smac/DIABLO

Survivin is the smallest element of the inhibitor of apoptosis protein (IAP) family [145]. Along with survivin, seven more IAP proteins were identified in humans: XIAP/hILP

(X-chromosome-linked IAP), C-IAP1/HAIP-2 (cellular inhibitor of apoptosis 1), C-IAP2/HAIP-1 (cellular inhibitor of apoptosis 2), ILP2 (IAP-like protein 2), MLIAP (livin), NAIP (neuronal apoptosis inhibitor protein) and BRUCE [146]. IAP family members are characterized by the presence of two to three baculoviral IAP repeat (BIR) domains, which can bind and inactivate caspases and prevent caspase-induced apoptosis; however, survivin only contains a BIR domain [145, 146].

Survivin plays an important role in apoptosis. In response to cell death stimuli, mitochondrial survivin is released to the cytosol, inhibiting apoptosis by either directly or indirectly interfering with caspase function (Figure 2) [147]. Indeed, it was reported that overexpression of survivin suppress apoptosis induced by multiple stimuli including TNF and Fas [148-150]. Some authors refer that survivin can directly bind and inhibit caspase-3, -7 and -9 activity, since survivin co-immunoprecipitated with these, and since apoptosis was suppressed with co-expression of survivin and overexpression of caspase-3, -7 and -9 [150]. On the other hand, others demonstrated that survivin inhibits apoptosis through the interaction with Smac/DIABLO, which is pro-apoptotic factor that promotes caspase activation, binding and inhibiting other IAPs activity [151]. By binding with Smac/DIABLO, survivin delays the release of Smac/DIABLO into the cytoplasm and increases its cytosolic stability, which results in prolonged cell survival [152]. Indeed, Song and colleagues [153] proposed that survivin binds Smac/DIABLO released from the mitochondria, decreasing antagonism of Smac/DIABLO to XIAP, and therefore the free XIAP directly inhibits caspase and consequently apoptosis. Survivin also blocks apoptosis downstream pro-apoptotic mediators Bax, Bik, Bak, and cyt c [148, 150]. Furthermore, it was suggested that under apoptotic stimuli involving mitochondria, survivin is also able to forms a complex with XIAP, enhancing its stability and inhibiting caspases-9 activation [154].

Apart from its role in apoptosis, survivin also regulates cell division and viability. Survivin seems to have a cell-cycle dependent expression in mitosis, being upregulated during G2/M phase of the cell cycle [155]. It was demonstrated to be essential in karyokinesis and cytokinesis through the formation of chromosomal passenger complex along with Aurora B kinase, INCENP and Borealin [156]. Indeed, survivin is necessary in multiple points of mitosis through the regulation of spindle assembly checkpoint, metaphase spindle formation and microtubule stability [155, 157, 158]. Therefore, inhibition of survivin expression/function results in multiple mitotic defects due to cell

cycle arrest, formation of aberrant multipolar spindles, absence of cytokinesis and development of multinucleated polyploidy cells [155, 157, 159].

BIRC5, the gene that encodes survivin, produces five major isoforms of the transcript through alternative splicing of survivin pre-mRNA: survivin (wild type), survivin-2B, survivin- $\Delta$ Ex3, survivin-3B and survivin-2 $\alpha$ . Survivin-2 $\alpha$  seems to support the induction of apoptosis, while survivin (wild type), survivin-3B and survivin- $\Delta$ Ex3 seem to be cytoprotective [160-163]. Some isoforms, such as survivin-3B, appears to also be implicated in cell cycle regulation [163]. It has been hypothesised that the dual role of survivin might be related with its different compartmentalization. Although nuclear survivin is associated to the regulation of cell division, mitochondrial survivin is linked to the inhibition of apoptosis, preventing the activation of effectors caspases which confers resistance to apoptosis [147, 154, 164]. On the contrary, survivin released to the cytoplasm after apoptosis initiation, loses its protective ability against apoptosis, probably due to posttranslational modifications [154, 165]. In a study [166] realized in HeLa cells it was demonstrated that differential ablation of survivin through RNA interference influences cell death. Partial reduction of the cytosolic pool of survivin through RNA interference in association to UVB irradiation resulted in the increased catalytic activity of caspases-3/-7 as well as in the proteolytic cleavage of caspase-9. On the other hand, loss of mitochondrial membrane permeability and spontaneous apoptosis were observed after complete removal of cytosolic survivin. In the same study [166] it was verified that preferential suppression of the nuclear pool of survivin culminated in cell arrest followed by re-entry into the cell cycle and polyploidy along with other mitotic defects.

In the last years, PAH has been considered a cancer like disease since almost characteristics that defines cancer, such as deregulation of growth pathways, angiogenesis, dysfunctional  $K_v$  channels, loss of expression of PGI<sub>2</sub> synthase gene (anti-proliferative) and augmentation of survivin expression have been associated in the disease [167-170]. Indeed, survivin is practically undetectable in differentiated adult tissues, but is highly expressed in the most common human cancers, thus making it an interesting tumor marker [171]. However, little is known about its expression and role in PAH. Until now, it was reported that in an animal model of MCT-induced PH with vascular remodelling and PAs of patients with PH survivin was found overexpressed and was related with the inhibition of apoptosis and stimulation of PSMCs

proliferation [121]. Moreover, lung transfer of survivin gene in experimental animals resulted in pulmonary vascular remodelling while adenoviral gene therapy with a mutant survivin reversed MCT-induced PH through the activation of  $K_v$  channels, induction of apoptosis and suppression of proliferation [121]. Therefore a better characterization of survivin involvement in the development and progression of PAH will potentially open new perspectives on the treatment of PAH.

#### **4. Animal Models for the Study of PAH**

Animal models have allowed the study of the pathogenesis and therapeutic strategies of multiple diseases, including PAH [172]. Although an animal model should mimic the combine characteristics of human disease, including relevant clinical, hemodynamic, biochemical and histopathological features, until now no model is able to mimic all features of the PAH [173]. Different animal models have been used to study the distinct features of PAH, namely the various underlying genetic and environmental causes. MCT and chronic hypoxia are the most frequently used animal models in the study of PH [172]. In addition, transgenic and knockout models have been used to evaluate altered expression of specific genes in the development and/or progression of PH [174, 175]. MCT model has been used for over than 50 years, however the molecular mechanisms underlying MCT-induced PH need to be better clarified [176].

##### **4.1. Monocrotaline Animal Model of PAH**

Monocrotaline (MCT) (or 12 $\beta$ ,13 $\beta$ , -dehydroxy-12 $\alpha$ ,13 $\alpha$ ,14 $\alpha$ -trimethylcrotal-1-enine) is an 11-membered macrocyclic pyrrolizidine alkaloid derived from the stems, leaves and seeds of the *Crotalaria spectabilis* that can be administrated by intraperitoneal (60 mg/kg), subcutaneous (60 mg/kg) or intravenous injection (1-5mg/kg). Although when topically or injected applied it does not cause localized toxicity, MCT is known for its hepatic and cardiopulmonary toxicity, causing lesions in multiple organs after absorption and hepatic bioactivation [15, 177]. In fact, it was described that in chronic exposure or single high doses injections of MCT causes periacinar hepatic necrosis, hepatocyte necrosis, hepatocellular megalocytosis and hepatic fibrosis [178]. However, a single lower-dose injection does not cause hepatic pathology but results in PH after

hepatic bioactivation [178, 179]. In the liver, MCT is activated and converted to reactive metabolites by cytochrome P450 3A, including dehydromonocrotaline also called monocrotaline pyrrole (MCTP) [179]. It is considered that MCTP metabolite is responsible for the toxicity and injuries associated to MCT, since intravenous administration of chemically synthesized MCTP results in lung and/or liver lesions similar to those caused by MCT [15, 179, 180].

