

ORIGINAL RESEARCH

# Human Resistin Induces Cardiac Dysfunction in Pulmonary Hypertension

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**BACKGROUND:** Cardiac failure is the primary cause of death in most patients with pulmonary arterial hypertension (PH). As pleiotropic cytokines, human resistin (Hresistin) and its rodent homolog, resistin-like molecule  $\alpha$ , are mechanistically critical to pulmonary vascular remodeling in PH. However, it is still unclear whether activation of these resistin-like molecules can directly cause PH-associated cardiac dysfunction and remodeling.

**METHODS AND RESULTS:** In this study, we detected Hresistin protein in right ventricular (RV) tissue of patients with PH and elevated resistin-like molecule expression in RV tissues of rodents with RV hypertrophy and failure. In a humanized mouse model, cardiac-specific Hresistin overexpression was sufficient to cause cardiac dysfunction and remodeling. Dilated hearts exhibited reduced force development and decreased intracellular  $Ca^{2+}$  transients. In the RV tissues overexpressing Hresistin, the impaired contractility was associated with the suppression of protein kinase A and AMP-activated protein kinase. Mechanistically, Hresistin activation triggered the inflammation mediated by signaling of the key damage-associated molecular pattern molecule high-mobility group box 1, and subsequently induced pro-proliferative Ki67 in RV tissues of the transgenic mice. Intriguingly, an anti-Hresistin human antibody that we generated protected the myocardium from hypertrophy and failure in the rodent PH models.

**CONCLUSIONS:** Our data indicate that Hresistin is expressed in heart tissues and plays a role in the development of RV dysfunction and maladaptive remodeling through its immunoregulatory activities. Targeting this signaling to modulate cardiac inflammation may offer a promising strategy to treat PH-associated RV hypertrophy and failure in humans.

**Key Words:** cardiac remodeling ■ HMGB1 ■ inflammation ■ RELM $\alpha$  ■ right ventricle

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Cardiac failure is the primary cause of death in most patients with pulmonary arterial hypertension (PH). As pleiotropic cytokines, human resistin (Hresistin) and its rodent homolog, resistin-like molecule (RELM)  $\alpha$ , are mechanistically critical to pulmonary vascular remodeling in PH.<sup>1</sup> However, it is still unclear whether activation of these RELM molecules can directly cause PH-associated cardiac dysfunction and remodeling.

Development of right ventricular (RV) dysfunction (RVD) or RV failure in patients with PH determines their clinical outcome. However, therapies targeting

pulmonary vasculature and flow have little success in reducing morbidity and mortality in these patients.<sup>2</sup> Therefore, effective treatment strategies directly targeting RVD are needed. One critical barrier to developing an effective therapy that targets the right ventricle is the poor understanding of the associated pathogenetic mechanisms.

Resistin-like molecules (RELMs) are pleiotropic cytokines with pro-proliferative, pro-inflammatory, pro-fibrotic, vasoconstricting, and chemokine activities. Previous studies point toward their role not only in

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## CLINICAL PERSPECTIVE

### What Is New?

- Human resistin (Hresistin) protein is upregulated in right ventricular (RV) tissues of patients with PH, and cardiac-specific Hresistin overexpression in humanized mice causes heart dysfunction and remodeling.
- Hresistin activates damage-associated molecular pattern signaling to trigger cardiac inflammation in RV.
- Anti-Hresistin human antibody ameliorates cardiac hypertrophy and failure in rodent PH models.

### What Are the Clinical Implications?

- By using RV tissues from patients with PH, a humanized mouse model of cardiac-specific Hresistin overexpression, and a human antibody targeting Hresistin, we found that Hresistin drives development of RV dysfunction and maladaptive remodeling through its inflammation-promoting activities.
- Our data suggest that Hresistin has a unique role in the immune interaction between lung and RV in humans, mediating remodeling in both organs. Thus, targeting Hresistin signaling may offer a promising strategy to treat PH-associated RV hypertrophy and failure clinically as well as pulmonary vascular remodeling.

## Nonstandard Abbreviations and Acronyms

<b>AMPK</b>	AMP-activated protein kinase
<b>DAMP</b>	damage-associated molecular pattern
<b>EP</b>	ethyl pyruvate
<b>HMGB1</b>	high mobility group box 1
<b>Hresistin</b>	human resistin
<b>MHC</b>	myosin heavy chain
<b>NRCM</b>	neonatal rat cardiomyocyte
<b>PH</b>	pulmonary arterial hypertension
<b>PKA</b>	protein kinase A
<b>RELM</b>	resistin-like molecule
<b>RVD</b>	right ventricular dysfunction
<b>SSc</b>	scleroderma
<b>tTA</b>	tetracycline trans-activator

pulmonary vascular remodeling,<sup>3–5</sup> but also in cardiac left ventricle (LV) dysfunction during PH. Systematic injection of experimental rats with an adeno-associated virus with rat resistin gene reportedly induced myocardial dysfunction, especially in the LV.<sup>6</sup>

However, the homolog of rodent resistin could not be identified in humans.<sup>1</sup> Unlike rodent resistin, rodent RELM $\alpha$  and Hresistin are primarily derived from immune cells and have similar inflammation-modulatory functions.<sup>1</sup> Thus, RELM $\alpha$  was recognized as the rodent counterpart of Hresistin.<sup>1</sup> We have shown that rodent RELM $\alpha$  (also known as found in inflammatory zone 1<sup>7</sup> or hypoxia-induced mitogenic factor)<sup>8</sup> and Hresistin have a causal role in PH pathogenesis, because these proteins are mechanistically critical to pulmonary vascular inflammation and remodeling.<sup>3,5,7–17</sup> However, it remained unclear whether RELM $\alpha$  or Hresistin is present in cardiac tissue during PH development and whether activation can directly induce RVD and remodeling.

In the present study, we detected upregulated Hresistin protein in the RV tissue of patients with PH. We also found that RELM $\alpha$  expression was induced in the RV tissue of rodents with cardiac hypertrophy and failure during PH development. In this case, RELM $\alpha$  was present in cardiomyocytes and infiltrating immune cells. To further dissect the cardiac involvement of RELM signaling and more accurately mimic the human disease, we developed the humanized, cardiac-specific,  $\alpha$ -myosin heavy chain (MHC)-directed Hresistin knock-in mouse model. We found that Hresistin overexpression in the heart is sufficient to cause cardiac dysfunction and remodeling in mice. These humanized mice had dilated, impaired RVs with decreased intracellular Ca<sup>2+</sup> availability. Ca<sup>2+</sup> was associated with the Hresistin regulation of protein kinase A (PKA) and AMP-activated protein kinase (AMPK) in the hearts. Mechanistically, Hresistin overexpression triggered activation of the key damage-associated molecular pattern (DAMP) molecule known as high mobility group box (HMGB)1. The presence of HMGB1 was associated with increased Ki67 proliferation in leukocytes that infiltrated damaged right hearts. Moreover, an antibody that we developed to Hresistin prevented RV hypertrophy in PH and prolonged the survival of experimental rats with failing hearts. Our data indicate that Hresistin plays a direct cardiac role in the pathogenesis of RVD. This knowledge can benefit the search for a new therapeutic approach to PH-accompanied RVD and other related heart diseases in humans.

## METHODS

We present a sequence of studies aimed at identifying the role of Hresistin in the pathogenesis of RVD. First we examined stored human cardiac samples of patients with scleroderma (SSc)-associated PH (SSc-PH) and of control patients with non-SSc PH. Next we created and examined induced PH rodent models, both rat and mouse, and compared them with controls. We created a transgenic mouse line to target the mechanistic responses in cardiac tissue. Finally, we

administered anti-Hresistin monoclonal antibodies to rats before induction of PH.

The Johns Hopkins Medicine Institutional Review Board issued type 4 exemption approval before the start of these studies. Animal housing and experimental protocols were approved by the Animal Care and Use Committee of Johns Hopkins University.

The data that support the findings of this study are available from the corresponding author upon reasonable request.

## Human Cardiac Tissue

We obtained RV tissue from patients  $\geq 18$  years old with SSc-associated PH from stored samples through collaboration with the Johns Hopkins Department of Pathology. For comparison, we also obtained heart tissue from control patients with no signs of PH. These heart samples were collected through biopsy. Clinical data pertaining to these patients with SSc-PH, including right heart catheterization parameters, were published in our previous study.<sup>18</sup>

## Generation of a Humanized Transgenic Mouse Model That Overexpresses Hresistin in Cardiomyocytes

We generated a knock-in transgenic MHC-hRETN-overexpressing mouse line that overexpresses cardiac-specific Hresistin. To spatially and temporally control the expression of Hresistin in mouse heart tissue, we generated a dual transgenic inducible overexpression system. Briefly, the model has 2 transgenes. The first contains a promoter-specific sequence encoding a tetracycline trans-activator (tTA). The tTA is a fusion protein made up of a mutated tet Repressor protein and the herpesvirus VP-16 trans-activator,<sup>19</sup> a powerful activator of gene transcription. The tTA protein binds to a tetracycline operator [(*otet*)7CMV] site, also known as the tetracycline response element, upstream of the Hresistin gene sequence (RETN). The cardiomyocyte-specific MHC promoter was used to drive the expression of the tTA as previously described.<sup>20</sup> The second transgene in these mice contains the RETN gene construct. The construct is composed of (1) a tetracycline response element, (2) a “tight” control sequence to reduce expression of RETN in the absence of tTA, and (3) the RETN cDNA followed by a FLAG tag. The tetracycline response element-Tight-RETN transgene was microinjected into fertilized eggs as described previously.<sup>21</sup> RETN transgene-positive founders were identified and cross-bred with the MHC mice to obtain the double-transgenic mice having cardiac-specific Hresistin overexpression. Trabecular muscles from the RV tissues of MHC-hRETN-overexpressing mice were dissected, and twitch force and intracellular  $\text{Ca}^{2+}$

concentration ( $[\text{Ca}^{2+}]_i$ ) were measured as described in our previous studies.<sup>22,23</sup>

## Anti-Hresistin Antibody Development and Administration

We used our previously reported techniques to generate anti-Hresistin monoclonal antibodies.<sup>24</sup> They were identified through phage screening of a human library, and screened by ELISA and through plasmon resonance. Potential antibodies were validated for their in vitro antiproliferative function against Hresistin in human primary pulmonary smooth muscle cells. They were further screened for immunogenicity, manufacturability, and toxicity.<sup>24</sup> The human antibodies exhibited cross reactivity to block rodent RELM $\alpha$  in smooth muscle cell assays.<sup>24</sup> For therapeutic antibody administration in this study, rats received twice weekly intraperitoneal injections of 4 mg/kg before we began induction of PH by hypoxia or monocrotaline.

