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## **Down-Regulation of TRPM8 in Pulmonary Arteries of Pulmonary Hypertensive Rats**

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## Abstract

**Background**—Pulmonary hypertension (PH) is characterized by profound vascular remodeling and alterations in Ca<sup>2+</sup> homeostasis in pulmonary arterial smooth muscle cells (PASMCs). Multiple transient receptor potential melastatin-related (TRPM) subtypes have been identified in vascular tissue. However, the changes in the expression and function of TRPM channels in pulmonary hypertension have not been characterized in detail.

**Methods**—We examined the expression of TRPM channels and characterized the functions of the altered TRPM channels in two widely used rat models of chronic hypoxia (CH)- and monocrotaline (MCT)-induced PH.

**Results**—CH-exposed and MCT-treated rats developed severe PH and right ventricular hypertrophy, with a significant decrease in TRPM8 mRNA and protein expression in pulmonary arteries (PAs). The downregulation of TRPM8 was associated with significant reduction in menthol-induced cation-influx. Time-profiles showed that TRPM8 down-regulation occurred prior to the increase of right ventricular systolic pressure (RVSP) and right ventricular mass index (RVMI) in CH-exposed rats, but these changes were delayed in MCT-treated rats. The TRPM8 agonist menthol induced vasorelaxation in phenylephrine-precontracted PAs, and the vasorelaxing effects were significantly attenuated in PAs of both PH rat models, consistent with decreased TRPM8 expression.

**Conclusion**—Downregulation of TRPM8 may contribute to the enhanced vasoreactivity in PH.

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This is an Open Access article licensed under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs 3.0 License (www.karger.com/OA-license), applicable to the online version of the article only. Distribution for non-commercial purposes only. Mo-Jun Lin, M.D. Ph.D. and James S.K. Sham, Ph.D., Department of Physiology and Pathophysiology, Fujian Medical University, 1 Xueyuan Road, Shangjie Zhen, Minhou County, Fuzhou, Fujian Province 350108 (P.R. China) and Division of Pulmonary and Critical Care Medicine, Johns Hopkins Medical Institutions, 5501 Hopkins Bayview Circle, Baltimore, MD 21204 (USA), mjlin@mail.fjmu.edu.cn (Mo-Jun Lin), jsks@jhmi.edu (James S.K. Sham) <sup>d</sup>The authors have equal contribution to this publication

## Keywords

Transient receptor potential melastatin; Pulmonary hypertension; Calcium signaling; Monocrotaline; Chronic hypoxia; Menthol

## Introduction

Pulmonary hypertension (PH) is characterized by profound vascular remodeling, right ventricular hypertrophy and alterations in Ca<sup>2+</sup> homeostasis in pulmonary arterial smooth muscle cells (PASMCs) [1]. It is associated with a board spectrum of diseases with different pathological features and etiological mechanisms. They are classified into five categories, namely, pulmonary arterial hypertension (PAH), PH owing to left heart disease, PH owing to lung disease and/or hypoxemia, chronic thromboembolic pulmonary hypertension, and PH with unclear multifactorial mechanisms [2]. The etiological mechanisms of the various forms of PH are different, but they all have the common features of abnormalities in pulmonary vascular function, vascular cell proliferation, and remodeling, suggesting that they may share some important signaling mechanisms in the development of the disease.

 $Ca^{2+}$  signaling plays vital roles in regulation of numerous physiological and pathophysiological processes, including contraction, proliferation, hypertrophy and migration of PASMCs [3]. Previous studies indicate that there are major alterations in ion channel expression and  $Ca^{2+}$  homeostasis, such as membrane depolarization, downregulation of voltage-gated K<sup>+</sup> channels, upregulation of canonical transient receptor potential (TRPC) channels, increase in  $Ca^{2+}$  influx, and elevation of resting intracellular  $Ca^{2+}$  concentration ( $[Ca^{2+}]_i$ ) in PASMCs of idiopathic PAH patients and animal models of PH [4–8].  $Ca^{2+}$  influx in PASMCs is mainly regulated by voltage-dependent  $Ca^{2+}$  channels (VDCC) and voltage-independent nonselective cation channels.

Members of transient receptor potential (TRP) gene family are known to encode different types of nonselective cation channels. They play critical roles in numerous cellular functions activated by a wide variety of chemical and physical stimuli including agonist, mechanical stress, temperature, light and sound to modulate calcium influx. To date, 28 mammalian TRP homologues have been found in a wide variety of cells and tissues, and at least 10 TRPs have been identified as functional channels in vascular smooth muscle cells (VSMCs) [9]. They are involved in many different vascular functions, such as myogenic response, agonist-induced vasoconstriction, VSMCs proliferation, and vascular remodeling.

