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Endogenous sulfur dioxide is a novel inhibitor of hypoxia-induced mast cell degranulation



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HIGHLIGHTS

- Endogenous SO₂/AAT pathway exists in mast cells (MCs).
- Endogenous SO₂ is a novel MC membrane stabilizer under hypoxic circumstance.
- MC-derived SO₂ upregulates cAMP level, thereby suppressing MC degranulation.

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G R A P H I C A L A B S T R A C T

Endogenous sulfur dioxide is a novel inhibitor of hypoxia-induced mast cell degranulation. Endogenous SO₂/AAT pathway exists in MCs. Mast cell-derived SO₂ upregulates cAMP level through activation of AC and inhibition of PDE, thereby suppressing degranulation of MCs.*Introduction*:



ABSTRACT

Introduction: Mast cell (MC) degranulation is an important step in the pathogenesis of inflammatory reactions and allergies; however, the mechanism of stabilizing MC membranes to reduce their degranulation is unclear.

Methods: SO₂ content in MC culture supernatant was measured by HPLC-FD. The protein and mRNA expressions of the key enzymes aspartate aminotransferase 1 (AAT1) and AAT2 and intracellular AAT activity were detected. The cAMP level in MCs was detected by immunofluorescence and ELISA. The release rate of MC degranulation marker β -hexosaminidase was measured. The expression of AAT1 and cAMP, the MC accumulation and degranulation in lung tissues were detected.

Objectives: To exam whether an endogenous sulfur dioxide (SO₂) pathway exists in MCs and if it serves as a novel endogenous MC stabilizer.

Results: We firstly show the existence of the endogenous SO_2/AAT pathway in MCs. Moreover, when AAT1 was knocked down in MCs, MC degranulation was significantly increased, and could be rescued by a SO_2 donor. Mechanistically, AAT1 knockdown decreased the cyclic adenosine monophosphate

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(cAMP) content in MCs, while SO₂ prevented this reduction in a dose-independent manner. Pretreatment with the cAMP-synthesizing agonist forskolin or the cAMP degradation inhibitor IBMX significantly blocked the increase in AAT1 knockdown-induced MC degranulation. Furthermore, in hypoxia-stimulated MCs, AAT1 protein expression and SO₂ production were markedly down regulated, and MC degranulation was activated, which were blunted by AAT1 overexpression. The cAMP synthesis inhibitor SQ22536 disrupted the suppressive effect of AAT1 overexpression on hypoxia-induced MC degranulation. In a hypoxic environment, mRNA and protein expression of AAT1 was significantly reduced in lung tissues of rats. Supplementation of SO₂ elevated the cAMP level and reduced perivascular MC accumulation and degranulation in lung tissues of rats exposed to a hypoxic environment *in vivo*.

Conclusion: SO_2 serves as an endogenous MC stabilizer via upregulating the cAMP pathway under hypoxic circumstance.

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Introduction

Mast cells (MCs) play important physiological and pathological roles in host defense, immune regulation, allergic reactions and chronic inflammation [1]. During MC degranulation, activated mast cells rapidly release particles stored in the cytoplasm into the extracellular environment in response to external stimuli [2]. The release of IgE-dependent degranulation and the release of proinflammatory mediators during an allergic reaction play important roles in host defense and nonallergic inflammatory diseases [3]. Studies have shown that the activation of MC degranulation is also an important step in the pathogenesis of nonallergic chronic diseases such as myocardial infarction [4], cardiomyopathy [5], cirrhosis [6,7] and other chronic diseases. Clarifying the mechanisms for MC degranulation has important implications in the life science and medical science fields.

The activation of MC degranulation is mainly induced by immune stimulation caused by antigens and nonimmune stimulation by drugs, which in turn causes the activation of downstream signaling pathways. This activation ultimately leads to the release of inflammatory mediators such as histamine, heparin and various cytokines stored in MCs. During allergic reactions, IgE binds to the IgE receptor and induces MC activation [8,9]. MC degranulation depends on the action of calcium ions. An increase in the cytosolic calcium ion concentration leads to changes in calmodulin (CaM) configuration, which in turn activates CaM protein kinase, and activated tubulin and myosin promote significant fusion and efflux of vesicle particles and cell membranes in MCs [10,11]. After MC activation, protein kinase C (PKC) is significantly activated and phosphorylates cell membrane proteins and cytoskeletal proteins, thereby triggering MC degranulation [12]; however, the mechanisms underlying MC degranulation and the responsible signaling pathways are unclear. Therefore, exploring endogenous factors that steadily regulate MC activation and stabilization, as well as the regulatory mechanisms, would be of great significance for the control of allergic diseases and inflammation.