## Animal Models of PH

Our PH rodent models were induced by monocrotaline<sup>22,25</sup> or hypoxia<sup>10</sup> as we have described previously. Animals were maintained at 20 to 24 °C with a 12:12-hour light–dark cycle with access to normal laboratory diet (Teklad global 18% protein rodent diet; Envigo) and chlorinated water ad libitum. Cage bedding was also from Envigo (7097 Teklad corncob bedding).<sup>16</sup> For induction of the hypoxia-induced PH model, 8- to 12-week-old male C57BL/6 mice or 250 to 300 g male Sprague–Dawley rats (Charles River Laboratories, Wilmington, MA) were exposed to 10.0%  $\text{O}_2$  (hypoxia) for 4 or 28 days and then euthanized and processed as described previously.<sup>10</sup> Control mice were exposed to normal room air (20.8%  $\text{O}_2$ , normoxia). As we have explained previously,<sup>16</sup> given the reports that estrogens exhibit protective effects in classical rodent PH models,<sup>26,27</sup> and the finding that the damage-associated inflammatory response is more profound in male rodents with PH than in corresponding female rodents,<sup>28</sup> we used only male rodents to produce a more severe disease and investigate the Hresistin-regulated DAMP signaling.<sup>16</sup> Each animal was identified using a numbered ear tag. Investigators performing procedures on the animals were blinded to the group assignment. Animals in each group were unblinded once data analysis was completed. The group sizes were determined by power calculations. We ensured that experiments were unbiased by following the recent guidelines for PH preclinical and translational research.<sup>29,30</sup> For the monocrotaline-induced RVD model, 250 to 300 g male Sprague–Dawley rats from Charles River were randomized to subcutaneous injection with monocrotaline 60 mg/kg (Sigma Aldrich) or vehicle control. For

the echocardiographic analysis, before anesthesia for hemodynamic and other analyses, each rat underwent transthoracic echocardiography in an awake state on the indicated time points. The hearts were harvested at the following time points: (1) control, no intervention; (2) RV hypertrophy, defined as compensated hypertrophic RV (significantly increased RV wall thickness and RV size, as compared with the nonmonocrotaline control) with maintained contractility (as indicated by the RV ejection fraction) (2.5–4.0 weeks postmonocrotaline injection)<sup>22</sup>; (3) RV failure, defined as decompensated (hypertrophic) RV with significantly decreased contractility (4.5 weeks postmonocrotaline injection); in this decompensated stage, the rats clinically exhibited signs of RV failure including low food intake, retarded weight gain, and lack of daily activity.<sup>25</sup> Between 5 and 6 weeks after monocrotaline treatment, the key hemodynamic index cardiac output, calculated by echocardiography, was >70% decreased.<sup>22,25</sup> This is comparable to the cardiac output values calculated by catheterization (~50% decrease) in the same decompensated stage and the same monocrotaline rat model by other groups.<sup>31–33</sup> The rats entering the decompensating phase exhibited increased morbidity and mortality. Their survival curves were recorded.

### RV Systolic Pressure and Fulton Index

For measurement of RV systolic pressure, the hypoxia-treated (or control normoxia) rats were weighed, and sustained anesthesia was induced by intraperitoneal injections of ketamine (100 mg/kg) and xylazine (10 mg/kg). A trachea cannula was inserted, and the animal was ventilated with a Harvard Rodent Ventilator Model 683 (1.2–2.5 mL/breath and 70–81 breaths/min); rocuronium was injected intraperitoneally (2 mg/kg) as a paralytic. Rat body temperature was monitored and controlled via rectal probe connected to a TCAT 2LV Controller (Physitemp, Clifton, NJ) to provide continuous feedback. The chest was opened to expose the heart. RV systolic pressure was measured via apical stab followed by insertion of a 1.4F pressure-volume catheter (SPR-839, Millar Instruments, Houston, TX) into the right ventricle. Data were collected with the AD Instruments Powerlab 8/35 (AD Instruments, Colorado Springs, CO) and Millar MPVS Ultra. Rats were allowed to reach a stable baseline (~10–15 minutes) before we recorded 3 nonventilated pressures to determine mean RV systolic pressure per animal. After hemodynamic measurements, rats were euthanized by exsanguination, and the heart was removed en bloc and bisected into the RV and LV plus septum. Each portion of the heart was weighed, and the RV and LV plus septum ratio was determined as the Fulton index. For the echocardiographic analysis of these rats exposed to hypoxia or control normoxia, before anesthesia for

hemodynamic and other analyses, each rat underwent transthoracic echocardiography in an awake state. Here we report M-mode evaluation of RV thickness and pulmonary artery Doppler flow analysis of pulmonary acceleration time/pulmonary ejection time ratio.

### Immunoprecipitation

We assessed binding of the human therapeutic antibody against Hresistin to rodent RELM $\alpha$  by immunoprecipitation. We incubated 2  $\mu$ g of the anti-Hresistin antibody with 100 ng of laboratory-made FLAG-tagged recombinant rat RELM $\alpha$  protein and Dynabeads Protein A (10001D, Thermo Fisher, Waltham, MA). Protein–antibody binding was detected by western blotting with anti-Hresistin antibody (AF1359, R&D Systems, Minneapolis, MN). Recombinant rat RELM $\alpha$  protein (30 ng) served as a positive loading control.

### Trabecular Muscles

Twitch force and intracellular Ca<sup>2+</sup> concentration ([Ca<sup>2+</sup>]<sub>i</sub>) were measured as described in our previous studies.<sup>20,24</sup> Trabecular muscles from the right ventricle were dissected, mounted between a force transducer and a motor arm, superfused with Krebs–Henseleit solution at a rate of ~10 mL/min, and stimulated at 0.5 Hz. The dissecting Krebs–Henseleit solution was composed of (in mmol/L) NaCl 120, NaHCO<sub>3</sub> 20, KCl 5, MgCl 1.2, glucose 10, CaCl<sub>2</sub> 0.5, and 2,3-butanedione monoxime 20 (pH 7.35–7.45 at room temperature [21–22 °C]). Force was measured by a force transducer and [Ca<sup>2+</sup>]<sub>i</sub> was measured by using fura-2. The muscles underwent isometric contractions with the resting muscle length set such that resting force was 15% of total force development (optimal muscle length).

### In Vitro Culture, Treatments, and Assessment of the Cardiomyocytes Isolated From Neonatal Rats

Neonatal rat cardiomyocytes (NRCMs) were isolated from 1- to 3-day-old Sprague–Dawley pups and cultured in DMEM as we previously described.<sup>34</sup> NRCMs were serum starved in DMEM containing 0.1% insulin transferrin selenium (Thermo Fisher) for 24 hours and then incubated with the anti-Hresistin antibody at 3  $\mu$ g/mL (a dose that has blocking activities against Hresistin recombinant protein without cytotoxic effects<sup>24</sup>) or with 5  $\mu$ mol/L HMGB1 inhibitor ethyl pyruvate (EP) (#E47808, Sigma). After stimulation, NRCMs were transfected with MOI-300 green fluorescent protein–tagged adeno-associated virus expressing the Hresistin (*hRETN*) (the viral vectors containing the ubiquitous CB promoter were produced by the University of Florida Vector Core Laboratory, as we previously

described<sup>3,9</sup>). After 24 hours, some treated/transfected NRCMs were harvested for western blotting analysis. For the cell surface area measurement, after treatment and transfection of the cultured NRCMs, real-time live cell images were captured using the IncuCyte S3 Live-Cell Analysis System and analyzed using the IncuCyte software to calculate cardiomyocyte area, as we previously described.<sup>34</sup>

### Quantitative Real-Time Polymerase Chain Reaction

Total RNA was isolated with the RNeasy kit (Qiagen) according to the manufacturer's protocol. Total RNA (500 ng) was reverse transcribed into cDNA. Quantitative polymerase chain reaction was carried out on an ABI 7500 fast real-time polymerase chain reaction system (Applied Biosystems, Foster City, CA). Fold changes in gene expression were acquired by using the delta method and normalization to 18S rRNA. The primers used in this process were 5'-CAATCCCATGGCGTATA AAAGCATC-3' (RELM $\alpha$  forward), 5'-TCATTCTTAGG ACAGTTGGCAGCAG-3' (RELM $\alpha$  reverse), 5'-TGAGC AAGAGAGGCCCTATC-3' (GAPDH forward), and 5'-AGGCCCTCCTGTTATTATG-3' (GAPDH reverse).

### Immunohistology

We used immunofluorescence staining to detect the expression of FLAG-tagged Hresistin in the RV tissue of knock-in mice. After deparaffinization of tissue, rehydration, and antigen retrieval, sections were treated with anti-heavy chain cardiac myosin (ab185967, Abcam) and anti-FLAG (F1804, Sigma) antibodies overnight at 4 °C and then with Alexa Fluor488-donkey anti-mouse IgG (715–545-150, Jackson ImmunoResearch) and Cy3-donkey anti-rabbit IgG (711–166-152, Jackson ImmunoResearch) for double fluorescence staining. In the monocrotaline model, rat RV sections were treated with anti-RELM $\alpha$  (AF1523, R&D Systems) and anti-myosin (ab185967, Abcam) antibodies, whereas human RV tissue from SSc-PH patients were treated with anti-Hresistin antibody (AF1359, R&D Systems) for immune staining. In other immunofluorescent studies, RV tissue from monocrotaline-treated wild-type rats, hypoxic wild-type mice, and MHC-hRETN mice were sliced and stained with anti-Ki67 (ab16667, Abcam) or anti-HMGB1 (ab18256, Abcam) antibodies, with or without co-staining of anti-heavy chain cardiac myosin (ab185967, Abcam), F4/80 (ab6640, Abcam), myeloperoxidase (MPO; AF3667, R&D), or vimentin (ab8978, Abcam). Sections were then incubated with the appropriate fluorochrome-coupled secondary antibody (Jackson ImmunoResearch). Finally, all sections were mounted in ProLong Gold anti-fade reagent with DAPI (Invitrogen). Staining was imaged and tissue sections

were analyzed by confocal microscopy (Leica SPE DMI8).