The canonical TRP (TRPC) channels have been implicated as store-operated calcium entry (SOCC) and receptor-operated calcium entry (ROCC) in vascular smooth muscle cells. The expression of TRPC1 and TRPC6 mRNAs and proteins are upregulated, and SOCE and ROCE are potentiated in PASMCs of hypoxic PAH rats [4]. Furthermore, expression of TRPC1 and SOCE are also enhanced in PASMCs of MCT-induced PH rats [8]. These results suggest that TRPC1-SOCE may be a common signaling pathway for the development of PH. Compared to TRPC, the physiological roles of other TRP subfamilies are much more elusive in pulmonary vascular smooth muscle. A recent study shows that the vanilloid related TRP (TRPV4) channel operates as a mechanosensitive cation channel in

PASMCs, and the expression of TRPV4 channel is upregulated in PAs of chronic hypoxic rats [10]. Moreover, the upregulation of TRPV4 in PAs is associated with the appearance of myogenic tone, and deletion of TRPV4 gene delayed and suppressed the development of PH.

The melastatin TRP subfamily, which consists of eight mammalian members (TRPM1-8), are known to participate in tumor suppression, reactive oxygen species (ROS)-induced apoptosis, Mg<sup>2+</sup> homeostasis, and thermo-sensing (hot and cold) in nonvascular tissues [11-13]. A few recent studies have revealed that some of these TRPM channels are expressed and may play different physiological roles in vascular smooth muscles [10, 14–15]. It has been proposed that TRPM4 contributes to membrane depolarization and vasoconstriction associated with increased intraluminal pressure in cerebral arteries [16-18]. TRPM7 was identified as a functional regulator of Mg<sup>2+</sup> homeostasis in mouse and rat mesenteric and aortic smooth muscle cells [19]. Application of the TRPM8-specific agonist menthol or icilin induced small vasoconstriction in relaxed vessels, but caused vasodilation in endothelium-denuded rat tail arteries pre-contracted with KCl or the a-adrenoceptor agonist phenylephrine (PHEN) [15]. We have previously shown that TRPM2, TRPM3, TRPM4, TRPM7 and TRPM8 were expressed in intralobar pulmonary arteries (PAs) and aorta of rats, and TRPM8 was the most abundantly expressed TRPM subtype [14]. Activation of TRPM8 channels in PASMCs with menthol induced robust increase in  $[Ca^{2+}]_i$  which could be abolished by removal of extracellular  $Ca^{2+}$  or by applying the non-selective cation channel blocker Ni<sup>2+</sup>. These results indicate that multiple functional TRPM channels are expressed in rat intralobar PAs and these novel Ca<sup>2+</sup> entry pathways may play important roles in the regulation of pulmonary and systemic circulation.

However, alternations of the expression and function of TRPM channels in PH have not been studied in detail [10]. The present study used two classical rat models of chronic hypoxia-induced PH and MCT-induced PH to examine the changes in the expression and functional activities of TRPM channels, and to explore their contribution to the development of PH.

## **Materials and Methods**

### PH models

Chronic hypoxia (CH)-induced pulmonary hypertension was produced by the established method. Male Sprague-Dawley rats (200~250g) were placed in a hypoxic chamber and exposed to either normoxia or normobaric hypoxia for 3~4 weeks. The chamber was continuously flushed with either room air or a mixture of room air and N<sub>2</sub> to maintain oxygen level at 10%. Chamber O<sub>2</sub> and CO<sub>2</sub> were monitored continuously (OM-11 oxygen analyzer). Rats were exposed to room air for 10 min twice a week to clean the cages, and to replenish food and water supplies. MCT-induced PAH were generated in adult male Sprague-Dawley rats (200~250g). Rats were given a single intraperitoneal injection of MCT (60 mg/kg) or an equivalent volume of saline (2ml/kg). MCT (Sigma) was dissolved in 1 M HCl, neutralized to pH 7.4 with 1 M NaOH, and diluted with saline [8]. Twenty-one days after MCT injection, rats in the MCT and sham injection groups were anesthetized with urethane (1g/kg). Right ventricle systolic pressure (RVSP) and mean systemic arterial