It is believed that pulmonary vascular endothelium is the early target of MCT injury probably due to circulatory proximity to the liver and because lungs are the major vascular bed after the liver [179, 181]. In fact, within 7-14 days it was verified increased DNA synthesis and hypertrophy of ECs that occurs after an initial induction of ECs apoptosis within 4 days pos-injection of MCT [181-184]. As early as 4h after injection platelet thrombi can be detected in small arteries, and 4 days pos-injection there is an augmentation in the number of swollen mitochondria as well as ROS generation, which might be endothelial-toxic [184-186]. At day 7, it has been described an early perivascular mononuclear inflammation in MCT-treated rats, mainly due to accumulation of macrophages in the adventitia [187, 188]. MCT treatment also causes the augmentation of extracellular space in the adventitia, probably due to accumulation of oedema fluid from altered microvascular EC permeability [178]. By 8 days after MCT injection there is extension of SMCs into peripheral normally non-muscular small PAs [189]. By 12 days it was described medial hypertrophy of small arteries with a concomitant increase in PAP and resistance [185, 190]. Only after 21 days of MCT injection RVH and dysfunction has been described [190, 191]. Unfortunately, there is evidence that MCT treatment causes myocarditis affecting both right and left ventricles, which will complicate the study of RVH/failure frequently associated with severe PH [192]. In addition, MCT treated rats do not developed plexiform lesions, a characteristic of later stages of human PH [193]. However, combined treatment of MCT with pneumonectomy revealed the development of plexiform lesions [194].

Finally, MCT is also able to induce multiple cellular and molecular alterations at all layers of pulmonary vessels. Lungs of rats exposed to MCT present increased expression of multiple pro-inflammatory cytokines [195-197]. In addition, after MCT treatment it was described an increase of medial SERT and survivin expression, whereas  $K_v$  channels expression, including  $K_v$  1.5 and  $K_v$  2.1, were diminished [37, 121, 198]. BMPRII expression along with its signalling pathway is reduced in MCT-

treated rats, however its restoration does not improve PH [199, 200]. At intima level, MCT cause an augmentation in endothelial expression of ET-1 with a downregulation of ET<sub>B</sub> (endothelin receptor type B) and eNOS (endothelial nitric oxide synthase) expression [201, 202]. In the adventitial layer there is an enhancement of ECM glycoproteins production, such as elastin, fibronectin, collagen and tenascin-C [203, 204].

Taken together, MCT-induced PH differs in some aspects of human PAH, but mimics several of its features, such as hemodynamic repercussions, histological alterations and high mortality. In comparison with other models of PH, MCT model offers technical simplicity, reproducibility and low cost, reasons that justify its general use in the investigation of PH [205].

## 5. Treatment

Until approximately 25 years ago, PAH was considered a fatal disease with poor survival, being its treatment only palliative [4]. The therapies were based on symptomatic relief and were adapted from other pulmonary and cardiac diseases and included oxygen supplementation, anticoagulants, diuretics, digoxin, inotropes and calcium-channel blockers (CCB) (Table 4) [206]. However, the intense research dedicated to the pathobiology of PAH has been helpful for the development of several disease-specific therapies. In fact, treatments targeting specific cellular and molecular pathways involved in the pathogenesis of PAH have being implemented and they allowed symptoms improvement and increased survival (Table 4) [207]. Despite slowing the progression of the disease, treatments do not lead to a cure but diminished the need of surgical procedures (e.g. heart/lung transplantation and atrial septostomy), performed in patients who fail to respond to the pharmacologic therapy [208-211]. As summarized in Table 4, current treatments include: i) supportive therapies, ii) calcium-channel blockers, iii) synthetic prostacyclin and prostacyclin analogues, iv) endothelin-1 receptor antagonists, v) type 5 phosphodiesterase inhibitors and vi) interventional procedures.



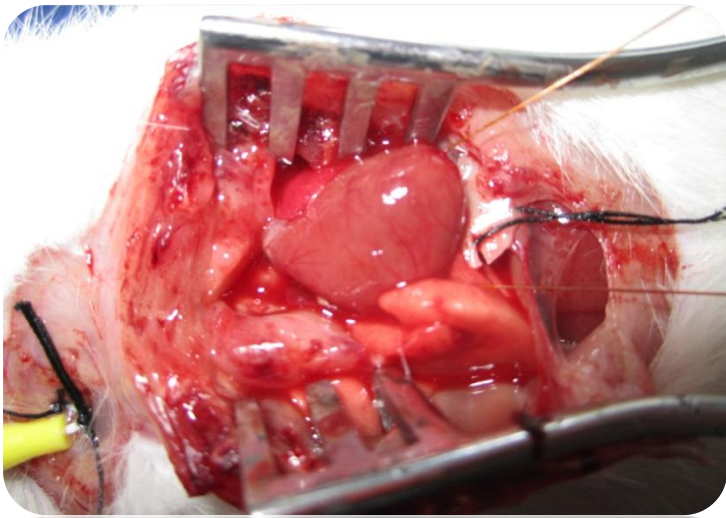
Table 4 | Current therapies for PAH

Drug	Route	Comments and recommendations	Adverse effects and contradictions	References
<b>Supportive therapies</b>				
<b>Anticoagulants</b>	O	Improved survival in small retrospective studies	Bleeding	[206, 207, 212]
<b>Diuretics</b>	O	Used to reduce fluid retention, indicated for RV volume overload	Hypotension	
<b>Oxygen</b>	In	Useful in patients with hypoxia		
<b>Digoxin</b>	O	Used for RV failure with benefit on RV function	Vomiting, arrhythmias	
<b>Inotropes</b>	IV	Mostly applied for end-stage RV failure	Tachycardia	
<b>Calcium-channel blockers</b>				
<b>Nifedipine</b>	O	Helpful as long term in patients with acute vasodilator responsiveness	Peripheral oedema, hypotension	[206, 213]
<b>Diltiazem</b>	O			
<b>Amlodipine</b>	O			
<b>Synthetic prostacyclin and prostacyclin analogues</b>				
<b>Epoprostenol</b>	cIV	Can have beneficial effects for years, used in combination therapy	Nausea, diarrhoea, headache, jaw pain, flushing, catheter-related sepsis/thrombosis	[207, 214-220]
<b>Treprostinil</b>	SC, O, IV, In	Used when oral therapy has failed; short-term benefits	Nausea, diarrhoea, rash, headache, infusion site pain	
<b>Berapost</b>	O	Short-term benefits; used in combination therapy	Flushing, headache	
<b>Iloprost</b>	In, IV	Short-term benefits; used in combination therapy	Headache, jaw pain, cough, flushing	
<b>Endothelin-1 receptor antagonists</b>				
<b>Bosentan</b>	O	Dual ET <sub>A</sub> /ET <sub>B</sub> antagonist, used as a first-line drug; short-term benefits; used in combination therapy	Headache, anaemia, oedema, hepatotoxicity	[207, 221, 222]
<b>Sitaxsentan</b>	O	Selective ET <sub>A</sub> receptor antagonists	Flushing, hepatotoxicity, INR augmentation, warfarin interaction	
<b>Ambrisentan</b>	O		Flushing, hepatotoxicity	
<b>Type 5 phosphodiesterase inhibitors</b>				
<b>Sildenafil</b>	O	Used as an initial agent and in combination therapy	Headache, visual disturbances, nasal congestion	[207, 223, 224]
<b>Tadalafil</b>	O		Nausea, headache, dyspnea	
<b>Interventional procedures</b>				
<b>Atrial septostomy</b>		Decompresses right heart, improves systemic output		[206, 207, 211]
<b>Lung transplantation</b>		Improves survival and quality of life		

Legend: cIV, continuous intravenous; In, inhaled; INR, international normalized ratio; IV, intravenous; O, oral; RV, right ventricle; SC, subcutaneous.

In addition to monotherapy, combination therapy has been an attractive option since it permits to target more than one pathophysiological mechanisms of PAH, allowing the improvement of clinical efficiency and minimizing side-effects [225, 226]. The main clinical combination therapy being considered includes: i) prostanoids and endothelin-1 receptor antagonist (e.g. bosentan + iloprost, bosentan + beraprost, bosentan + epoprostenol), ii) prostanoids and type 5 phosphodiesterase inhibitors (e.g. sildenafil + epoprostenol, sildenafil + iloprost, sildenafil + berapost, sildenafil + treprostinil) and iii) type 5 phosphodiesterase inhibitors and endothelin-1 receptors antagonists (e.g. bosentan + sildenafil, bosentan + tadalafil) [227-233]. Finally, with increasing studies on the pathobiology of PAH multiple potential therapeutic targets have been suggested: i) Rho kinases inhibitors (fasudil), ii) tyrosine kinase inhibitors (imatinib), iii) stimulators and activators of soluble guanylate cyclase (riociguat), iv) VIP, v) ADM, vi) HMG-CoA reductase inhibitors (statins such as simvastatin) and vii) angiotensin II type 1 receptor antagonists (losartan) [66, 234-243] . These potential therapeutic targets were or are being studied in patients with PAH and/or experimental models of the disease.