### Western Blotting

The collected mouse/rat heart tissues or NRCMs in RIPA buffer (Sigma) (supplemented with 1 mmol/L phenylmethylsulfonyl fluoride, 1 mmol/L Na<sub>4</sub>VO<sub>3</sub>, and protease inhibitor mixture [Roche, 116974980011]) were lysed with homogenization beads (0.9–2.0 mm; SSB14B, Bullet Blender) in a Bullet Blender at 4 °C, vortexed, and then centrifuged. The concentration of isolated proteins was measured by the bicinchoninic acid method (BCA kit, Bio-Rad). The supernatants were mixed in SDS sample loading buffer (NuPAGE, Invitrogen) at 99 °C for 10 minutes and then subjected to SDS-PAGE. After electrophoresis, proteins were transferred onto polyvinylidene difluoride membranes and immunoblotted with antibodies to RELM $\alpha$  (AF1523, R&D Systems), total PKA C (4782, Cell Signaling), phospho-PKA C (5661, Cell Signaling), total AMPK (Ab131512, Abcam), phospho-AMPK (50081, Cell Signaling), HMGB1 (ab18256, Abcam), Ki67 (ab16667, Abcam), or GAPDH (G8795, Sigma) overnight at 4 °C, and then probed with horseradish peroxidase-conjugated secondary antibodies (Bio-Rad) for 2 hours. Protein bands were visualized by chemiluminescence (ECL; Amersham Pharmacia Biotech, Arlington Heights, IL).

### Statistical Analysis

Data are presented as the mean $\pm$ SEM. Comparisons between 2 groups were analyzed by Student *t* test, and comparisons of multiple groups were analyzed by 1-way ANOVA followed by the Newman-Keuls post hoc test unless otherwise noted: the Tukey post hoc test was used for the immunoblotting analysis in the RV of antibody-treated rats, and the log-rank test was used for the survival analysis. All statistical analyses were performed with Prism 7.0e (GraphPad Software, La Jolla, CA). A *P* < 0.05 was considered statistically significant.

## RESULTS

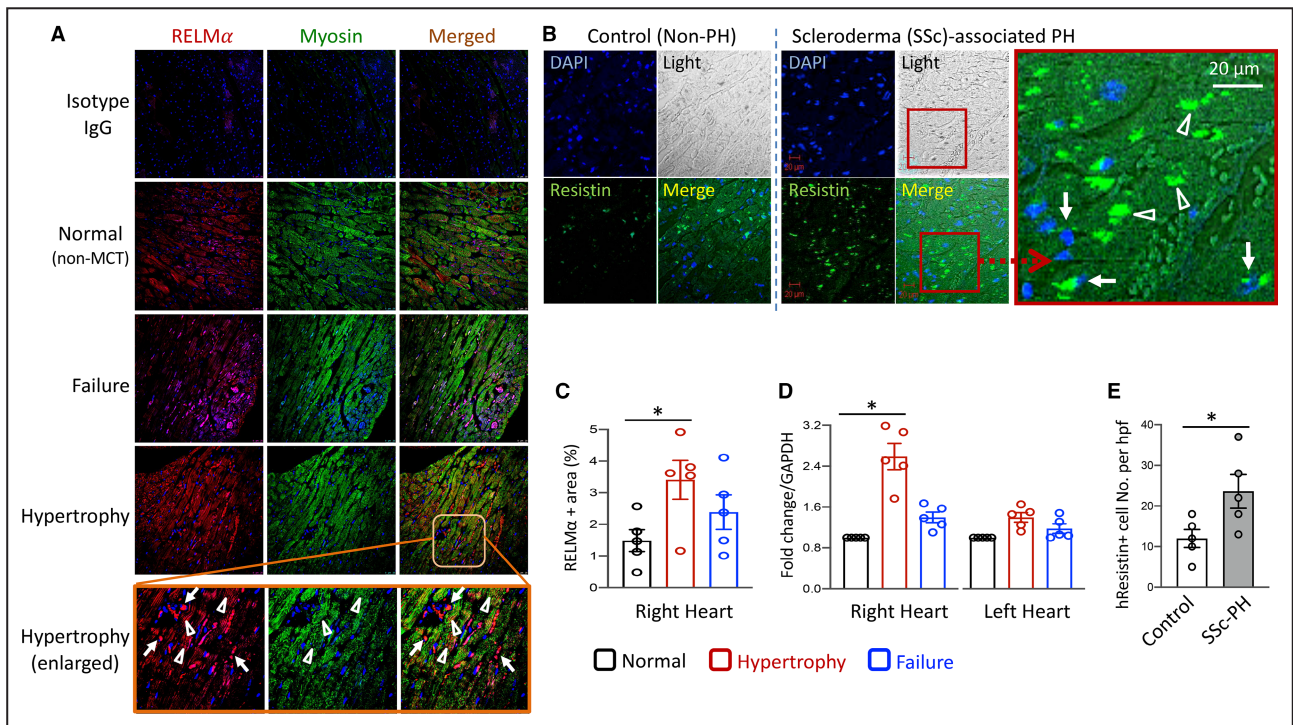
We found evidence to suggest that RELM $\alpha$ /Hresistin signaling contributes to RVD in parallel with PH development in rodents as well as humans. In addition, Hresistin knock-in mice displayed RVD, as seen in RV remodeling, impaired contractility, and induced myocyte hypertrophy and fibrosis. We also examined evidence of HMGB1 signaling in monocrotaline-induced rats and Hresistin knock-in mice, identifying a pathological temporal progression in rats. In addition, administration of the antagonist EP inhibited HMGB1 release

in rats. Finally, we administered an anti-Hresistin human antibody in the chronic hypoxia-induced PH rat model with promising results for therapeutic application.

### Hresistin/RELM $\alpha$ Expression in RV Tissues of PH Rodent Models and Patients

Before examining the expression of Hresistin/RELM $\alpha$ , the monocrotaline-treated rats were tested by echocardiography (Figure S1A) and categorized as normal (RV wall thickness:  $0.49 \pm 0.01$  mm; RV chamber diameter:  $2.46 \pm 0.08$  mm; ejection fraction:  $70.6 \pm 6.4\%$ ),

RV hypertrophy (RV wall thickness:  $0.59 \pm 0.02$  mm; RV chamber diameter:  $2.82 \pm 0.20$  mm; ejection fraction:  $75.2 \pm 2.1\%$ ), and RV failure (RV wall thickness:  $0.60 \pm 0.02$  mm; RV chamber diameter:  $3.22 \pm 0.10$  mm; ejection fraction:  $61.9 \pm 3.5\%$ ). These echocardiography data were consistent with that of our previous studies on the monocrotaline rat model.<sup>22,25</sup> We also characterized the experimental rats at the histologic level with Masson's trichrome analysis (Figure S1B), which showed that monocrotaline drove progressive and severe RV fibrotic remodeling. Following the study on the RV status, we found that the rodent homolog of Hresistin was upregulated in the RV tissues of the



**Figure 1. Expression of Hresistin/RELM $\alpha$  during right ventricular dysfunction.**

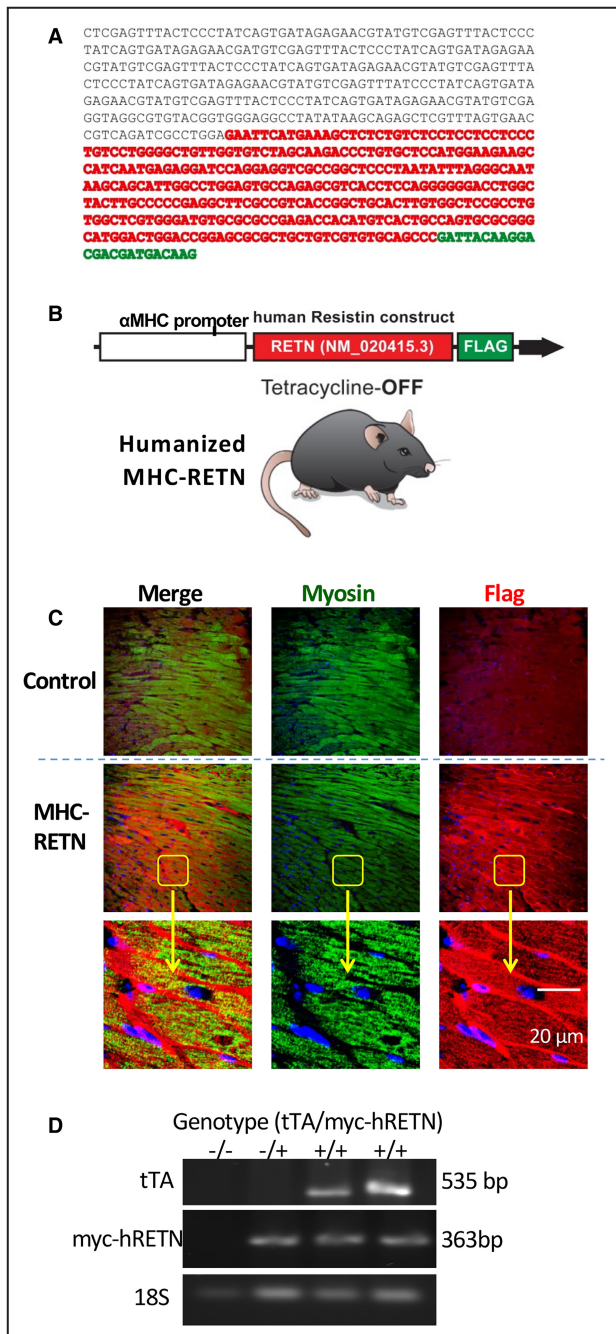
**A**, Immunofluorescence images of RV tissue slices from monocrotaline (MCT)-treated rats and normal controls. Sections were stained with anti-RELM $\alpha$  (red), co-stained with anti-myosin (green), and counterstained with DAPI. Original magnification:  $\times 200$ . The bottom panels show the outlined area from the 2-week post-MCT time point at higher magnification ( $\times 400$ ). The arrowheads point to the myosin (green)-labeled cardiomyocytes positively stained for RELM $\alpha$  (red), whereas the arrows point to the myosin-negative RELM $\alpha$ -positive immune cells that infiltrated into the myocardial interstitium. Representative images are from 5 individual RV samples per group. **B**, Hresistin detection in human RV tissue ( $n=5$  subjects per group). Confocal images show Hresistin (green signal) staining in the RV heart biopsy of non-PH control subjects and patients with scleroderma (SSc)-associated pulmonary arterial hypertension. Light micrograph of fluorescence images to show structure. Signals of light micrograph (showing tissue structure) and fluorescence images (of Hresistin and DAPI staining) are digitally merged and the boxed area is enlarged and displayed in the right panel. The arrowheads point to the Hresistin protein signals in cardiomyocytes whereas the arrows point to the Hresistin-positive staining in the infiltrating immune cell. **C**, Quantitative analysis of data in **(A)**. Percentage of areas positive for RELM $\alpha$  in rat RV tissues was determined by the histogram tool with Adobe Photoshop software. Data are presented as mean $\pm$ SEM ( $n=5$  animals per group). \* $P<0.05$  vs normal group. **D**, Quantitative real-time PCR analysis of RELM $\alpha$  genes in the right heart and left heart tissues of MCT-injected rats ( $n=5$  animals per group; the dots in the graphs represent single individuals) with cardiac hypertrophy (2 weeks post-MCT) and failure (4 weeks post-MCT). **E**, Quantitative analysis of data in **(B)**. The Hresistin-positive (+) cells were counted and expressed as numbers per high-power field (hpf). Data are presented as mean $\pm$ SEM ( $n=5$  subjects per group). \* $P<0.05$ . DAPI indicates 4'-6-diamidino-2-phenyl-indole; Hresistin, human resistin; PCR, polymerase chain reaction; PH, pulmonary arterial hypertension; RELM $\alpha$ , resistin-like molecule- $\alpha$ ; and RV, right ventricular.