pressure (MSAP) were measured by accessing the right ventricle through the jugular vein and right carotid artery, respectively, using polyethylene catheters connected to pressure transducers (YPJ01; Chengyi, China). Pressure signals were displayed continuously on an RM6240 polygraph (Chengyi, China). Heart rate was determined from the right ventricle pressure pulse. At the end of hemodynamic measurement, the rat was sacrificed with an overdose of urethane. The heart was removed, and right ventricular mass index (RVMI) was calculated as the ratio of wet weight of the right ventricle to the left ventricular wall plus septum [RV/(LV+S)]. In the experiment on the time-course of change of TRPM8 expression and PH development, animals were sacrificed on 0, 1, 3, 5, 7, 14, and 21 days after exposure to CH exposure or MCT treatment. All procedures were approved by the Animal Care and Use Committee of Fujian Medical University.

## Isolation and culture of PASMCs

PASMCs were enzymatically isolated and transiently cultured as previously described [8]. Briefly, male Sprague-Dawley rats were injected with heparin and anesthetized with sodium pentobarbital (130mg/kg i.p.). They were exsanguinated and the lungs were removed and transferred to a petri dish filled with HEPES-buffered salt solution (HBSS) containing (in mM) 130 NaCl, 5 KCl, 1.2 MgCl<sub>2</sub>, 1.5 CaCl<sub>2</sub>, 10 HEPES, 10 glucose, pH 7.4 (adjusted with NaOH). Third and fourth generation intrapulmonary arteries (~300 to 800µm) were isolated and cleaned free of connective tissue. The endothelium was removed by gently rubbing the luminal surface with a cotton swab. Arteries were then allowed to recover for 30 min in cold (4°C) HBSS, followed by 20 min in reduced-Ca<sup>2+</sup> (20µM) HBSS at room temperature. The tissue was digested at 37°C for 20min in 20µM Ca<sup>2+</sup> HBSS containing collagenase (Type I, 1750U/ml), papain (9.5 U/ml), bovine serum albumin (2mg/ml), and dithiothreitol (1mM), then removed and washed with  $Ca^{2+}$ -free HBSS to stop digestion. PASMCs were gently dispersed by trituration with a small-bore pipette in Ca<sup>2+</sup>-free HBSS at room temperature. The cell suspension was then placed on 25mm glass coverslips in Ham's F-12 medium (with L-glutamine) supplemented with 0.5% fetal calf serum, 100U/ml of streptomycin, and 0.1 mg/ml of penicillin. PASMCs from chronic hypoxic and normoxic animals were transiently (~24 hours) cultured inside a modular incubator chamber (billups-rothenberg, Inc.) under 4% O<sub>2</sub>/5% CO<sub>2</sub> and 21% O<sub>2</sub>/5% CO<sub>2</sub>, respectively.

### Quantitative real-time RT-PCR

De-endothelialized PAs frozen in liquid nitrogen were homogenized mechanically. Total RNA was extracted using an RNeasy mini kit (Qiagen, Valencia, CA). Total RNA (1µg) was used for first-strand cDNA synthesis using Prime Script reverse transcriptase kit (DRR037A, Takara, Japan) according to the manufacturer's protocol. Quantitative real-time RT-PCR was used to quantify the changes in the expression of TRPM subtypes. PCR reactions were set up with iQ SYBR Green PCR Supermix (Bio-Rad, Hercules, CA) (Roche, Germany) using 0.25µl of cDNA as the template in each 10µl reaction mixture. Gene-specific real-time PCR primers for TRPM channels are listed in Table 1. The PCR protocol, consisting of an initial step at 95°C for 10min, followed by 40 cycles at 95°C for 15 s, 60°C for 1 min, and 72°C for 1 min, was performed using the Multicolor Real-Time PCR Detection System (ABI, USA). In the experiments for determining the time course of change in TRPM8 expression, the relative quantities of TRPM8 were calculated using the CT

method with  $\beta$ -actin as the internal control. TRPM8 values measured from samples at various time points were normalized to the averaged value of the control samples.