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**III – Aims**

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Vascular remodelling is an important event in the progression of PAH, characterized by an excessive cellular proliferation and a reduced apoptosis; however, little is known about the molecular events underlying the imbalance between cellular proliferation and apoptosis. So, the aim of the present work was to study the contribution of apoptosis to the pathogenesis of PAH, through the analysis of the apoptotic proteins survivin and Smac/DIABLO, in an animal model of MCT-induced PAH. To achieve this goal, morphometric analysis of heart and lungs of controls and animals with MCT-induced PAH was related with the pulmonary and cardiac expression of survivin and Smac/DIABLO, evaluated by immunohistochemistry and western blot.





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## **IV – Methods and Materials**

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## 1. Experimental Design

In order to study the contribution of apoptosis to the pathogenesis of PAH, an experimental protocol was designed and is summarized in Figure 4.

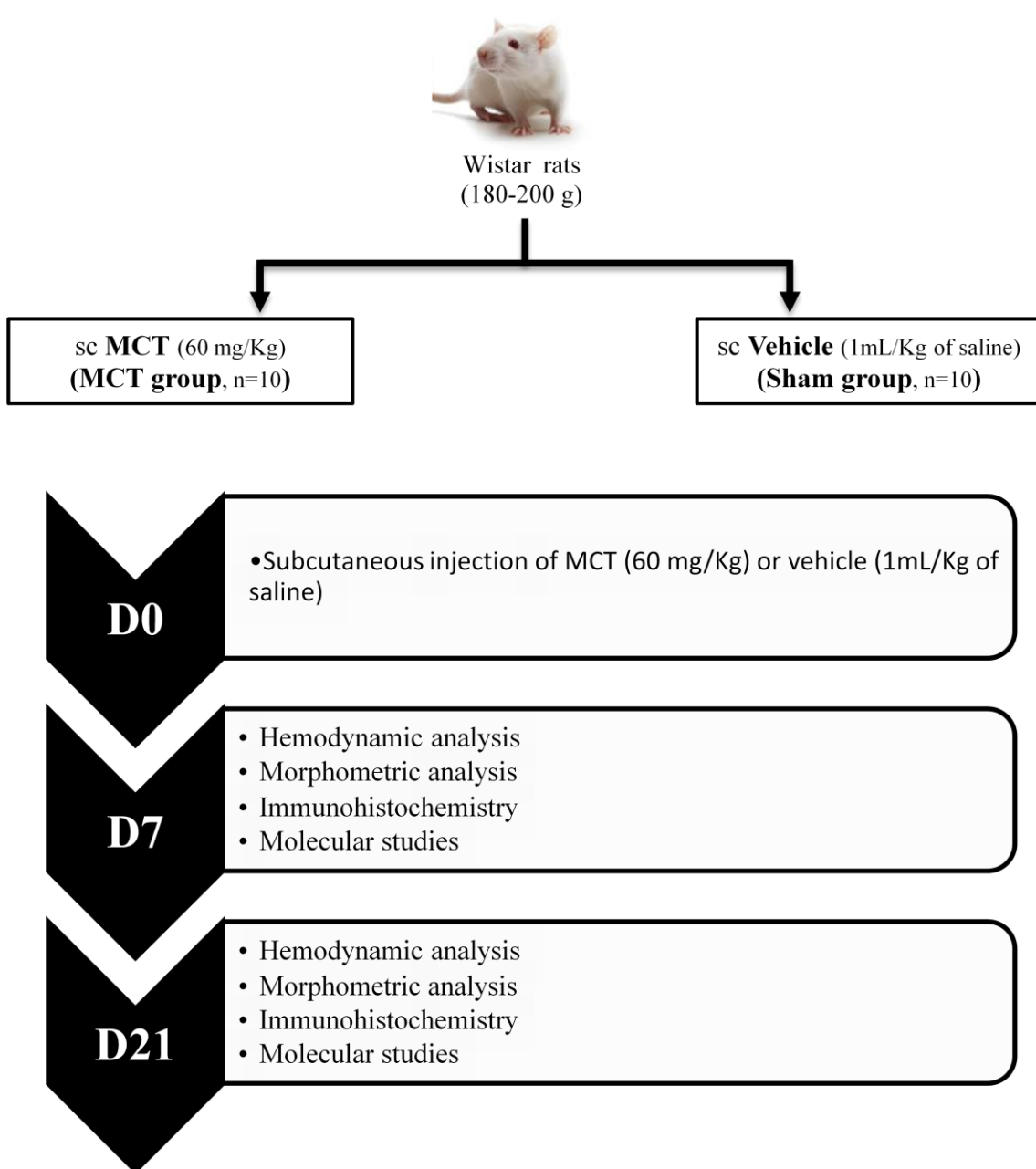


Figure 4 | Experimental protocol design (sc, subcutaneous injection)

## 2. Animal Protocol

Animal experiments were performed according to the Portuguese law for animal welfare (DL 129/92, DL 197/96; P 1131/97) and conform to the National Institutes of Health Guide for the Care and Use of Laboratory Animals (NIH Pub. No. 85-23, Revised 1996). Male Wistar rats (Charles River Laboratories, Barcelona, Spain) weighing 180-200g were housed in groups of 5 rats/cage, in a controlled environment under a 12:12-h light-dark cycle at room temperature of 22°C, with free supply of food and water. Rats randomly received a subcutaneous injection of MCT (60 mg/kg, Sigma-Aldrich, Barcelona, Spain) (MCT group, n=10 per time point) or an equal volume of vehicle (1 mL/kg of saline) (Sham group, n=10 per time point).

## 3. Hemodynamic Analysis

In order to evaluate and confirm the presence and progression of the disease, hemodynamic assessment was performed 7 and 21 days after MCT/vehicle injection. The rats were anesthetized by inhalation of a mixture of sevoflurane (4%) and oxygen, intubated for mechanical ventilation (Dual Mode, Kent Scientific, Connecticut, USA) and placed over a heating pad (body temperature is maintained at 37°C). Under binocular surgical microscopy (Wild M651.MS-D, Leica; Herbrugg, Switzerland), the right jugular vein was cannulated for fluid administration (prewarmed 0.9% NaCl solution) to compensate for perioperative losses. The heart was exposed by a median sternotomy and the pericardium was widely opened. Right ventricular hemodynamic function was measured with pressure-volume (PV) catheter (PVR-1045, Millar instruments, Houston, USA). Data was continually acquired (MPVS 300, Millar Instruments, Houston, USA) and digitally recorded at 1000Hz (ML880 PowerLab 16/30, Millar TM instruments, Houston, USA). After complete instrumentation, the animal preparation was allowed to stabilize for 15 minutes. Hemodynamic recording were made under basal with respiration suspended at end-expiration. Heart rate (HR) and RV peak systolic pressure ( $P_{\max}$ ) were obtained and analysed using PVAN 3.5 (Millar Instruments, Houston, USA).

#### 4. Tissue Preparation for Morphometric, Immunohistochemistry and Molecular Analysis

The heart, lungs and right *gastrocnemius* muscle were excised and weighted. The right tibia was also excised and its length was measured with a millimetric ruler. Under binocular magnification ( $\times 3.5$ ), the RV free wall was dissected from the left ventricle (LV) + septum (S) and weighted separately. Heart, lungs, RV and LV + S weights were normalized to body weight (BW) and *gastrocnemius* weight was normalized to tibia length. Samples from RV, LV and lung were fixed and included in paraffin for light microscopy, or frozen with liquid nitrogen for molecular studies.