monocrotaline-induced PH model. Protein (Figure 1A and 1C) and gene (Figure 1D) expression of RELM $\alpha$  in rats was especially clear in the hypertrophic remodeling RV, as compared with that in the RV of control rats or in the unaffected LV of the same heart. Furthermore, costaining revealed that endogenous RELM $\alpha$  was localized to both infiltrating immune cells and cardiomyocytes of rats with RVD and right ventricular failure (Figure 1A), indicating cellular sources of this cytokine. Of note, in cardiac biopsy samples from patients with SSc-PH, immunofluorescence analysis also detected the upregulated Hresistin protein signal in RV tissues

of patients with PH compared with those of non-PH control subjects (Figure 1B and 1E). The expression of Hresistin was observed in the immune cells and cardiomyocytes of patients with PH (Figure 1B), consistent with that of RELM $\alpha$  in PH rats (Figure 1A). These data indicate that RELM $\alpha$ /Hresistin signaling in the RV contributes to RVD development in parallel with pulmonary vascular remodeling in lungs during PH development.<sup>16,17</sup>

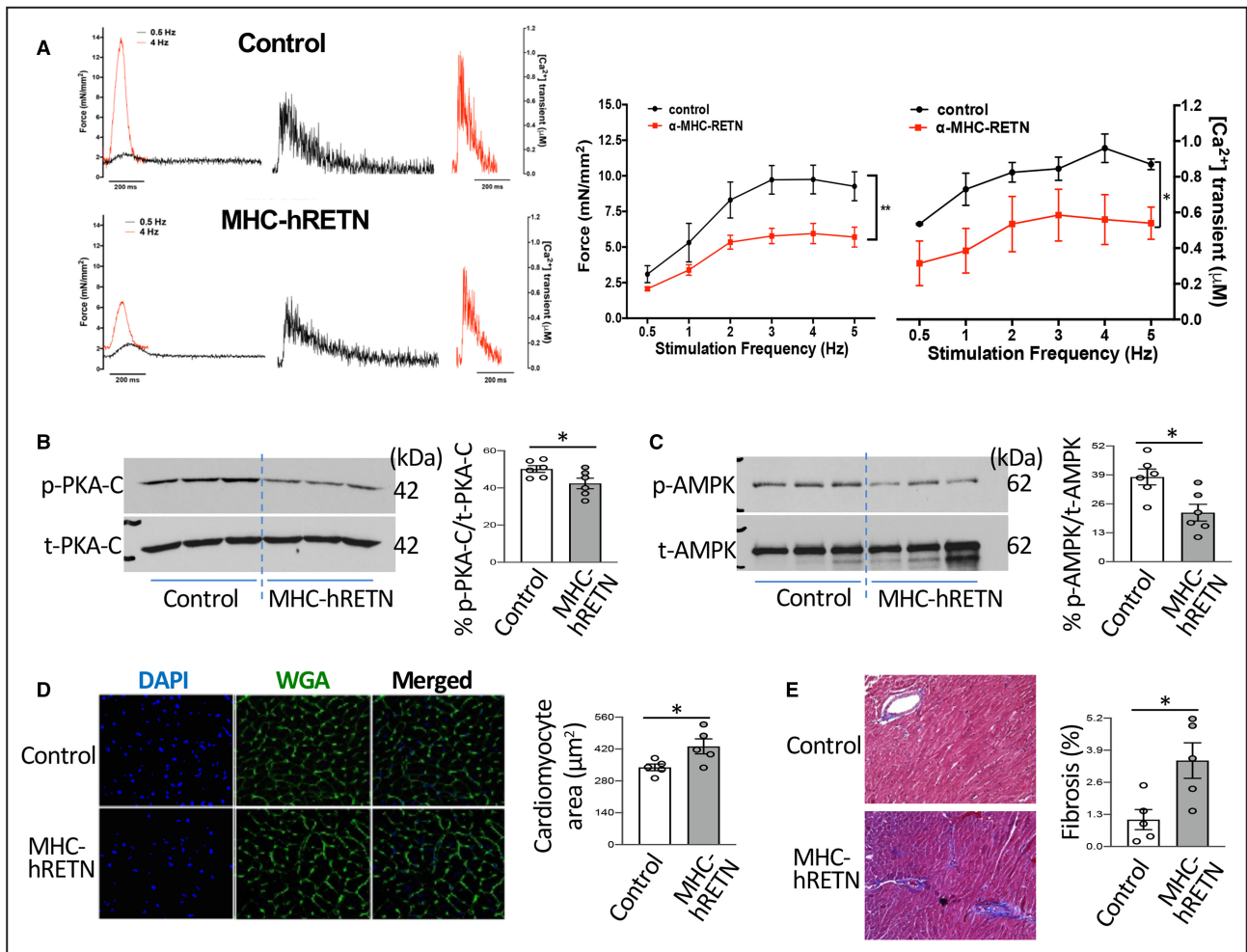
### Cardiac-Restricted Hresistin Overexpression Causes Cardiac Dysfunction and Remodeling

The Hresistin knock-in mice were generated through a direct cardiac overexpression (cross with MHC promoter) of recombinant Hresistin (Figure 2A and 2B). We validated Hresistin protein (Figure 2C) and gene (Figure 2D) expression in the RV tissues of these cardiac-specific Hresistin humanized mice. Functional assessment of trabecular muscle samples isolated from the MHC-Hresistin-overexpressing RVs revealed decreased twitch force and intracellular Ca<sup>2+</sup> transients (Figure 3A). This indicates depression of excitation–contraction coupling. We then tested key calcium-modulating protein kinases and found that phosphorylation of both PKA (Figure 3B) and AMPK (Figure 3C) were mitigated in the Hresistin-overexpressing cardiac tissues. In isolated myocytes, wheat germ agglutinin cell boundary staining revealed that Hresistin-overexpressing myocytes were larger than control myocytes (Figure 3D). Moreover, hearts that overexpressed Hresistin exhibited increased fibrosis, another key feature of remodeling (Figure 3E). These results indicate that cardiac Hresistin overexpression



**Figure 2. Generation of cardiac-specific Hresistin transgenic mice.**

**A** and **B**, Images present the nucleotide sequence for the Hresistin (hRETN) construct (**A**) and a schematic representation of the  $\alpha$ MHC-hRETN transgene structure (**B**). **C**, Immunofluorescence images of heart tissue samples from hRETN cardiac-overexpressing humanized mice and their corresponding littermate controls. Sections were stained with anti-myosin (green), co-stained with anti-FLAG (red), and counterstained with DAPI to validate the expression of the FLAG-tagged hRETN protein. Separate channels are displayed in middle and right panels, and digitally merged in left panels. Original magnification:  $\times 200$ . Boxed areas are shown at higher magnification ( $\times 400$ ) in the lower panels. Images are representative of 3 individual heart samples. **D**, Genotyping by PCR analysis of genomic DNA. Amplification of a 363-bp product encoding the myc-RETN epitope region of the transgene indicates that the humanized animals carry the knock-in hRETN gene in hearts. A 535-bp product was specifically amplified from animals carrying the tTA transgene. The 494-bp 18S housekeeping gene served as a control. DAPI indicates 4'-diamidino-2-phenyl-indole; MHC, myosin heavy chain; PCR, polymerase chain reaction; and tTA, tetracycline trans-activator.



**Figure 3. Cardiac dysfunction and remodeling in humanized mice that overexpress MHC-hRETN.**

**A**, Pooled data of force-frequency (left panel) and intracellular Ca<sup>2+</sup> transient-frequency (right panel) relationships in trabecular muscles from RV of wild-type and cardiac-specific MHC-hRETN-overexpressing mice (in force-frequency test: n=6 and 5 for control and overexpressing mice, respectively; in Ca<sup>2+</sup> transient-frequency test: n=4 animals per group; the 2 assays used samples from different animals). \*P<0.05, \*\*P<0.01. **B** and **C**, Changes in phosphorylation of PKA (**B**) and AMPK (**C**) in cardiac tissue from cardiac-specific MHC-hRETN-overexpressing mice and control littermates were determined by western blotting. Left panels show representative immunoblots. Right panels show quantitative analysis of expression. Data are shown as mean±SEM (n = 6 animals per group). \*P<0.05 vs control littermates. P- indicates phosphorylated protein; t- indicates total protein. **D**, Wheat germ agglutinin (WGA) cell boundary staining in the Hresistin-overexpressing myocytes. Left panels: heart tissue samples show cell nuclei (blue) and cell boundary (green). Magnification: ×400. Right panels: quantification of cell surface area based on histologic analysis of cardiomyocytes. Data represent means±SEM (n=5 animals per group). \*P<0.05 vs littermate control group. **E**, Masson's trichrome staining of heart tissue samples from MHC-Hresistin (RETN) humanized and littermate control groups. Magnification: ×200. Representative images (left panels) and quantitative analysis (right panels) were presented. Pooled data for quantification of fibrosis from 5 randomly selected histological fields at a magnification of ×400 on each slide. Data are shown as mean±SEM (n=5 animals per group). \*P<0.05 vs control group. AMPK indicates AMPK-activated protein kinase; DAPI, 4'6-diamidino-2-phenyl-indole; hRETN, human resistin; MHC, myosin heavy chain; and PKA, protein kinase A.