#### Western blot analysis

De-endothelialized PAs were quickly frozen in liquid nitrogen. The frozen tissues were crushed and homogenized using a mortar and pestle and then resuspended in ice-cold cell lysis buffer containing 50 mM Tris-HCl (pH 7.4), 150mM NaCl, 1% deoxycholic acid, 0.1% SDS, 0.5% Nonidet P-40, and protease inhibitor cocktail (Roche, Mannheim, Germany). The homogenate was centrifuged at 4°C and 14,000g for 15min, then the supernatant was collected, and the protein concentration was estimated using the bicinchoninic acid assay. The protein sample (40µg) was resolved in an 8% SDS-PAGE and electrotransferred onto a PVDF membrane. The membrane was blocked with 5% (wt/vol) nonfat dry milk in TBS containing 0.05% Tween 20 (TBST) for 2h at room temperature, followed by incubation at 4°C overnight with a monoclonal anti-TRPM8 antibody (ACC-049, 1:300 dilution; Almone). The PVDF membrane was then washed with TBST. After being washed, the membrane was incubated with goat-anti-rabbit secondary antibody (no. 7074, 1:5000 dilution; Cell Signaling Technology) at room temperature for 1 h. Excess secondary antibody was washed again, the bound secondary antibody was detected with enhanced chemiluminescence (Pierce, Rockford, IL), and images were taken using a Gel Logic 200 image system (Kodak, New Haven, CT).

#### **Isometric contraction**

Intralobar PAs (~300 to 800  $\mu$ m) were isolated and placed in oxygenated modified Kreb's solution containing (in mM) 118 NaCl, 4.7 KCl, 1.2 MgCl<sub>2</sub>, 10 HEPES, 11.1 dextrose, and 2.0 CaCl<sub>2</sub>. They were cleaned of connective tissue and cut into 4 mm lengths. Endothelium was disrupted by gently rubbing the lumen with a small wooden stick, and the arterial rings were suspended between two stainless steel stirrups in organ chambers filled with modified Kreb's solution for isometric tension recording. The solution was gassed with 95% O<sub>2</sub>+5% CO<sub>2</sub> to maintain a pH of 7.4 and the temperature at 37°C. Isometric contraction was measured using a strain gauge connected to a polygraph. Resting tension was adjusted to 0.8~1.0g. Arteries were exposed to 60mM KCl to establish maximum contraction and to phenylephrine (3×10<sup>-7</sup>M) followed by acetylcholine (10<sup>-6</sup>M) to verify complete disruption of endothelium.

## Mn<sup>2+</sup> quenching of Fura-2

Rate of Ca<sup>2+</sup> entry through TRPM8 was quantified by quenching of fura-2 with  $Mn^{2+}$ . PASMCs from control, CH- and MCT-treated rats were loaded with fura-2AM as described above. Fura-2 was excited at the Ca<sup>2+</sup>-insensitive isobestic point of 360nm, and emission light was recorded at 510 nm (PTI, USA). PASMCs were then bathed in a Ca<sup>2+</sup> free (with 0.1 mM EGTA) Tyrode solution containing 10  $\mu$ M nifedipine. After a stable baseline fluorescent measurement was attained, 500  $\mu$ M Mn<sup>2+</sup> was applied through a concentrationclamp system with the multi-barrel pipette positioned <50  $\mu$ m from PASMCs. The rates of quenching of fura-2 fluorescence in PASMCs with/without drug treatments were determined and compared.

## Statistical analysis

Data are expressed as means  $\pm$  SE, and n indicates the number of animals, cell samples, or PA rings used. Statistical significance was assessed using unpaired or paired Student's t-tests and ANOVA wherever appropriate. Differences were considered significant at P<0.05. Curve fitting was performed using the SigmaPlot11.0 software.

## Results

## Verification of CH- and MCT-induced PH models

CH- and MCT-treated rats exhibited profound PH and right ventricular hypertrophy when examined 3 weeks after hypoxia exposure or MCT injection, respectively. PH was generally more severe in the MCT-treated rats. RVSP increased by two-folds in CH-exposed rats, and almost tripled in MCT-treated rats (control:  $25.1\pm0.9$  mmHg, n=29; MCT:  $60.6\pm4.2$  mmHg, n=18, P<0.01, CH:  $52.6\pm3.2$  mmHg, n=14; P < 0.01). RVMI was about 40% higher for CH rats and was nearly doubled in MCT rats when compared with that of normoxic controls (control:  $27.5\pm0.5\%$ , n=29; MCT:  $59.9\pm4.9\%$ , n=18, P<0.01, CH:  $38.5\pm0.7\%$ ; n=14, P<0.01) (Fig. 1, A and B). Consistent with previous reports, there were no significant changes in mean SAP (control:  $104.0\pm3.6$ mmHg; MCT:  $102.5\pm3.4$  mmHg; CH:  $101.0\pm2.3$  mmHg) and heart rate (control:  $395\pm15$  beats/min; MCT:  $359\pm13$  beats/min; CH:  $361\pm7$  beats/min, Fig. 1, C and D) in the three groups of rats. Histological examination of lung sections of CH- and MCT-treated rats showed that the medial wall of the muscular small PAs (vessel outer diameter of <100µm) was thickened significantly (data not shown).