#### 5. Morphometric Analysis

Samples of RV, LV (midway between the apex and base) and lung were fixed in 4% (v/v) buffered paraformaldehyde followed by dehydration with graded ethanol, diaphanization with xylene and included in paraffin blocks. Serial sections (4  $\mu\text{m}$  thickness) of paraffin blocks were cut by a microtome (RM2125RTS, Leica, Nussloch, Germany) and mounted on silane-coated slides. The slides were dewaxed in xylene and hydrated through graded alcohols. Sections were stained for haematoxylin-eosin by immersing slides in Mayer's haematoxylin solution for 5 minutes followed by immersion in aqueous eosin solution for 5 minutes. Slides were still submitted to graded alcohols and xylene and mounted with Entellan. Studied samples were observed at light microscopy (Dialux 20, Leitz, Wetzlar, Germany), photographed with a digital camera (XC30, Olympus, California, USA) and measured with a digital image analyzer (cell<sup>^</sup>B life science basic imaging software, Olympus, California, USA). Five images with random microscopic fields (magnification of  $\times 400$ ) were obtained from each section to compensate for variations within sections. Only round to ovoid muscle fibers with a nuclear profile were counted to measure the cardiomyocytes surface area (CSA). Around 500 cardiomyocytes per group per time point were analyzed. On pulmonary specimens, external diameter and medial wall thickness in muscular arteries (20-25 arteries/lung) were analyzed. Orthogonal intercepts were used to generate eight random measurements of external diameter (distance between the external lamina) and sixteen random measurements of medial thickness (distance between the internal and external lamina). For each artery medial hypertrophy was expressed as follows: % wall thickness =  $[(\text{medial thickness} \times 2) / (\text{external diameter})] \times 100$ .

## 6. Immunohistochemistry

Immunohistochemistry was performed to determine survivin and Smac/DIABLO expression in the RV. Sections (4 µm) were placed on SuperFrost®Plus slides (Thermo Scientific, Massachusetts, USA). After deparaffinization and rehydration, slides were subjected to heat induced antigen retrieval by immersion in 10 mM sodium citrate buffer (pH 6.0) in the microwave for 30 minutes. Peroxidase activity was blocked by 3% hydrogen peroxide for 10 minutes. Blockage of non-specific binding was performed with 5% normal goat serum (G9023-10mL, Sigma-Aldrich, Barcelona, Spain) in TBS-T (100 mM Tris, 1.5 mM NaCl, pH 8.0; 0.1% Tween-20) for 1 hour at room temperature followed by 15 minutes of washes in TBS-T. Sections were encircled with a pap pen (Vector Laboratories, California, USA) to prevent splitting leakage of the flowing incubation solutions. Endogenous avidin-biotin expression was blocked using an endogenous avidin + biotin blocking system (ab3387, abcam, Cambridge, UK) according to manufacturer's instructions, being followed by incubation with the primary antibodies (1:500 dilution; rabbit anti-Survivin (ab469, abcam, Cambridge, UK) or dilution 1:250; rabbit anti-Smac/DIABLO (ab8115, abcam, Cambridge, UK)) overnight at 4°C. After incubation with primary antibodies, slides were washed 3 times, 5 minutes each with TBS-T and incubated with goat anti-rabbit IgG secondary antibody (1:250 dilution; ab6720, abcam, Cambridge, UK) for 2 hours at room temperature. Slides were submitted to another 3 washes in TBS-T prior being incubated with Streptavidin protein, HRP (1:1000 dilution; ab7403, abcam, Cambridge, UK). To visualize the peroxidase activities in sections 3,3-diaminobenzidine (DAB, ab94665, abcam, Cambridge, UK) was used. Finally, slides were counterstained with Mayer's haematoxylin, submitted to graded alcohols and xylene and mounted with Entellan. Negative control reactions included omission of the primary antibody. The slides were observed and photographed with a microscope (Dialux 20, Leitz, Wetzlar, Germany) under ×400 magnification. Survivin and Smac/DIABLO expression was qualitatively determined as positive (cytoplasmatic staining) or negative in around 500 cardiomyocyte per group per time point.

## 7. Western Blot

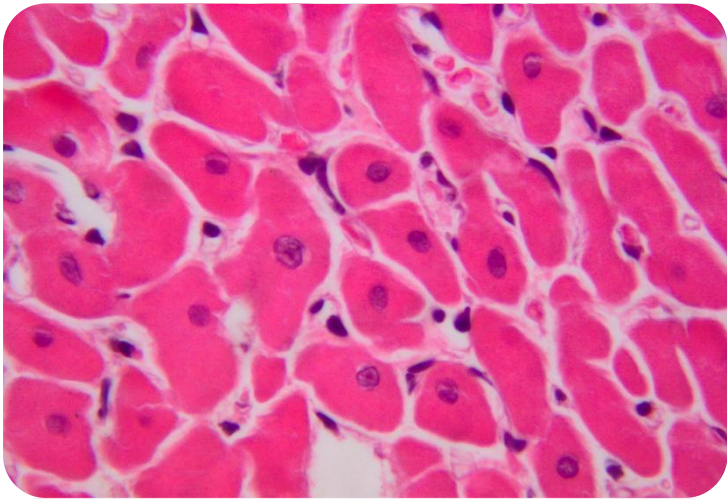
RV, LV and lung samples (n=6 per group per time point) previously frozen with liquid nitrogen were homogenised in phosphate buffer (13 mM  $\text{KH}_2\text{PO}_4$ , 54mM  $\text{NaHPO}_4$ , pH 7.4) (in the proportion of 1:20) with a Bio-Gen PRO200 homogeniser (Pro 200, Pro Scientific, Connecticut, USA). Total protein concentration was spectrophotometrically determined with the colorimetric method RC-DC protein assay (Bio-Rad, California, USA). The optic density was determined at 750nm in a microplate reader (UVM340, Asys, Cambridge, UK). Simultaneously, a calibration curve was performed using different concentrations of bovine serum albumin (BSA).

Equivalent amounts of total protein from the homogenised RV, LV and lung of each group were electrophoresed on a 12.5 % SDS-PAGE at 200 V at room temperature as described by Laemmli [244]. Proteins were electrotransferred to a nitrocellulose membrane (0,2  $\mu\text{m}$ , Bio-Rad, California, USA) in 25 mM Tris, 192 mM and 20 % methanol at 200 mA. Successful transfer was confirmed by staining the membranes with Ponceau S. Nonspecific binding sites were blocked with 5 % (w/v) dry non-fat milk in TBS-T (100 mM Tris, 1.5 mM NaCl, pH 8.0 (TBS) and 0.5 % Tween 20). Membranes were incubated with primary antibody (rabbit anti-Survivin (ab469, abcam, Cambridge, UK), 1:1000 dilution or rabbit anti-Smac/DIABLO (ab8115, abcam, Cambridge, UK), 1:500 dilution) overnight at 4°C with agitation. Afterwards, blots were washed in TBS-T, incubated with a secondary antibody (LI-COR IRDye@ 800CW, Nebraska, USA) and washed in TBS-T. Immunoreactive bands were observed under fluorescence using an Odyssey system (LI-COR Odyssey, Nebraska, USA) and the results were analysed with Quantity One software v. 4.6.3 (Bio-Rad, California, USA)

## 8. Statistical Analysis

Statistical analysis was performed using GraphPad Prism software v. 5.0 (GraphPad Software, California, USA). All data are presented as mean  $\pm$  SEM and were compared using Two Way ANOVA. When treatments were significantly different, Students-Newman Keuls post-hoc test was selected to perform pairwise multiple comparisons. Results were considered significantly different when  $p < 0.05$ .





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**V – Results**

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## 1. Hemodynamic Evaluation and Morphometric Analysis

In order to confirm and evaluate RV dysfunction and the development of PAH through MCT injection, hemodynamic evaluation and morphometric analysis were performed 7 and 21 days after MCT injection. As represented in Table 5, at D7 no significant hemodynamic alterations were noted. However, at D21, RV peak systolic pressure in the MCT treated group was significantly enhanced when compared with Sham group.