Previous studies reported that endogenously generated gaseous signal molecules are widely present in mammals and involved in the fine and continuous regulation of disease development [13,14]. Sulfur dioxide (SO₂), a novel endogenous sulfur-containing gaseous signaling molecule catalyzed by aspartate aminotransferase (AAT) with the substrate of cysteine [15], significantly reduced the levels of inflammatory factors in endothelial cells [16]. Also, it was reported that SO₂ pretreatment reduced serum IL-1 and IL-6 levels in rats with acute lung injury induced by lipopolysaccharide [17]. The abovementioned reports suggest that endogenous SO₂ has the ability to control the inflammatory response. Previous findings demonstrated that the endogenous sulfur-containing gaseous signaling molecule hydrogen sulfide (H₂S) is produced in MCs [18] and that both H₂S and SO₂ are products of the same sulfur-containing amino acid metabolic family

[19,20]. Therefore, we hypothesized that a potential endogenous SO_2 production pathway exists in MCs. In the case of such a pathway, whether endogenous SO_2 regulates allergic or inflammationinduced degranulation of MCs is an important mechanism to address.

After the activation of human MCs, active substances such as βhexosaminidase and histamine, release and participate in the inflammatory reaction, and the cyclic adenosine monophosphate (cAMP)/protein kinase A (PKA)/phosphodiesterases (PDEs) pathway impacts the activation of human MCs [21]. The intracellular cAMP was found to inhibit excessive inflammatory or allergic reactions. cAMP activates cAMP-dependent PKA to inhibit IgE/antigendependent signaling activation and calcium influx, thereby inhibiting MC degranulation [21,22]. Increased intracellular cAMP levels caused by stimuli, such as adenosine and beta-adrenergic receptors, were shown to inhibit MC degranulation [23]. Previous studies on smooth muscle cell proliferation showed that endogenous SO₂ activates cAMP/PKA and thereby inhibits the activation of Raf/MEK/ERK and cell proliferation [24]. Whether endogenous SO₂ controls the degranulation of MCs and the underlying mechanism remains unclear. Therefore, we investigated the existence of an endogenous SO₂/AAT pathway in MCs, its possible role in MC degranulation and its potential mechanisms.

Materials and methods

Reagents

MCs, vascular smooth muscle cells (VSMCs) and vascular endothelial cells (VECs) were acquired from the China Infrastructure of Cell Line Resources Center. Sodium bisulfite and sodium sulfite (NaHSO₃/Na₂SO₃, freshly mixed at a mole ratio 1:3, pH 7.4) was used as the SO₂ donor. Lentivirus carrying AAT1 shRNA, AAT2 shRNA, scramble shRNA and AAT1 cDNA, and vehicle lentivirus were constructed by Cyagen (Guangzhou, China). The AAT activity detection kit was purchased from Jiancheng (Nanjing, China). The cAMP ELISA kit was from Neweast(PA, USA). The Lglutamine, streptomycin penicillin and fetal bovine serum were purchased from Gibco (USA). SYBR Green was purchased from Thermofisher (USA). The Oligo(dT)15 primer and reverse transcription kit were purchased from Promega (USA). Toluidine blue dye was purchased from Solarbio (Beijing, China).

Cell culture and treatment

The MCs and VECs used in the experiments were humanderived, and the VSMCs were rat-derived. The MCs were cultured in Iscove's Modified Dulbecco's Medium (IMDM), the VECs were cultured in F12/Dulbecco's modified Eagle's medium (F12/DMEM) and the VSMCs were cultured in DMEM. To prepare complete culture medium for MCs, IMDM was supplemented with the fetal bovine serum (10%), streptomycin penicillin (1%) and L-glutamine (2 mM). The cell synchronization culture medium IMDM was the same as the complete culture medium except for the addition of fetal bovine serum. The MCs were cultured in a moist environment at 37 °C and 5% CO₂.

The concentration of forskolin [25] and SQ22536 used in the experiment was 30 μ mol/L. The concentration of IBMX was 500 μ mol/L [26]. The oxygen concentration in the hypoxic treatment group was set at 1%.

The plasma SO₂ concentration in rats was approximately 0– 30 μ mol/L [27]. A previous study showed that the SO₂ concentration in rat vascular tissues was approximately 127 μ M [28]. Therefore, the SO₂ donor in the present study was used at concentrations of 50, 100 and 200 μ mol/L for dose-dependent experiments and at 100 μ mol/L for the remaining cellular experiments.