### TRPM8 expression was downregulated in PAs of CH- and MCT-induced PH rats

The relative mRNA expression of various TRPM subtypes was determined using quantitative real-time RT-PCR. TRPM1-8 transcripts were detected in endothelium-denuded PAs isolated from control rats. TRPM7 ( $0.018\pm0.005$ ) and TRPM8 ( $0.010\pm0.003$ ) transcripts were predominantly expressed, with a lower level of TRPM2, TRPM3, TRPM4 and TRPM6 transcripts, and almost no expression of TRPM1 and TRPM5 in control PAs. Expression of various TRPM channels was unchanged in PAs of CH- and MCT-treated rats, except for TRPM8. (Fig. 2). Compared to control, CH and MCT-treatment dramatically suppressed TRPM8 mRNA expression (control:  $0.010\pm0.003$ , n=9; MCT:  $0.005\pm0.0001$ , n=5, P<0.01; CH:  $0.0015\pm0.002$ , n=9, P<0.01). Consistent with mRNA results, TRPM8 protein level relative to GAPDH was also significantly decreased in both CH- and MCT-treated PAs (Fig. 2, B and C).

### TRPM8-mediated cation entry was reduced in PASMCs of CH and MCT-treated rats

Down-regulation of TRPM8 mRNA and protein in both models PASMCs were accompanied by a reduction in TRPM8-mediated cation entry determined by  $Mn^{2+}$ quenching of fura-2 fluorescence. Activation of TRPM8-mediated cation entry by exposing PASMCs to 300 µM menthol for 15 minutes caused a dramatic increase in the rate of  $Mn^{2+}$ induced quenching. The percent reduction in Fura-2 fluorescence measured 300 sec after application of 500 µM  $Mn^{2+}$  and max slope of  $Mn^{2+}$ -induced quenching was decreased from  $-32.6\pm3.4\%$  and  $-8.28\pm1.05\times10^{-3}$ /second (n=10, Fig. 3) in control to  $-12.6\pm1.5\%$  and

 $-4.03\pm0.47\times10^{-3}$ /second (n=12, P<0.01, Fig. 3) in hypoxic PASMCs, and  $-16.1\pm1.4\%$  and  $-3.91\pm0.72\times10^{-3}$ /second (n=10, P<0.01, Fig. 3) in MCT-treated PASMCs. These results clearly indicate that the expression and the functional activity of TRPM8 channel were both reduced in PASMCs of CH- or MCT-treated rats.

# Time-course of TRPM8 downregulation and the development of pulmonary hypertension in CH- and MCT-treated rats

The association between TRPM8 expression and PH was further examined by determining TRPM8 mRNA level, RVSP and RVMI in PAs at different time-points after MCT injection or CH exposure (Fig. 4). In PAs of MCT-treated rats, TRPM8 mRNA level decreased after 7 days post-injection and eventually declined to the minimal level after 2 weeks (0.06±0.02, n=6, P<0.01, Fig. 4, A). Significant increase in RVSP and RVMI was observed 3 to 5 days after MCT injection (36.5±3.1 mmHg, n=13, P<0.01, and 30.7±2.8%, n=11, P<0.05, Fig. 4, B and C), and they continued to increase to  $60.6\pm4.2$  mmHg (n=16, P<0.01, Fig. 4, B) and 59.9±4.9%, (n=14, P<0.01, Fig. 4, C), respectively, after 3 weeks. In contrast, TRPM8 mRNA level decreased significantly on the first day of CH exposure (normoxia: 1.00±0.2, CH: 0.08±0.01, n=6, P<0.01, Fig. 4, D), and the reduced level was maintained throughout the 3 week period. RVSP increased gradually, reaching a plateau on day 7 and sustained thereafter (Day 0: 27.4±1.0 mmHg, n=16; Day 7: 54.2±3.1 mmHg, n=8, P<0.01, Fig. 4, E). The time-course of increase in RVMI was similar to that of RVSP. Significant increase in RVMI was observed in day 7 (38.5±0.7%, P<0.01, Fig. 4, F) and it was followed by more gradual increase in the second and third week of CH exposure. Overall speaking, the downregulation of TRPM8 expression preceded the increase of RVSP and RVMI in CHtreated rats, but slightly lagged behind the changes in RVSP and RVMI in MCT-treated rats.