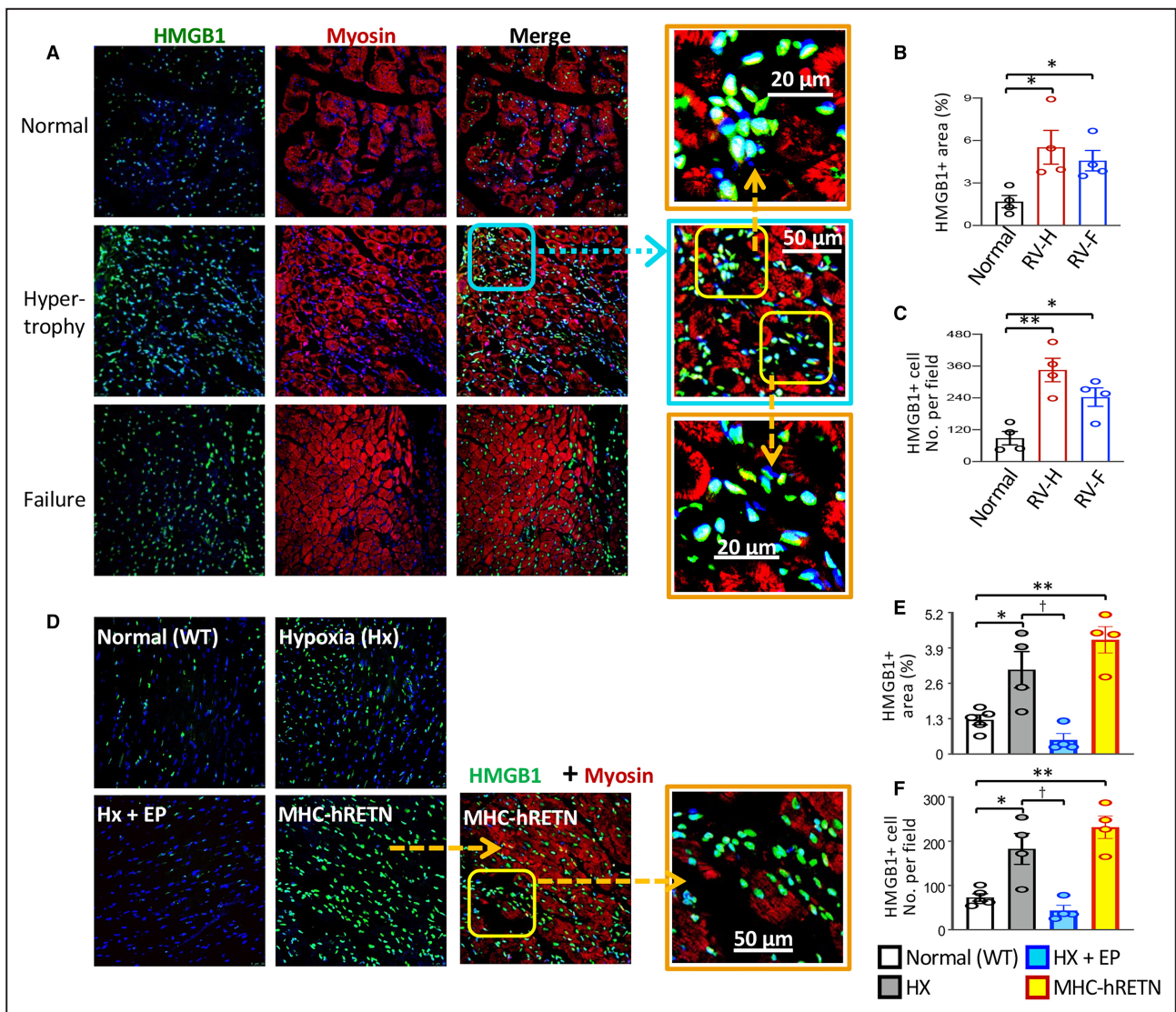
impairs cell function and induces myocyte hypertrophy and fibrosis.

### Hresistin Triggers DAMP-Dependent Inflammatory Signaling in RV

The Hresistin-inhibited AMPK phosphorylation (Figure 3C) suggested the DAMP pathway. As in a rodent lipopolysaccharide-induced acute lung injury model,

Hresistin-induced AMPK attenuation was linked to the activation of the alarmin HMGB1.<sup>35</sup> We also identified Hresistin/RELMα as an upstream regulator of HMGB1 in the hypoxic lung.<sup>16,17</sup> We explored the proinflammatory pathways triggered by Hresistin by examining HMGB1 signaling and observed high numbers of HMGB1-expressing cells in the hypertrophic and failing RV tissue of monocrotaline-treated rats (Figure 4A through 4C). Interestingly, a substantial number of HMGB1-positive,





**Figure 4. Hresistin activates HMGB1 signaling in the RV during RV r dysfunction.**

**A**, Immunofluorescence images of hypertrophic RV (RV-H, 2 weeks post-MCT induction) and failing RV (RV-F, 4 weeks post-MCT induction) from MCT-treated rats. Sections were stained with anti-HMGB1 antibody (green), costained with anti-myosin (red), and counterstained with DAPI (blue). Representative photographs of 4 individual animals per group. Original magnification:  $\times 200$ . Boxed region in the RV-H group is shown at higher magnification to the right ( $\times 400$ ). Further magnification ( $\times 1000$ ) of the 2 framed areas are shown in the upper and lower panels to illustrate the HMGB1-positive, myosin-negative cells that infiltrated the myocardial interstitium. **B** and **C**, Quantitative analysis of data in **A**. Percentage of area positive for HMGB1 signal (HMGB1+) in rat right heart determined with Adobe Photoshop software (**B**) and the number (No.) of HMGB1-positive cells counted (**C**) on 5 randomly chosen RV fields in each animal at 200-fold magnification. Data are presented as mean $\pm$ SEM (n=4 animals per group). \* $P < 0.05$ , \*\* $P < 0.01$  vs normal (non-MCT-treated) rats. **D**, Immunofluorescence images of RV tissue from wild-type (WT) mice on post-hypoxia day 4 and from MHC-hRETN humanized mice. Some hypoxic (Hx) mice received daily intraperitoneal injections of the HMGB1-specific inhibitor ethyl pyruvate (EP, 50mg/kg) for 4 days. Representative photographs of 4 individual animals per group. Original magnification:  $\times 200$ . In the MHC-hRETN group, the corresponding co-staining for HMGB1 (green) with myosin (red) is presented in the lower panel, and the boxed region in it is shown at higher magnification ( $\times 400$ ) on the left. **E** and **F**, Quantitative analysis of data in **D**. Percentage of area positive for HMGB1 signal in mouse right hearts was determined (**E**), and HMGB1-positive cells (per observed field) were counted (**F**). Data are presented as mean $\pm$ SEM (n=5 animals per group for the control [normal WT] group and n=4 animals per group for the other 3 groups [Hx, Hx+EP, and MHC-hRETN]; the dots in the graphs represent single individuals). \* $P < 0.05$ , \*\* $P < 0.01$  (increase) vs normal WT mice; † $P < 0.05$  (decrease) vs the hypoxia (Hx)-only group. DAPI indicates 4'-6-diamidino-2-phenyl-indole; EP, ethyl pyruvate; HMGB1, high mobility group box 1; hRETN, human resistin; hx, hypoxia; MCT, monocrotaline; MHC, myosin heavy chain; RV-F, failing right ventricle; and RV-H, hypertrophic right ventricle.

myosin-negative cells infiltrated into the interstitium of the RV tissues, indicating that accumulating immune cells are also a source of HMGB1. This DAMP signal peaked

in the hypertrophic myocardium (Figure 4A through 4C) and then declined slightly but remained elevated in the failing stage (Figure 4A through 4C).

We demonstrated that HMGB1-positive cell infiltration, also observed in the early onset of RVD (posthypoxic day-4, during the early PH inflammation phase)<sup>12,14</sup> (Figure 4D through 4F), was reversed by the specific HMGB1 inhibitor ethyl pyruvate (EP; Figure 4D through 4F). The upregulation of HMGB1 expression was even more striking in RV tissue that overexpressed Hresistin (Figure 4D through 4F).

Associated with HMGB1 elevation, we observed a marked increase in Ki67-positive cells in hypertrophic and failing RV tissue in the monocrotaline-induced rat PH model compared with that in the non-monocrotaline control rats (Figure 5A and 5B). The Ki67-positive cells were present during the early posthypoxia inflammatory phase (Figure 5C and 5D). Intriguingly, Hresistin knock-in induced a much stronger proliferation of Ki67-positive cells in RV tissues than did hypoxia exposure in wild-type mice (Figure 5C and 5D). These Hresistin-induced Ki67 signals were also observed in the interstitium of myosin-positive cardiomyocytes (Figure 5C) and had an expression pattern similar to that of HMGB1 (Figure 4D). Further colocalization analysis revealed that most of the Ki67-positive cells were the infiltrating leukocytes (F4/80+ macrophages and MPO+ neutrophils), and a few Ki67-positive cells were also the vimentin+ fibroblasts,<sup>36</sup> in the Hresistin-overexpressing RV tissues (Figure 5E), indicating the mechanism that Hresistin signaling induces RV remodeling through the inflammatory cytokine-associated cardiac fibrosis. The enhancement of this proliferative marker was reversed by administration of the HMGB1 antagonist EP (Figure 5C and 5D). Thus, Ki67 signal may mediate the effects of Hresistin/HMGB1 pathway axis to further amplify RV inflammation.

Collectively, these data suggest that Hresistin signaling triggers the HMGB1-dependent inflammatory response to initiate RVD. Moreover, it is a self-replenishing mechanism that sustains this proinflammatory state and continuously fuels RVD, thereby leading to RV remodeling and failure in chronic PH.