**Table 5 | Hemodynamic evaluation of MCT-induced PAH.**

	D7		D21	
	Sham	MCT	Sham	MCT
<b>Heart rate (bpm)</b>	349 ± 22	357 ± 10	340 ± 14	376 ± 23
<b>RV P<sub>max</sub> (mmHg)</b>	27.3 ± 2.2	31.5 ± 1.5	26.3 ± 1.2	38.9 ± 2.7 *

Data are present as mean ± SEM. Sham, sham group; MCT, monocrotaline group; P<sub>max</sub>, peak systolic pressure; RV, right ventricle. \*p < 0.05 vs Sham of the same day

**Table 6 | Morphometric alterations in MCT-induced PAH**

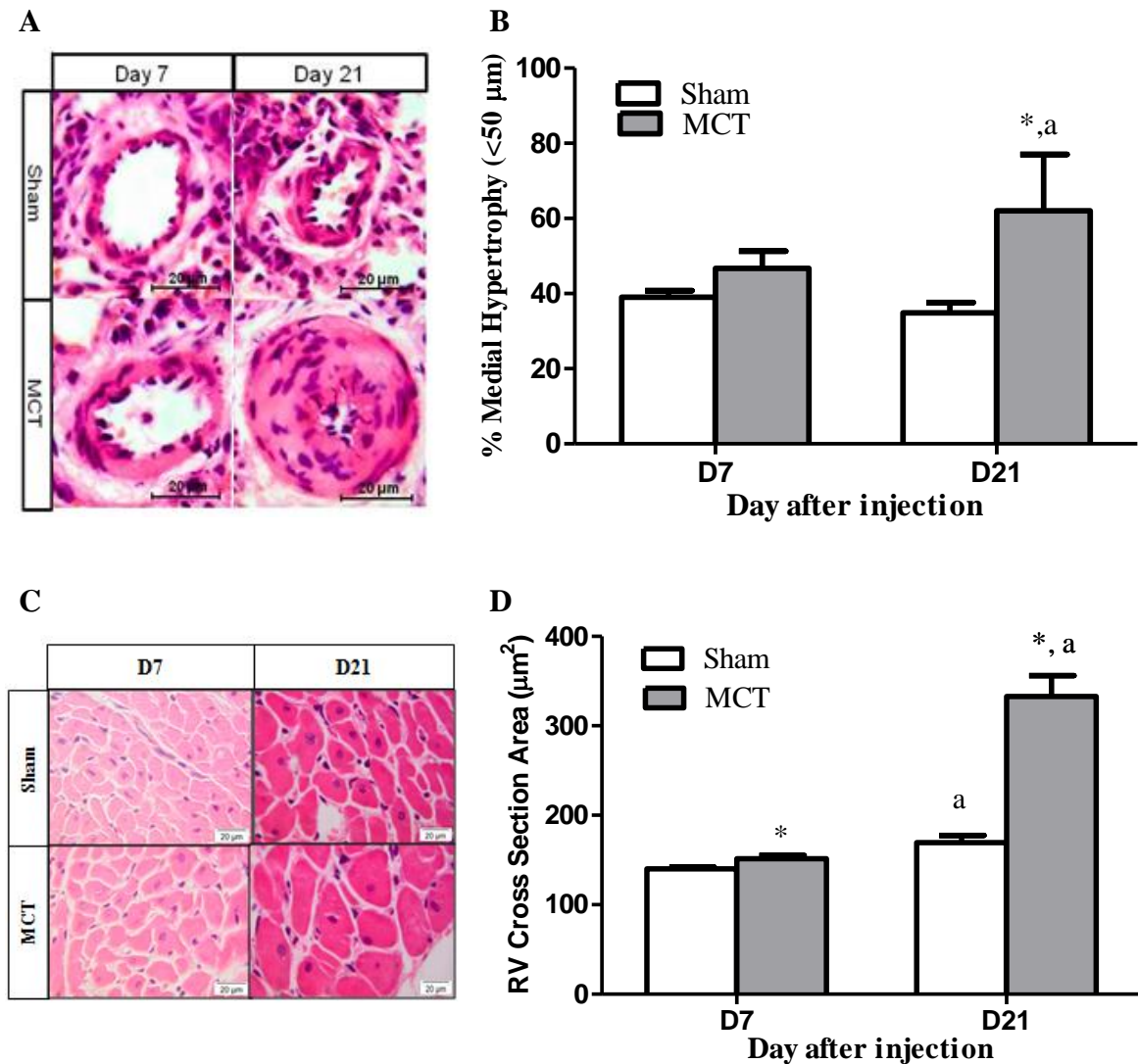
	D7		D21	
	Sham	MCT	Sham	MCT
<b>Body weight (g)</b>	230.4 ± 8.2	214.6 ± 3.2	290.6 ± 4.5 <sup>a</sup>	260 ± 4.9 <sup>*, a</sup>
<b>HW/BW (g/Kg)</b>	3.031 ± 0.079	3.129 ± 0.061	2.794 ± 0.069	3.282 ± 0.113 *
<b>RV/(LV+S) (g/g)</b>	0.282 ± 0.017	0.332 ± 0.021	0.302 ± 0.012	0.467 ± 0.049 <sup>*, a</sup>
<b>RV/BW (g/Kg)</b>	0.583 ± 0.038	0.332 ± 0.021	0.584 ± 0.013	0.911 ± 0.095 <sup>*, a</sup>
<b>(LV+S)/BW (g/Kg)</b>	2.070 ± 0.058	2.011 ± 0.047	1.957 ± 0.060	1.956 ± 0.042
<b>L/BW (g/Kg)</b>	5.436 ± 0.307	5.896 ± 0.400	4.782 ± 0.323	7.241 ± 0.464 <sup>*, a</sup>
<b>G/tib (g/cm)</b>	0.382 ± 0.012	0.374 ± 0.004	0.440 ± 0.016	0.430 ± 0.008

Data are present as mean ± SEM. Sham, Sham group; MCT, monocrotaline group; HW, heart weight; BW, body weight; RV, right ventricle; LV+S, left ventricle plus septum; L, lungs; G, gastrocnemius; tib, tibia. \*p < 0.05 vs Sham of the same day; <sup>a</sup>p < 0.05 vs D7 of the same treatment group.

The morphometric alterations registered in MCT-induced PAH are summarized in Table 6. At day 21, animals from the MCT group evidenced a significant body weight loss in comparison to the Sham group. There was a significant augmentation of body weight in both groups between days 7 and 21. In addition, there were no significant alterations in the ratio G/tib.

In MCT treated animals, lung weight increased between time-points, and was significantly higher in MCT group than in Sham group. As expressed in Figure 5, at D21 media of pulmonary artery is significantly hypertrophied in MCT-treated animals comparing with Sham group.

The ratio HW/BW was significantly higher in MCT-treated animals compared to Sham on day 21 after injection. Moreover,  $RV/(LV+S)$  and  $RV/BW$  indexes were both significantly increased on day 21 in MCT-treated animals, indicating that these ones developed RV hypertrophy. Also, as expressed in Figure 5, MCT-treated animals developed cardiomyocytes hypertrophy since D7. The RV hypertrophy increased with the progression of the disease since RV cardiomyocytes cross sectional area was significantly higher on day 21 than on day 7. There were no differences in the  $(LV+S)/BW$  parameter in both MCT and Sham groups.