### Menthol-induced vasodilation was attenuated in CH- and MCT-induced PH

To further evaluate the effect of TRPM8 downregulation on pulmonary vascular function, the vasorelaxant effects of the TRPM8 agonist menthol was determined in phenylephrine pre-contracted endothelium denuded PA rings. Application of 1  $\mu$ M to 3 mM caused concentration-dependent relaxation in PAs of control, CH- or MCT-treated rats. Compared to control PAs, menthol induced vasodilation in phenylephrine pre-contracted PAs was significantly decreased in CH rats and in MCT rats with the maximal percentage relaxation of 68.2±2.2% (n=13, P<0.01) and 70.6±3.9% (n=14, P<0.01), respectively (Fig. 5). EC<sub>50</sub> of vasodilation was 97.8±14.7 $\mu$ M in control PAs (n=14), 81.5±14.4 $\mu$ M in PA of MCT-treated rats (n=13), and 185.0±41.8 $\mu$ M (n=14, P<0.05, Fig. 5) in PAs of CH rats. The attenuated vasorelaxing response of menthol in PAs of CH- and MCT-treated rats is consistent with of the downregulation of TRPM8 expression in CH- and MCT-induced PH rats.

## Discussion

In this study, we examined the changes in TRPM mRNA and protein expression as well as their functions in PAs of two widely used rat models of PH. The major findings of this study are: (1) among the various TRPM channels expressed in PAs, TRPM8 was the only channel subtype decreased significantly in both PH models; (2) the down-regulation in the expression of TRPM8 mRNA and protein in PASMCs of both models were accompanied by

a decline in menthol-induced cation entry; (3) significant down-regulation of TRPM8 mRNA was observed PAs of rats on the first day of CH-exposure and 1–2 weeks after MCT-injection, suggesting that the role of TRPM8 may be different in the development of PH in the two models; (4) the TRPM8 agonist menthol was capable of causing relaxation of de-endothelialized PAs and the maximal menthol-induced vasodilatation was reduced in the CH- and MCT-treated rats compared to those of control groups. These findings provide the molecular and pathophysiological evidence suggesting that TRPM8 may play a contributing role in the development of CH- and MCT-induced PH.

TRPM8 was originally identified as a prostate-specific gene. Its expression is androgenresponsive and upregulated in human prostate carcinoma [20], suggesting possible involvement in cell proliferation/metastasis. TRPM8 is also a menthol- and cold-sensitive ion channel in sensory neurons for the detection of cold temperature [21, 22]. It has also been shown to be expressed in skeletal and smooth muscle, lungs, bladder and urogenital tract, even though its functions in these tissues are not clear. Functional expression of TRPM8 has been demonstrated in rat intralobar pulmonary arteries and aortic smooth muscle [14]. Menthol elicits significant  $[Ca^{2+}]_i$  increase in PASMCs and aortic smooth muscle cells. The response could be abolished in the absence of extracellular Ca<sup>2+</sup> or in the presence of Ni<sup>2+</sup> but was unaffected by nifedipine, suggesting TRPM8 is a functional Ca<sup>2+</sup> influx channel in vascular myocytes [14]. TRPM8 mRNA and protein were also detected in rat tail, mesentery and femoral arteries [15].

Consistent with our previous studies [10, 14, 23], analysis of TRPM mRNA expression reveals multiple TRPM subtypes are coexpressed in rat PASMCs with TRPM7 and TRPM8 being most abundantly expressed in PAs of normal Sprague-Dawley rats. Among these TRPM subtypes, TRPM8 was the only channel downregulated in PAs of both CH- and MCT-treated rats. Our previous studies indicate that TRPC1, TRPC6 and TRPV4 are upregulated in PAs of CH rats [4, 10] and TRPC1 and TRPC4 expression are increased in PAs of MCT-induced PH rats [8], resulting in enhanced  $Ca^{2+}$  influx in PASMCs which contributes to the elevated pulmonary vascular tone and resting  $[Ca^{2+}]_i$ , as well as enhanced vascular reactivity in PH rats. TRPM8 therefore is the first TRP channel recognized by its down-regulation during PH [10], and the decrease in TRPM8 expression is reflected functionally by the significant reduction in non-selective cation entry detected by Mn<sup>2+</sup> quenching of Fura-2 fluorescence.