Lentivirus transfection

The lentivirus carrying AAT1 or AAT2 shRNA (Cyagen, China) was transfected into the MCs to construct AAT1 or AAT2 knockdown MCs according to manufacturer's instructions. Briefly, the MCs were seeded in 25 cm² culture flasks and infected with lentiviral AAT1 or AAT2 shRNA at a multiplicity of infection of 10 when the cell density was about 60%-70%. After a 24-hour transfection, the medium was replaced with fresh complete culture medium and incubated at 37 °C overnight. Then, the infected cells were screened with puromycin (4 µg/mL) for 1 week to acquire the stably transfected AAT1 or AAT2 shRNA was used to infect the MCs as the control according to the same protocol. Similarly, overexpression of AAT1 in MCs was achieved by transfecting with lentivirus containing AAT1 cDNA and vehicle lentivirus was used as the control.

Animal model

Twenty-four Wistar male rats (150–160 g) were randomly divided into normoxic, hypoxic and hypoxic + SO_2 groups (n = 6 each group). The rats in hypoxic group and hypoxic + SO_2 group breathed in the normobaric hypoxic air $(10\% O_2)$ in a hypoxia chamber (Biospherix, USA) for 3 weeks and 6 h every day, while the rats in the normoxic group breathed in the room air $(21\% O_2)$ for the same period. Moreover, rats in the hypoxic + SO_2 group were injected with NaHSO₃/Na₂SO₃ mixture (0.18 mmol and 0.54 mmol per kilogram body weight) before hypoxia exposure each day. The rats of the normoxic and hypoxic groups were injected with the same volume of physiological saline. The dose of SO₂ donor was chosen based on the previous finding that the administration of this dosage of NaHSO₃/Na₂SO₃ mixture could significantly restore the hypoxia-inhibited pulmonary SO₂ content and alleviate pulmonary hypertension and pulmonary vascular remodeling in hypoxic rats [29]. The experimental rat hypoxic model was established according to the previous literatures [29-32]. In the abovementioned studies [29–32], the rats exposed to hypoxia for 21 days showed a significant pulmonary hypertension and pulmonary vascular inflammation.

All rats were euthanized after 21 days of hypoxic exposure and the lung tissues were collected. One side of the lung lobe was removed and fixed with 10% formalin. Then, paraffin-embedded lung tissues were sectioned at a thickness of 5 μ m for toluidine blue staining and immunohistochemistry. The remaining lung tissue was rapidly frozen in liquid nitrogen and stored at – 80 °C for subsequent examinations.

Determination of aspartate aminotransferase (AAT) 1 and AAT2 mRNA by real-time RT-PCR

The AAT1 and AAT2 mRNA expression in MCs were measured by RT-PCR. The following primers were used:

AATT-Inditially, J-GTCAGAATCCGGTCAGCCATT-J (R),
5'-GTGGCCAGCACCCTCTCTAA-3' (F);
AAT2-human, 5'-CGCAGGCATGCAGAAGAA-3' (R),
5'-GTGGCCAGCACCCTCTCTAA-3' (F);
GAPDH-human, 5'-AGCCTTCTCCATGGTGGTGAAGAC-3' (R),
5'-CGGAGTCAACGGATTTGGTCGTAT-3' (F).
AAT1-rat, 5'-GCCATTGTCTTCACGTTTCCTT-3' (R),
5'-CCAGGGAGCTCGGATCGT-3' (F);
AAT2-rat, 5'-GTTTCCCCAGGATGGTTTGG-3' (R),
5'-GAGGGTCGGAGCCAGCTT-3' (F);
Actin-rat, 5'-TATCGTCATCCATGGCGAACT-3' (R),
5'-ACCCGCGAGTACAACCTTCTT-3' (F).

Total RNA was extracted from the MCs and lung tissues using Invitrogen TRIzol reagent. A total of 2 µg of RNA was used for complementary deoxyribonucleic acid (cDNA) synthesis by reverse transcription with an oligo(dT)15 primer. cDNA synthesis conditions included a 25 µl of reaction system for denaturation at 70 °C for 5 min followed by 42 °C for 60 min. The real-time PCR reaction conditions were set at 95 °C pre-denaturation for 10 min, 95 °C denaturation for 15 s, 60 °C annealing for 1 min, and 72 °C polymerization extension for 30 s, with a total of 40 cycles. The relative mRNA expression of AAT1 or AAT2 was analyzed using the formula $2^{-\Delta\Delta CT}$ method. GAPDH or β-actin mRNA was used as internal reference, respectively.