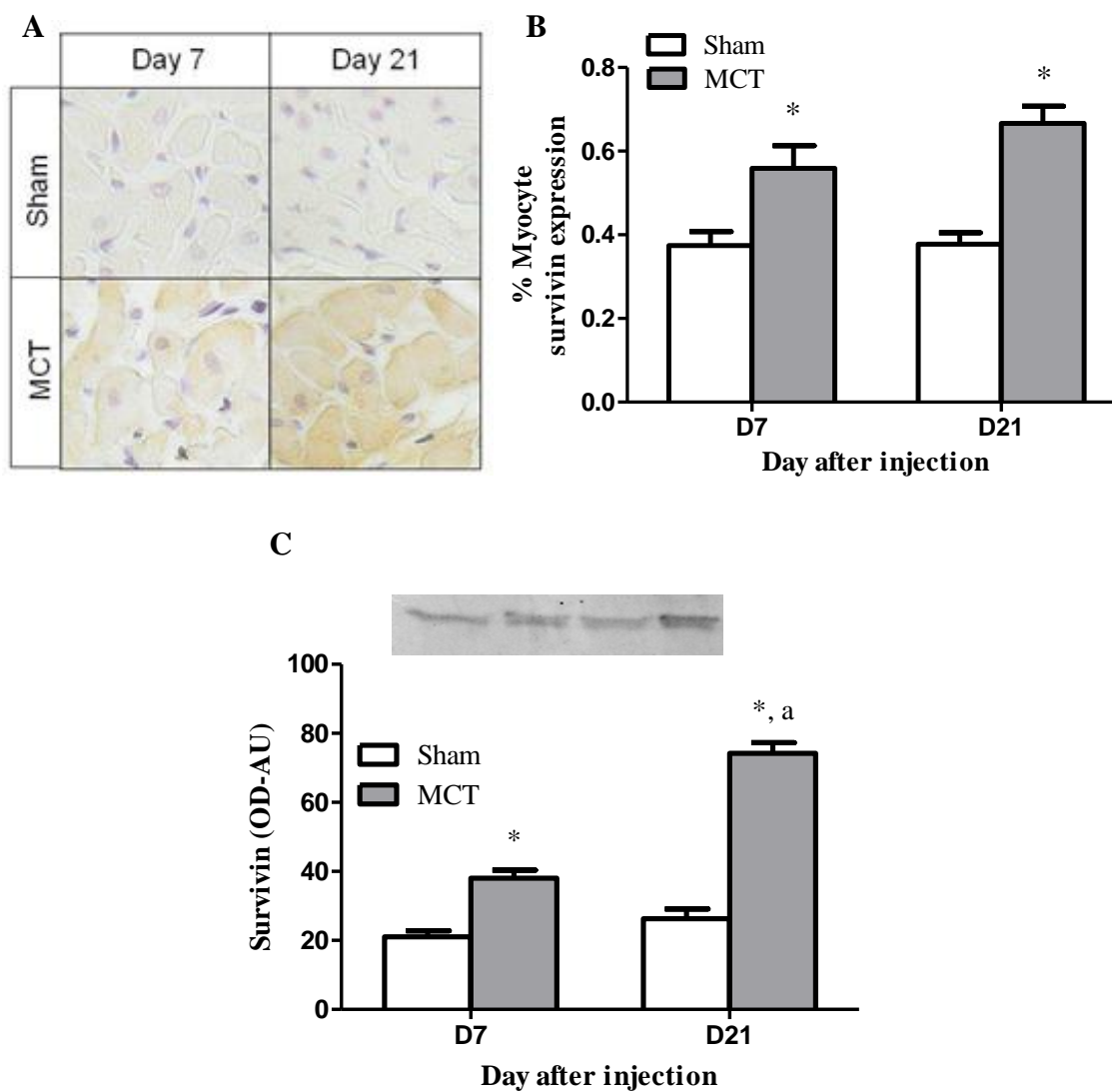
**Figure 5 | Pulmonary arterial and right ventricular cardiomyocyte hypertrophy.** A and C: Histological appearance of small pulmonary arteries and right ventricular cardiomyocytes, respectively, stained with hematoxylin and eosin of Sham and MCT groups 7 and 21 days after injection. B: Medial layer thickness expressed as percentage of all thickness. D: Right ventricular cardiomyocytes hypertrophy expressed as cardiomyocytes cross-sectional area ( $\mu\text{m}^2$ ). Data are present as mean  $\pm$  SEM. Sham, Sham group; MCT, monocrotaline group. \*  $p < 0.05$  vs Sham of the same day; <sup>a</sup>  $p < 0.05$  vs D7 of the same treatment group.

## 2. Survivin and Smac/DIABLO Expression

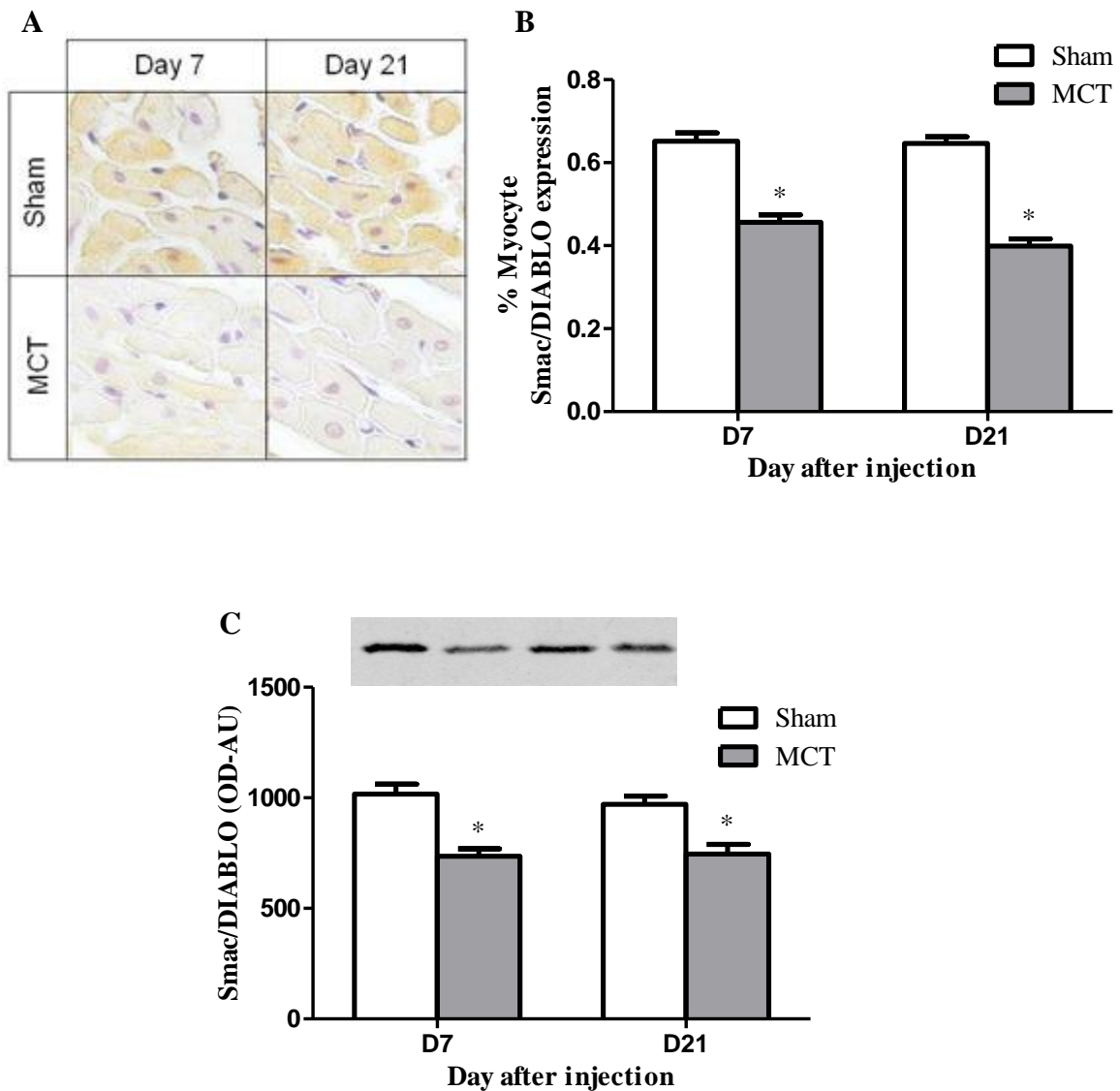
With the aim to study the role of apoptosis in PAH, the expression of survivin and Smac/DIABLO were analysed in RV, LV and lungs.

Survivin and Smac/DIABLO expression in RV were analysed through immunohistochemistry and western blot, and the results are shown in Figure 6 and Figure 7, respectively. MCT treatment significantly raised survivin expression in both day 7 and 21 after injection as compared with Sham group. Sham groups did not present

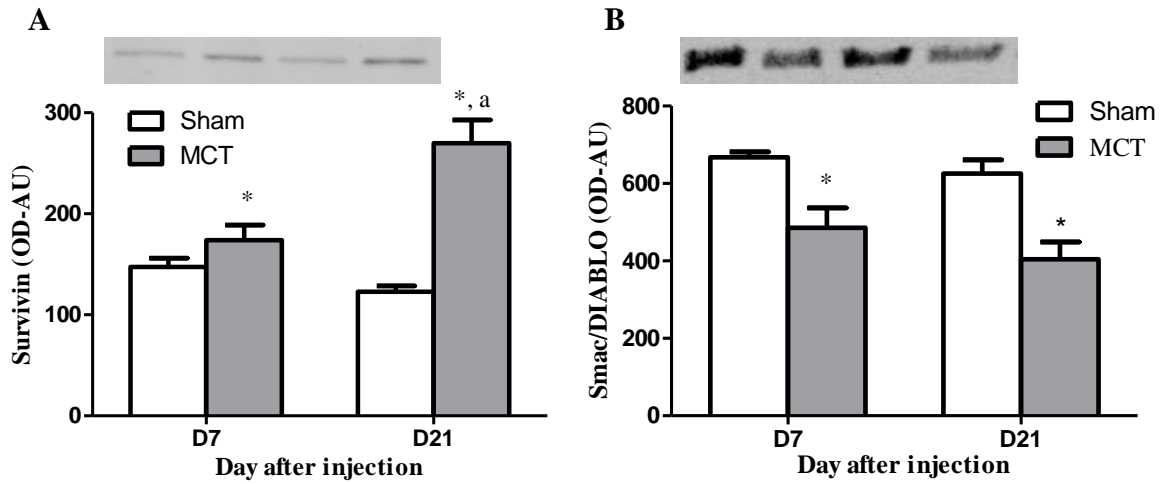
any alterations in survivin expression. Contrarily, MCT treatment significantly decreased Smac/DIABLO expression in comparison with Sham groups. As present in Figure 8, MCT treatment has induced an increase in survivin levels and a decrease in Smac/DIABLO levels in LV, in a similar way of those noted in RV.