The association of TRPM8 down-regulation and the development of PH is evident in the time-courses of change in TRPM8 expression and the increase in RVSP and RVMI in the two different experimental models of PH. In the CH model, significant decrease in TRPM8 mRNA level was observed within one day of hypoxia exposure preceding the significant increase in RVSP and RVMI. This suggests that down-regulation of TRPM8 is an early response to hypoxia and maybe involved in the early development of PH. In contrast, the decrease in TRPM8 expression was considerably delayed in the MCT-model with significant difference observed seven days after MCT injection. This is in line with the slow progressive development of PH in the MCT-model, where MCT is first metabolized into the toxic metabolite MCT pyrrole which causes inflammation, pulmonary endothelial cell damage and apoptosis, medial thickening and adventitial remodeling [24, 25]. Since the

reduction in TRPM8 expression was observed after significant increase in RVSP and RVMI, it is likely consequential to the MCT-induced inflammation and cell injury and maybe related to the later stage of pulmonary vascular remodeling. The differences in the time-profiles of change in TRPM8 expression and RVSP in the two models of PH suggest that TRPM8 may participate in CH- and MCT-induced PH in dissimilar manners.

Information on the physiological functions and regulations of TRPM8 in vascular tissue is scant, and how TRPM8 down-regulation contributes to pulmonary hypertension is unclear. Our results showed that the TRPM8 agonist menthol induces concentration-dependent relaxation of phenylephrine-contracted PAs of control, CH- and MCT-treated rats. This is consistent with a previous report in systemic arteries showing that activation of TRPM8 with menthol or icilin causes a dual effect of small contraction in relaxed arteries but relaxation in vessels precontracted with KCl or phenylephrine [15]. The relaxation effect was mediated in part through  $Ca^{2+}$ -independent phospholipase A<sub>2</sub> activation. It has been shown that TRPM8 are localized on plasma membrane and sacroplasmic reticulum [26]. Activation of TRPM8 in plasma membrane may lead to increase in  $Ca^{2+}$  influx to cause vasoconstriction, but it may also activate large conductance Ca<sup>2+</sup>-activated potassium channels, leading to repolarization of cell membrane as shown in rat tail arterial myocytes [27]. Activation of TRPM8 on sarcoplasmic reticulum membrane, on the other hand, may lead to release/leak of SR Ca<sup>2+</sup> and Ca<sup>2+</sup> store depletion [28, 29]. In addition, Ca<sup>2+</sup> influx via TRPM8 may also activate Ca<sup>2+</sup>-activated Cl<sup>-</sup> channels in PASMCs to cause membrane-depolarization. The balance of these processes determines the manifestation of vasoconstriction or vasodilatation, depending on the vasomotor tone. However, the possible interactions between TRPM8, Ca<sup>2+</sup>-activated K<sup>+</sup> and Cl<sup>-</sup> channels have not been examined, even though the expression of these channels are known to be altered by PH [30–32]. Nevertheless the profound vasorelaxant effect of TRPM8 activated by menthol in PAs suggests that the channel may operate as a regulatory mechanism for counterbalancing vasoconstriction. In this regard, the down-regulation of TRPM8 expression may contribute to the enhanced pulmonary vascular reactivity in PH by reducing the vasodilatory influence against vasoconstriction. This is supported by the significant attenuation in the vasorelaxing effect of menthol observed in phenylephrine-precontracted PAs of CH- and MCT-treated rats.

Previous studies of TRPM8 in other cell systems showed that TRPM8 is regulated by phosphatidylinositol 4, 5-bisphosphate (PIP<sub>2</sub>). Direct activation of TRPM8 by PIP<sub>2</sub> has been demonstrated in purified reconstituted TRPM8 protein in planar lipid bilayers [33–34]. Activation of phospholipase C, as occurs during vasoconstriction induced by many Gq-coupled receptors, can inhibit TRPM8 activity through PIP<sub>2</sub> hydrolysis/depletion and protein kinase C dependent mechanisms [35–37]. In addition, TRPM8 is regulation by phospholipaseA2 [38–40]. Lysophospholipids, the end products of Ca<sup>2+</sup>-independent phospholipaseA2 have been shown to stimulate TRPM8 activity [38, 39]. In CH- and MCT-induced PH, PLC activities are known to be greatly enhanced due to the release of numerous vasoactive agonists and mitogenic factors, as well as the upregulation of many agonist receptors. Hence, it is likely that the channel activity in addition to protein expression of TRPM8 is also greatly reduced in PAs of PH rats.