### Anti-Hresistin Antibody Ameliorates RV Dysfunction and Remodeling in the Rat PH Models

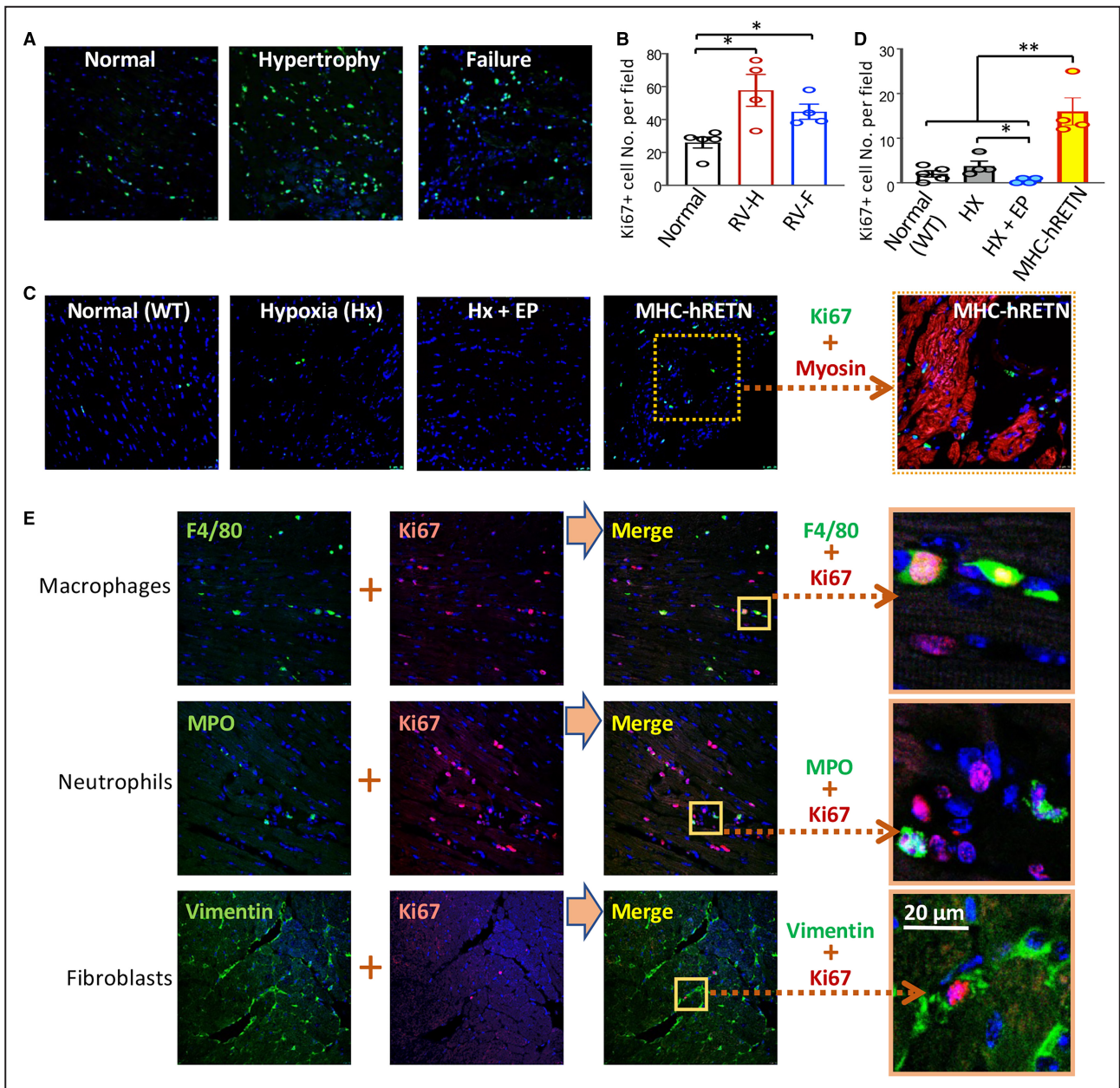
To validate the pathogenetic role of Hresistin in RV dysfunction and remodeling, we developed a series of therapeutic antibodies against Hresistin and identified a lead antibody that binds both Hresistin<sup>24</sup> and rodent RELM $\alpha$  (Figure 6C). When administered in the chronic hypoxia-induced PH rat model, this antibody ameliorated RV wall thickness (Figure 6A) and improved RV function (Figure 6B). Blocking Hresistin/RELM $\alpha$  with the neutralizing antibody also significantly reduced RV systolic pressure (Figure 6D) and RV hypertrophy (Figure 6E) in this rat PH model. Moreover,

in the monocrotaline-induced PH rat model, the anti-Hresistin therapeutic antibody prolonged survival of experimental animals in heart failure (Figure 6F). Mechanistically, in the RV tissues of these PH rats, hypoxia induced the production of RELM $\alpha$ , HMGB1, and the hypertrophic biomarkers atrial natriuretic peptide and  $\beta$ -MHC (Figure 7A and 7B). Consistently, in the cardiomyocytes isolated from neonatal rats, Hresistin overexpression mediated by adeno-associated virus transfection induced the expression of HMGB1, and  $\beta$ -MHC in vitro (Figure 7C and 7D). This was prevented by pretreatment with the anti-Hresistin/RELM $\alpha$  antibody (Figure 7C and 7D). In the neonatal rat cardiomyocytes, the Hresistin overexpression-induced hypertrophic proteins expression (atrial natriuretic peptide and  $\beta$ -MHC) (Figure 7E and 7F) and cell size increase (Figure 7G) were abrogated by pretreatment of the HMGB1 inhibitor EP, validating the role of HMGB1 in mediating the Hresistin-induced myocardial dysfunction in RV tissues during PH development. These findings suggest that our generated antibody against Hresistin/RELM $\alpha$  proteins has potential therapeutic value for human PH-associated RV dysfunction and failure.

## DISCUSSION

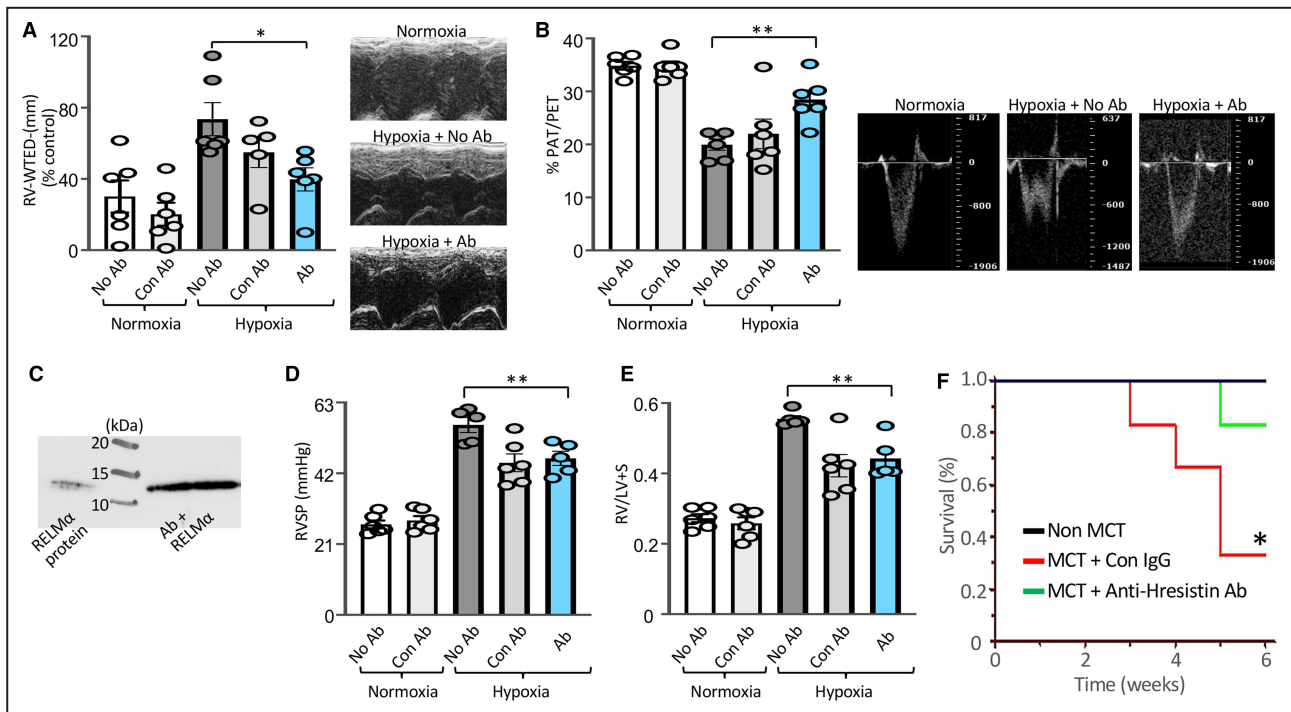
We present a novel concept that Hresistin signaling is a causal factor for the cardiac dysfunction and remodeling (Figure 8) that parallels the Hresistin/RELM $\alpha$ -induced vascular remodeling in lungs during PH. This study directly links RELMs to RVD, highlighting their unique role in the immune interaction between lungs and the RV in PH. The humanized mouse model of cardiac-restricted Hresistin overexpression enabled us to explore the specific cardiac action of Hresistin signaling from a more translational and mechanistic perspective. Moreover, the newly developed anti-Hresistin antibody had beneficial effects against RVD and may hold promise for preventing or reversing RV failure in human disease.

The RV has distinguishing physiologic properties in response to pathologic insults,<sup>37</sup> but the mechanism underlying PH-associated RV dysfunction is still unclear. Based on our mechanistic data, we believe Hresistin triggers DAMP-associated inflammation that eventually leads to RV failure. The kinases PKA and AMPK govern Ca<sup>2+</sup> channels, force-frequency response, and myofilament behavior in cardiomyocytes.<sup>38–48</sup> Dysregulation of these pathways could decrease cardiac contractility as we observed in the mouse Hresistin-overexpressing RV. These kinases have been implicated in inflammatory pathogenesis. PKA was involved in inflammasome activation in macrophages,<sup>49,50</sup> indicating a link to HMGB1 signaling, given the reported inflammasome-dependent release



**Figure 5. Hresistin/HMGB1 signaling axis upregulates Ki67 expression in RV.**

**A**, Sections of RV tissues during hypertrophy or failure (as described in above panel A) were stained with anti-Ki67 antibody (green) and counterstained with DAPI (blue). Representative photographs of 4 individual animals per group. Magnification:  $\times 200$ . **B**, Quantitative analysis of data from **A**. Ki67-positive (Ki67+) cells were counted on 5 randomly chosen fields of RV sections in each animal at  $\times 200$  magnification. Data are presented as mean  $\pm$  SEM ( $n=5$  animals per group for the normal control group and  $n=4$  animals per group for the groups of RV-H and RV-F; the dots in the graphs represent single individuals).  $*P<0.05$  vs normal (non-MCT-treated) rats. **C**, Immunofluorescence images of RV tissue from WT mice exposed to 4 days of hypoxia with or without the HMGB1 inhibitor EP and from cardiac-specific MHC-hRETN-overexpressing mice. The boxed region in the fourth panel is shown at higher magnification ( $\times 400$ ) on the right with costaining for Ki67 (green) and myosin (red). Representative photographs of 5 individual animals per group. Original magnification:  $\times 200$ . **D**, Quantitative analysis of data in **C**. The Ki67-positive cells were counted (per observed field). Data are presented as mean  $\pm$  SEM ( $n=5$  animals per group for the control [normal WT] group and  $n=4$  animals per group for the other 3 groups [Hx, Hx+EP, and MHC-hRETN]).  $*P<0.05$ ,  $**P<0.01$ . **E**, In the RV tissues of the MHC-hRETN-overexpressing mice, Ki67 (red, middle panels) was further costained with the markers (green, left panels) of macrophages (F4/80), neutrophils (MPO), or fibroblasts (Vimentin). Images were merged in the right panels. The boxed areas were further enlarged in the far right panels showing the double positive cells. Representative photographs of 4 individual animals per group. Original magnification:  $\times 400$ . DAPI indicates 4'-6-diamidino-2-phenyl-indole; EP, ethyl pyruvate; HMGB1, high mobility group box 1; h-RETN, human resistin; Hx, hypoxia; MCT, monocrotaline; MHC, myosin heavy chain; MPO, myeloperoxidase; RV-F, failing right ventricle; RV-H, hypertrophic right ventricle; and WT, wild type.