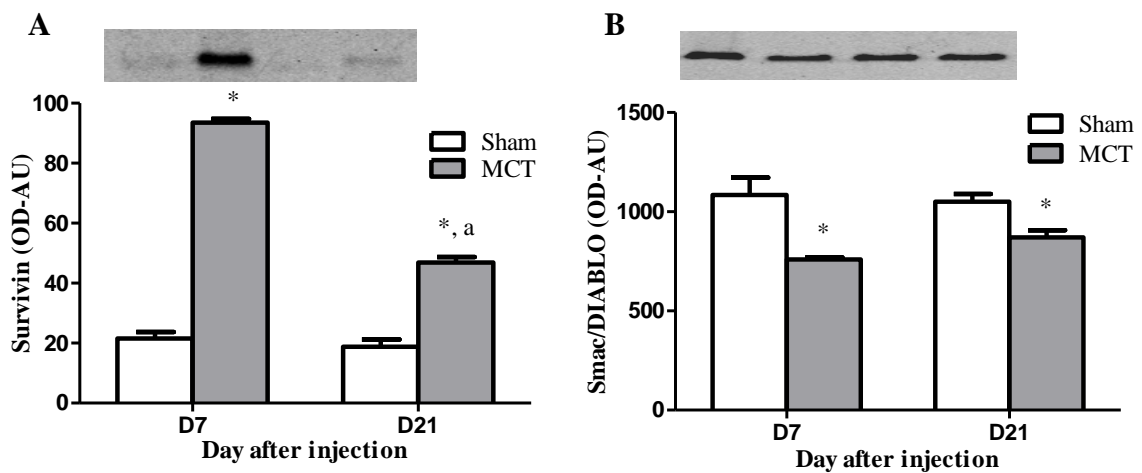
**Figure 6 | Right Ventricle survivin expression**, evaluated through immunohistochemistry and western blot. **A:** Representative image of immunohistochemistry for survivin expression in the right ventricle of Sham and MCT groups 7 and 21 days after injection. **B:** Survivin expression in right ventricle expressed as percentage of stained cardiomyocytes in immunohistochemistry. **C:** Right ventricle survivin expression evaluated through western blot with a representative image of the results. Data are present as mean  $\pm$  SEM. Sham, Sham group; MCT, monocrotaline group; RV, right ventricle. \* $p < 0.05$  vs Sham of the same day; <sup>a</sup> $p < 0.05$  vs D7 of the same treatment group.



**Figure 7 | Smac/DIABLO expression in right ventricle**, evaluated through immunohistochemistry and western blot. **A:** Representative image of immunohistochemistry for Smac/DIABLO expression in the right ventricle of Sham and MCT groups 7 and 21 days after injection. **B:** Smac/DIABLO expression in right ventricle expressed as percentage of stained cardiomyocytes in immunohistochemistry. **C:** Right ventricle Smac/DIABLO expression evaluated through western blot with a representative image of the results. Data are present as mean  $\pm$  SEM. Sham, Sham group; MCT, monocrotaline group; RV, right ventricle. \*  $p < 0.05$  vs Sham of the same day; <sup>a</sup>  $p < 0.05$  vs D7 of the same treatment group.



**Figure 8 | Survivin (A) and Smac/DIABLO (B) expression in left ventricle**, evaluated through western blot of Sham and MCT groups 7 and 21 days after injection, with respective representative images. Data are present as mean  $\pm$  SEM. Sham, Sham group; MCT, monocrotaline group. \*  $p < 0.05$  vs Sham of the same day; <sup>a</sup>  $p < 0.05$  vs D7 of the same treatment group.

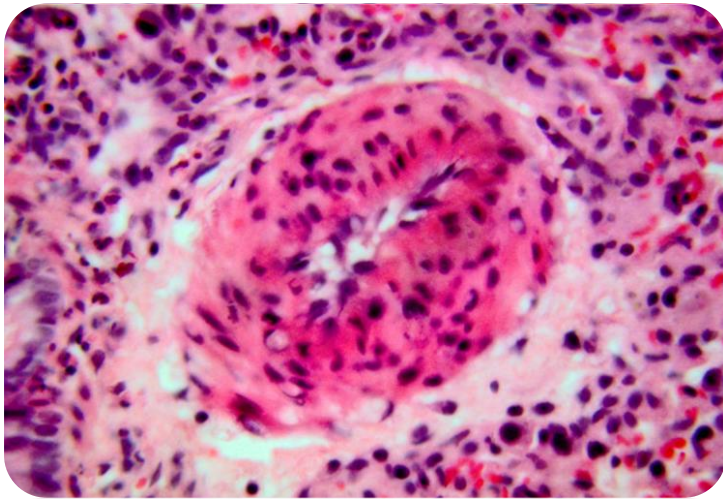


**Figure 9 | Lung Survivin (A) and Smac/DIABLO (B) expression**, evaluated through western blot of Sham and MCT groups, 7 and 21 days after injection, with respective representative images. Data are present as mean  $\pm$  SEM. Sham, Sham group; MCT, monocrotaline group. \*  $p < 0.05$  vs Sham of the same day; <sup>a</sup>  $p < 0.05$  vs D7 of the same treatment group.

In respect to lung (Figure 9), survivin expression was significantly augmented in MCT groups. However, in MCT-treated animals, between day 7 and 21 after injection, there was a significant reduction of survivin expression, while no differences were noted between Sham groups. In contrast, pulmonary Smac/DIABLO expression significantly lowered with MCT treatment in both time-points in comparison with Sham groups.







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## **VI – Discussion**

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PAH is a multifactorial, progressive disease with substantial mortality and morbidity. Over the years, cellular and molecular mechanisms underlying PAH have been studied in order to better understand PAH pathogenesis as well as to develop therapies that might lead to a better prognosis. Thus, studies focusing the development, progression and treatment of PAH are still essential. Most of the research performed with animal models of this disease has been focused on advanced stages of the disease (25-35 days after MCT injection), being scarce the ones conducted in the early phases, probably reflecting the fact that most patients with PAH are usually diagnosed at advance stages. We believe that the study of PAH in early stages will allow a better understanding of pathophysiologic pathways that might be therapeutically modulated aiming to cure PAH. For that reason, the present study analysed the role of apoptosis in two time-points of disease progression in a well characterized animal model of MCT-induced PAH.