In summary, we have demonstrated unequivocally that TRPM8 expression is downregulated in PAs of two etiologically different models of PH. The downregulation of TRPM8 expression is associated with reduced menthol-induced cation entry in PASMCs and menthol–induced vasorelaxation in agonist-precontracted PAs of PH rats. These alterations of TRPM8 expression and functions warrant further detail investigations, which will lead to novel information on the physiological and pathophysiological functions of TRP channels in pulmonary circulation.

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(*A*) Mean right ventricular pressure (RVSP), (*B*) right ventricular hypertrophy indexed by RVMI, (*C*) mean systemic arterial pressure (MSAP) and (*D*) heart rate (HR) in control, MCT-treated and CH-induced rats. The numbers of animals used in the control, MCT, and CH groups were 29, 18, and 14, respectively. \*\* P<0.01, difference between control and PH experimental model (MCT or CH); ## P<0.01, difference between MCT and CH groups.



# Fig. 2. Alterations in transient receptor potential melastatin (TRPM) expression in pulmonary arteries (PAs) of CH-exposured and MCT-treated rats

(*A*) quantitative real-time RT-PCR analysis of TRPM mRNA expression in endotheliumdenuded PAs of control (CON), MCT-treated and CH-exposured rats (n=9); (*B*) Western blots of TRPM8 protein in PAs of CON, MCT- and CH-induced PH rats (n=9); (*C*) normalized amount of TRPM8 proteins in PAs from 9 samples in CON, MCT and CH groups. \*, and \*\* denote significant differences of P<0.05 and P<0.01, respectively, between control rats and PH experimental models.



Fig. 3. Characterization of MT-induced cation entry by  $Mn^{2+}$  quenching of fura-2 in PASMCs (*A*) Representative tracings recorded in control, MCT- and CH-treated PASMCs. The averaged maximum quenching at 400s (*B*) and average maximum rate of quenching (*C*). 10 to 12 experiments from at least 3 different animals were performed for each group. \*\* indicates significant difference between control and PASMCs from CH- or MCT-treated rats.



## Fig. 4. Time-courses of change in TRPM8 mRNA expression, RVSP and RVMI in CH- and MCT-treated rats

(*A*) and (*D*) show the relative quantity of normalized TRPM8 mRNA expression in PA of rats isolated at various time-points after CH-exposure and MCT injection, respectively. (*B*) and (*E*): average values of RVSP measured at various time-points in the rat models. (*C*) and (*F*): average values of RVMI detected from CH and MCT rats at various time-points. The data were generated from  $6\sim17$  animals in each point, \* indicates P<0.05 and \*\* indicates P<0.01 when comparing with control.





Arrows indicate the application of menthol at various concentrations. 13 to 14 experiments from at least 3 different animals were performed for each group, \* indicates P<0.05, and \*\* indicates P<0.01 comparing to the control group.

## Table 1

Oligonucleotide sequences of the primers used for RT-PCR

Gene		Oligonucleotide Primers	Product(bp)
TRPM1	Sense	5'-CCTCCTGTGGACACCAAAGT-3'	
	Antisense	5'-AAGACAAGAGCGTCCAGCAT-3'	144
TRPM2	Sense	5'-CCATGCCACACATCAAACTC-3'	
	Antisense	5'-ATGATGGCCCAAATGAGAAG-3'	121
TRPM3	Sense	5'-ACCGGGTTGACTTTGTGAAG-3'	
	Antisense	5'-CAGATGAAGGGTGTTTGGT-3'	105
TRPM4	Sense	5'-ACCGGGTTGACTTTGTGAAG-3'	
	Antisense	5'-GCAGATGAAGGGTGTTTGGT-3'	141
TRPM5	Sense	5'-GGCACACCCATCTTACGACT-3'	
	Antisense	5'-ACTGTCTGGCTCCTGCAGAT-3'	126
TRPM6	Sense	5'-AAGCCATCCTGTCACCAAAG-3'	
	Antisense	5'-CAAAGAGGTAGACGGCTTGC-3'	180
TRPM7	Sense	5'-GGAAGGTGTGTGCAAAATGA-3'	
	Antisense	5'-GCTTCAGGTCCTTCACAAGC-3'	136
TRPM8	Sense	5'-GCTACGGACCAGCATTTCAT-3'	
	Antisense	5'-GCTTGTCAATGGGCTTCTT-3'	172
β-actin	Sense	5'-CCCATCTATGAGGGTTACGC-3'	
	Antisense	5'-TTTAATGTCACGCACGATTTC-3'	150