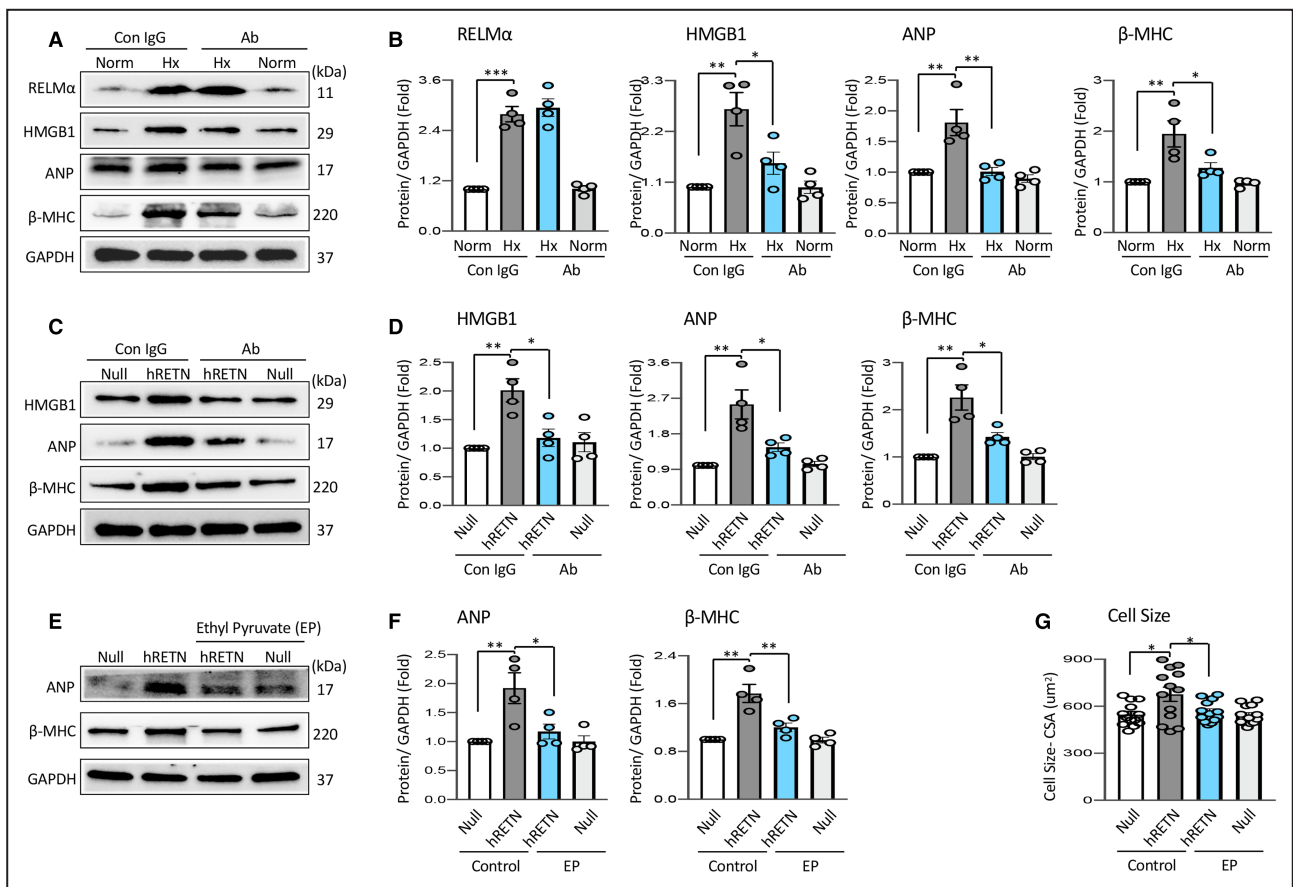
**Figure 6. Anti-Hresistin human antibody ameliorates RV dysfunction in rats with PH.**

The anti-Hresistin antibody (Ab) or the isotype-matched control IgG1 (Con IgG) at 4 mg/kg were administered intraperitoneally twice a week in the hypoxia-induced PH rats. **A** and **B**, Echocardiographic analysis of right ventricular (RV) wall thickness and pulmonary artery blood velocity in Ab-treated hypoxic rats. RV wall thickness external diameter (RV-WTED) was measured as the distance from the free wall to the interventricular septum (millimeter) in the parasternal long-axis view using M-mode (**A**). Data are expressed as a percentage of the value of normoxic control mice. The anti-Hresistin Ab treatment also lengthened pulmonary artery acceleration time (PAT). Results of pulsed wave Doppler measurement of PAT are shown in **B**. PAT values were normalized by pulmonary ejection time (PET). Data are expressed as means $\pm$ SEM ( $n=6$  animals per group). \* $P<0.05$ , \*\* $P<0.01$  vs hypoxia (no Ab) group. Representative echocardiographic images are shown in the right panels. **C**, Immunoprecipitation analysis of the binding of rat RELM $\alpha$  to the human therapeutic Ab targeting Hresistin. The protein-Ab binding was detected by western blotting with the anti-Hresistin antibody from R&D (AF1359). Recombinant rat RELM $\alpha$  protein was loaded as the positive control. **D** and **E**, Analysis of RV hypertrophy and hemodynamics in the hypoxia (Hx)-induced rat PH model. We measured the RV systolic pressure (RVSP) (**D**) and Fulton index (ratio of RV weight/LV+S weight) (**E**). Data are presented as means $\pm$ SEM (normal no Ab:  $n=6$ , normal con Ab:  $n=6$ , hypoxia no Ab:  $n=5$ , hypoxia con Ab:  $n=6$ , hypoxia Ab:  $n=5$ ;  $n$  refers to the number of animals per group). \*\* $P<0.01$  vs Hx-treated group without Ab treatment. **F**, Results of treatment with the human antibody targeting Hresistin to improve the survival rate among rats with monocrotaline-induced PH. \* $P<0.05$  by log rank test;  $n=6$  animals in each group. LV+S indicates left ventricle plus septum; MCT, monocrotaline; PH, pulmonary arterial hypertension; and RELM $\alpha$ , resistin-like molecule- $\alpha$ .

of HMGB1.<sup>51–54</sup> More notably, in humanized resistin mice with lipopolysaccharide-induced acute lung injury, Hresistin reduced AMPK and elevated HMGB1 in lungs,<sup>35</sup> consistent with our cardiac findings (Figures 3 and 4). These results suggest that HMGB1 is a key inflammatory element associated with RV dysfunction observed in Hresistin-induced RVD.

Inflammation is believed to be a mechanism that underlies RVD onset and progression.<sup>55</sup> Hresistin and its rodent homolog RELM $\alpha$  have been identified as TH2 inflammatory mediators in pulmonary diseases. Our recent studies in lungs showed that Hresistin/RELM $\alpha$  triggers HMGB1 signaling to mediate pulmonary vascular inflammation,<sup>16,17</sup> which is associated with the Hresistin/RELM $\alpha$ -induced recruitment of macrophages.<sup>17</sup> An alarmin, HMGB1, acts as an endogenous danger signal to regulate inflammatory response

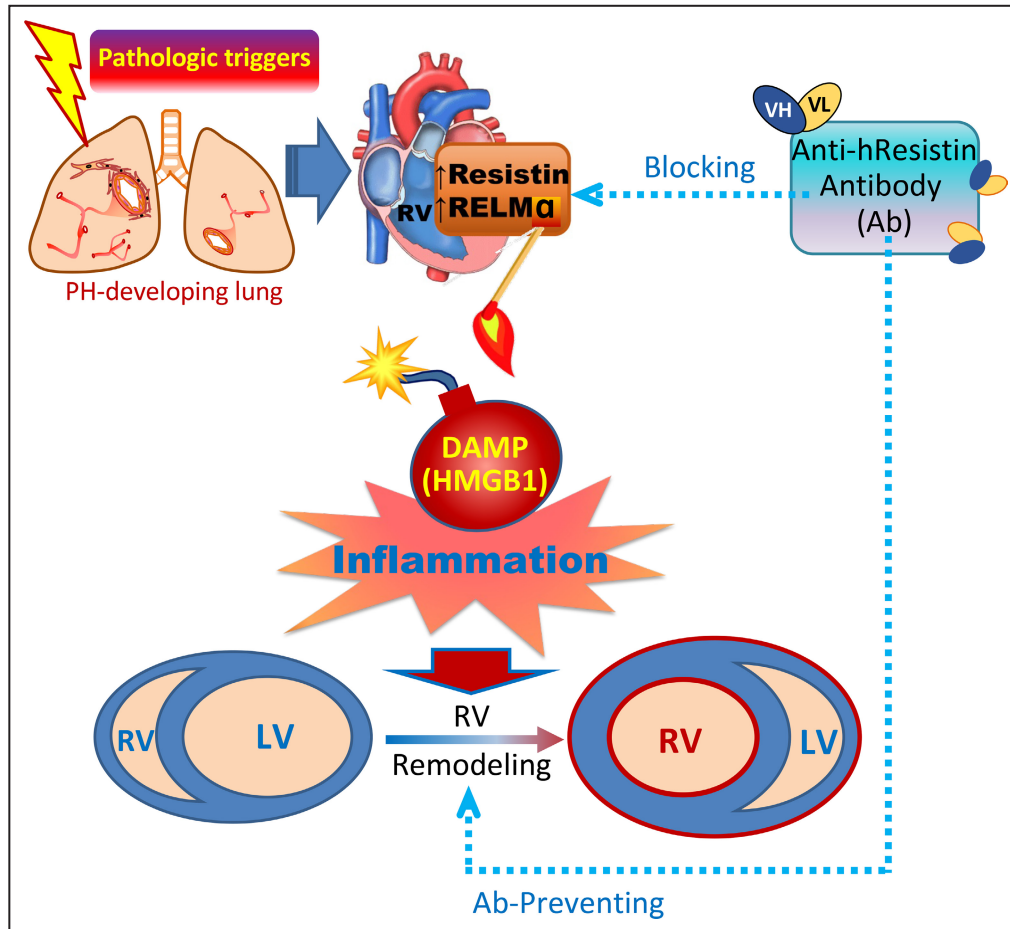
through an autocrine and paracrine manner by activating RAGE or TLR4.<sup>56,57</sup> Moreover, HMGB1 has been reported to have a critical role in inflammatory heart diseases.<sup>58,59</sup> Here, in our humanized mouse model, we observed activation of HMGB1 and the consequent ki67 in the RV. The proliferation of macrophages has been determined as a mechanism of RV failure in PH by studies on the monocrotaline-induced rat PH model.<sup>60</sup> In the current study, the Hresistin/HMGB1 signaling axis amplifies inflammation through enhancing the proliferation (Figure 5E) and recruitment of the immune cells in the RV and possibly expression in the myocyte itself. These findings suggest that Hresistin-triggered DAMP signaling not only ignites but also fuels the sterile inflammatory response during RVD onset and progression. Hresistin-initiated DAMP-dependent cardiac inflammation is likely an early response to injury