MCT administration affects mainly hepatic and cardiopulmonary systems, promoting vasoconstriction, ECs hypertrophy, SMCs hypertrophy and hyperplasia and inflammation at lung level and also RV hypertrophy [15, 182, 193]. Several studies reported that, in rats, the severity of these lesions depends on MCT dose, administration route and the animal age at the time of treatment [180, 245]. Although, not all the typical findings of human PAH are mimicked by this animal model (such as plexiform lesions present in human PAH and absent in the experimental model), it has identical hemodynamic and some morphological features, like pulmonary artery medial layer remodelling and RV hypertrophy, which are the focus of our study [15, 193]. Our results, showed that MCT treatment, after 21 days, induced an augmentation of RV peak systolic pressure (Table 5) in association with increased RV mass (Table 6) and cardiomyocyte hypertrophy (Figure 5), confirming the previous studies. Also, as described before, our results demonstrated that MCT injection promoted pulmonary arterial wall thickness (Figure 5), which is the primary cause of pressure overload, leading to the increase of PVR and PAP and consequently RV hypertrophy. Therefore, RV hypertrophy seems to reflect a compensatory response in which RV adapts to the sustained afterload elevation [21, 246]. Interestingly, although there were no hemodynamic neither morphological alterations (Table 5 and Table 6, respectively) at RV, our results demonstrated for the first time that cardiomyocytes were hypertrophied at day 7 after MCT injection (Figure 5). These results suggest that besides heart

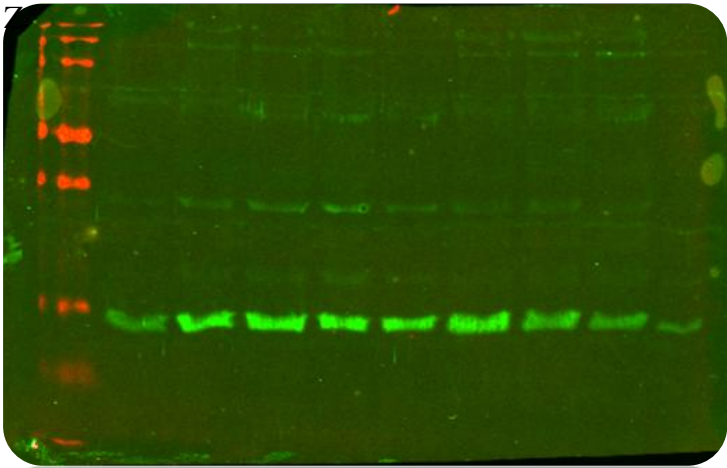
overwork due to pulmonary hypertrophy and PVR, other mechanisms, such as neurohumoral ones, might lead to RV hypertrophy.

In the last years, angiogenesis, dysfunctional  $K_v$  channels, loss expression of prostacyclin synthase gene (anti-proliferative), deregulation of growth signalling pathways and augmentation of survivin expression (anti-apoptotic) have been observed in the development and/or progression of PAH [167-170]. In order to evaluate the contribution of survivin pathway to PAH pathogenesis, we evaluated the cardiac and pulmonary expression of this anti-apoptotic protein as well as of its antagonist, Smac/DIABLO. In 2008, a study carried out by Levkau and colleagues [159] revealed several key features of survivin in the heart. They demonstrated that in the neonatal period, mice deficient for cardiac survivin presented a decrease number of total cardiomyocytes and a marked cardiomyocytes polyploidy due to multiple rounds of DNA replication without cytokinesis. This altered phenotype led to progressive heart failure and ultimately to death. In the same study survivin overexpression was associated with the induction of cell division and the protection of cardiomyocytes from doxorubicin-induced apoptosis. Moreover, they found only residual levels of survivin in normal human heart, but these were extraordinarily increased in the heart of patients with end-stage of heart failure. Interestingly, hemodynamic support with LVAD (left ventricular assist device) resulted in a marked reduction of expression of cardiomyocyte survivin, raising the hypothesis that cardiac survivin expression is load-dependent [159]. Wohlschlaeger and colleagues [247] also described that survivin was overexpressed in congestive heart failure but decreased after unloading. They also demonstrated a positive correlation between cardiomyocytes survivin expression and cardiac hypertrophy and DNA content. In the present study, survivin was significantly enhanced at day 7 after MCT injection and progressively increased throughout the development of MCT-induced PAH (Figure 6). Moreover, cardiomyocyte hypertrophy (Figure 5) was accompanied by increased expression of survivin (Figure 6) both of which progressively increased until day 21. LV expression of survivin had a similar pattern to RV (Figure 8). These results suggest that survivin might be involved in the cardiac remodelling process underlying PAH. Previously, McMurtry and colleagues [121] also demonstrated that lung survivin overexpression preceded hemodynamic alterations verified in the disease. Our results (Table 5 and Figure 9) corroborate McMurtry findings. Therefore, taking into consideration: i) the overexpression of RV,

LV and lung survivin at day 7 after MCT injection, ii) the similar survivin expression profile between LV and RV, and iii) the occurrence of survivin overexpression before hemodynamic alterations, makes us believe that besides pulmonary arterial hypertrophy other mechanisms, perhaps neurohumoral ones, are involved in RV hypertrophy. In particular, we hypothesised that pulmonary vascular remodelling phenomena may be signaled to RV, possibly through neurohumoral mediators, such as ET-1, ATII, PDGF and catecholamines, inducing survivin overexpression and cardiomyocytes hypertrophy, a favourable cardiac remodelling response to the disease, as previously suggested [21, 248].

The expression of Smac/DIABLO was the reverse of survivin expression with decreasing levels noticed from day 7 to day 21 after MCT administration, in both RV and LV as well as in lung (Figure 6, Figure 8 and Figure 9, respectively). Data suggest that expression of Smac/DIABLO decreases with the progression of PAH. Smac/DIABLO is known for its pro-apoptotic properties through its interaction with IAP proteins, including survivin, eliminating their inhibitory effect on caspases [119, 151, 249]. On the other hand, survivin expression is also able to delay the release of smac/DIABLO from the mitochondria, preventing its pro-apoptotic function [152]. Several studies also revealed a reciprocal expression of survivin and Smac/DIABLO in different types of cancers [250-253]. So, we believe that deregulation in the balance between survivin and Smac/DIABLO might be related with pulmonary vascular remodelling and cardiomyocytes hypertrophy in response to apoptotic stimuli.





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## VII – Conclusions

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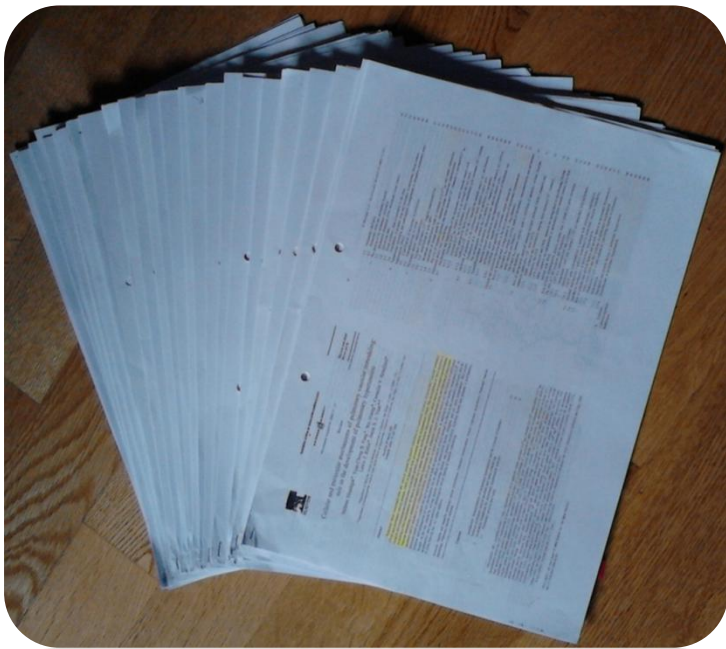
Aiming to evaluate the contribution of apoptosis to the pathogenesis of PAH, MCT animal model was used to analyse pulmonary and cardiac expression of survivin and Smac/DIABLO. Data allowed us to conclude that:

- i) cardiomyocytes hypertrophy was present 7 days after MCT injection, proceeding hemodynamic alterations;
- ii) RV and LV survivin overexpression was demonstrated 7 days after MCT injection and progressively increased throughout the development of MCT-induced PAH;
- iii) although increased 7 days after MCT injection, lung survivin expression progressively decreased between time-points;
- iv) RV, LV and lung Smac/DIABLO expression was upregulated 7 days after MCT injection but progressively decreased throughout the development of MCT-induced PAH.

Taken together, data suggest that a deregulation in the balance between survivin and Smac/DIABLO might be related with pulmonary vascular remodelling and cardiomyocytes hypertrophy in response to apoptotic stimuli. Therefore, we consider that a therapy targeting survivin and/or Smac/DIABLO might be a future option for the treatment of PAH.

In the future, we intend to study survivin expression in cellular fractions (nuclear, cytoplasmic and mitochondrial) as well as the existence of possible posttranslational modifications that regulate cellular location of this protein and, consequently, its activity. Data obtained in the present study also leads us to therapeutically target the balance between survivin and Smac/DIABLO using terameprocol which is a drug widely used in cancers therapeutics.





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## VIII – Bibliography

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