that leads to subsequent metabolic disarrangement. Thus, inflammation, metabolic derangement, impaired cardiac function, and adverse structural remodeling may occur on a time continuum that is initiated by the metabolism-regulating activity of HMGB1.<sup>61</sup> Additional study is required to elucidate this mechanism.

The time course over which Hresistin/RELM $\alpha$  activates inflammation may also help shed light on the transition from adaptive remodeling to maladaptive RV failure in patients with PH. Inflammatory cells are more numerous in the RV of patients with SSc-associated PH than in that of patients with other types of PH.

Additionally, the high number of inflammatory cells is associated with the morbidity of RV failure in these patients,<sup>55,62</sup> suggesting that inflammation contributes to development of RV failure. As a significant proportion of the WHO group-1 PH, the SSc-PH heralds a particularly poor prognosis compared with patients with subtypes of the group-1 PH including idiopathic PH, despite having similar measures of RV load.<sup>63,64</sup> Inflammation is a pathogenetic factor of cardiac involvement in SSc-PH.<sup>63</sup> Mechanistically, compared with idiopathic PH, SSc-PH exhibits depressed RV contractile reserve,<sup>65</sup> associated with reduced calcium



**Figure 8. Schematic illustration of Hresistin-induced cardiac inflammation and dysfunction.**

During pulmonary arterial hypertension (PH) development, Hresistin activates the damage-associated molecular pattern (DAMP), signaling triggering of inflammation in the right ventricle (RV) and contributing to RV dysfunction pathogenesis. Targeting the Hresistin signaling cascade may constitute a novel therapeutic approach to RV dysfunction and other related cardiac diseases in humans. HMGB1 indicates high mobility group box 1; Hresistin, human resistin; LV, left ventricle; and RELM $\alpha$ , resistin-like molecule- $\alpha$ .

recycling, which is caused by PKA suppression.<sup>65,66</sup> Given the amplified inflammation (Figure 4), fibrosis (Figure 3E), impaired contractility, decreased intracellular Ca<sup>2+</sup> availability (Figure 3A), and the suppressed PKA (Figure 3B) in the Hresistin-overexpressing RV tissues, it suggests the activation of Hresistin signaling in the RV of SSc-PH (Figure 1B) as the pathogenetic mechanism of RV dysfunction, remodeling, and failure of this PH subform. Moreover, we have observed RELM $\alpha$  activation as a key pathogenetic mechanism in hypoxia- and monocrotaline-induced models of PAH and RV dysfunction,<sup>1</sup> suggesting Hresistin as a critical pro-inflammatory mediator of RV remodeling and failure in a broad array of human PAH diseases. Nevertheless, further studies are warranted on the role of Hresistin signals in RV pathology of other PAH subtypes. The role of inflammation in LV failure has been better studied,<sup>67,68</sup> but anti-inflammatory therapy for LV failure has failed so far.<sup>55</sup> In our study, Hresistin/RELM $\alpha$

activation in the RV during PH development, and Hresistin/RELM $\alpha$ -mediated accumulation of HMGB1- and Ki67-producing inflammatory cells, all peaked in the hypertrophied RV and declined in the failing RV (Figures 1, 4, and 5). These findings suggest that RELM-DAMP signaling may facilitate the early, intensified damage-triggered inflammatory response that leads to later pathologic remodeling. This response may be responsible for the RV's transition from a compensated to a decompensated state. Hresistin thereby could constitute a therapeutic target for preventing RV dysfunction and the subsequent progression to RV failure. This possibility is supported by the promising results of the therapeutic anti-Hresistin antibody, which ameliorated cardiac remodeling and prolonged survival of PH rats (Figures 6 and 7), given that survival in PH is closely related to RV function.<sup>69</sup>

One limitation of our work is that although we focused on the RV, the MHC-Hresistin humanized model

is not RV-specific. It is difficult or impossible to create an animal model with exclusive RV expression. To date there are no RV-specific markers to target the genetic expression specifically to the right heart. In addition, given that immunoregulation by Hresistin/RELM $\alpha$  leads to fibrosis in lungs,<sup>70</sup> and that we observed Hresistin-induced heart fibrosis in the humanized mice (Figure 3), future studies are warranted to explore the proinflammation-to-profibrotic phenotype switch in RELM-DAMP-activated immune cells and myocytes, and to test the antifibrotic properties of the anti-Hresistin antibody. Nevertheless, we were able to show that Hresistin activation triggers RV inflammation, metabolic derangement, and subsequent hypertrophy that leads to the initiation and progression of right ventricular failure in PH and that blocking Hresistin signaling with a human antibody may prevent or reverse the PH-associated RVD in humans.

## ARTICLE INFORMATION

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### Disclosures

None.

### Supplemental Material

Figure S1

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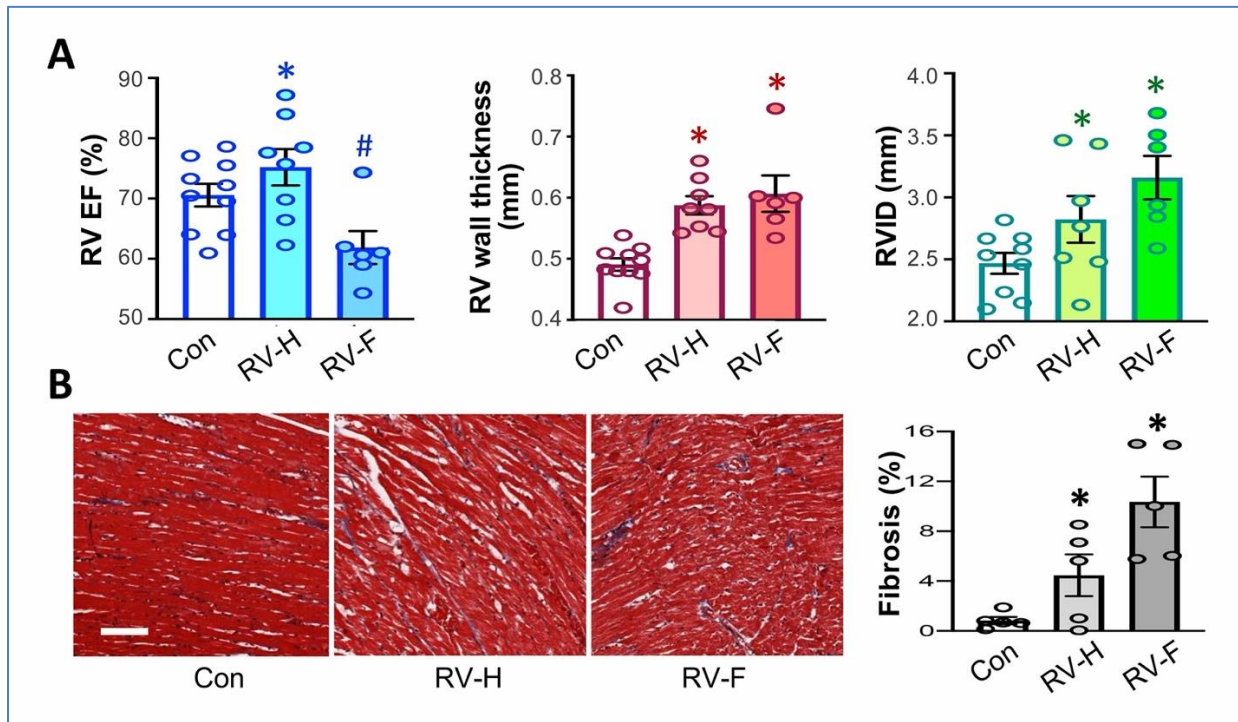
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# **SUPPLEMENTAL MATERIAL**

**Figure S1. Monocrotaline (MCT)-induced pulmonary hypertension causes right ventricular hypertrophy (RV-H) and failure (RV-F) in rats.**



**A**, RV ejection fraction (EF), wall thickness and chamber diameter during development of RV-H and RV-F. We designated RV-H as compensated right ventricular hypertrophy with maintained RV EF (i.e. between 2.5-3.5 weeks after MCT injection), and RV-F as decompensated RV failure (i.e. after 4.5 weeks MCT injection when RV EF was significantly reduced). In EF test, control (Con, normal): n = 10, RV-H: n = 8, RV-F: n = 6. In RV wall thickness test, Con: n = 10, RV-H: n = 8, RV-F: n = 6. In RV chamber diameter (RVID) test, Con: n = 9, RV-H: n = 7, RV-F: n = 6. The n refers to the number of animals per group; the dots in the graphs represent single individuals. \*,# $p < 0.05$  vs. 0 week (control). **B**, Masson staining of RVs of control, hypertrophied, and failed hearts (*left panels*) and quantification of fibrosis (*right panel*, data are means  $\pm$  SEM, n = 5 animals per group, \* $p < 0.05$  vs. control). Bar = 100  $\mu$ m.