Determination of SO₂ concentration by HPLC-FD

The SO₂ content in the cell supernatant and rat lung tissues was detected by high performance liquid chromatography with fluorescence detection (HPLC-FD) [22]. This experiment was quantified by the standard sodium sulfite. Briefly, sodium borohydride (70 μ l, 0.212 mol/L) was added to the 100 μ l of sample or standard. The mixtures were then incubated at 37 °C for 30 min. Next, mBrB (5 μ l, 70 mmol/L) was added to the above mixtures, which were then incubated at 42 °C for 10 min. Perchloric acid (40 μ l, 1.5 mol/L) was added to the mixture. The samples were centrifuged (12,400 × g) at 23 °C for 10 min to remove the protein precipitates. Tris-HCl (10 μ l, 2 mol/L, pH 3.0) was used to neutralize the mixture. Finally, the mixture was collected and then analyzed by HPLC.

Determination of AAT1 and AAT2 protein expression by Western blot analysis

The MCs were collected by centrifuging the cell culture medium, and washed twice with PBS. The cells were lysed in RIPA lysis buffer (50 mM Tris-Cl (pH 7.4), 150 mM NaCl, 1 mM EDTA, 1% NP40, 0.25% sodium deoxycholate and protease inhibitor cocktail) for 20 min in an ice bath, vortexed, ultrasonicated and centrifuged. The supernatant was collected as the protein lysate. The lung tissues were ground into a homogenate according to the mass volume ratio 1:10 $(mg/\mu L)$ with RIPA lysis buffer, and then the homogenate was centrifuged for 10 min at 4 °C to separate the supernatant. The protein lysate was separated on a 10% SDS polyacrylamide gel, and then transferred to a membrane. Following ponceau staining and using the reference protein marker, the target protein band was labeled. The membrane was blocked and then incubated with primary antibody. The primary antibodies used for the experiment were as follows: anti-AAT1 from Sigma, 1:1000; anti-AAT2 from Sigma, 1:1000; and anti-GAPDH from KangCheng, 1:4000. The following day, the membrane was removed from 4 °C and incubated with the following secondary antibodies: goat antimouse IgG, 1:5000; rabbit anti-goat IgG, 1:2000; and goat antirabbit IgG, 1:2000. Exposure was performed using the Alpha Imager 3400 gel image acquisition system. The strip gray value was analyzed by the Alpha Ease FS software and corrected by the gray value of the internal reference GAPDH strip.

AAT activity detected by colorimetric assay

The colorimetric method was used to test AAT protein activity in MCs [27]. The MCs were collected and lysed in 80 μ l of 0.01 M PBS by ultrasound under ice water bath conditions. AAT activity was detected following the manufacturer instructions. The substrate solution was added into the 96-well plate and briefly placed at 37 °C. A total of 20 μ l of each sample was added to the corresponding well, and the plate was placed at 37 °C for 30 min. The sodium hydroxide solution (200 μ l, 0.4 mol/L) was added and gently mixed into each well. The plate was incubated for 15 min at room temperature. The optical density (OD) value was detected at the absorption wavelength of 510 nm.

The β -hexosaminidase release rate detected by colorimetric assay

The release of β-hexosaminidase was used as a marker for MC degranulation. β-acetylglucosaminidase is an acid hydrolase in the cell lysosome. The substrate is hydrolyzed by βacetylglucosaminidase to release free p-nitrophenol. The reaction was terminated by adding an alkaline solution, in which the pnitrophenol was stained. The MCs were separated from the cell culture medium and resuspended in 4 °C PBS. The cells were sonicated and centrifuged, and the supernatant was collected for analysis. A total of 100 µl of the sample (cell culture supernatant or cell solution) was mixed with 500 µl of substrate buffer. After thoroughly mixing, the samples were incubated for 15 min at 37 °C. After stopping the reaction, the absorbance was detected at 400 nm to calculate the β -acetylglucosaminidase activity. Each liter of the sample was incubated with the substrate for 1 min at 37 °C. The enzyme activity unit was 1 µmol p-nitrophenol formed by hydrolysis. In this experiment, 0.6 mmol/L p-nitrophenol was used as the standard.

Immunofluorescence detection of cAMP expression in mast cells

A confocal laser scanning microscope was used to acquire the immunofluorescence images. The MCs were gently rinsed and fixed. The anti-cAMP antibody (1:50, CST, USA) was added to the cells then the secondary antibody (1:200, Thermo, USA) was added to conjugate to the primary antibody. Finally, the cells were observed under a confocal microscope [33].

The level of cAMP detected by enzyme-linked immunosorbent assay (ELISA)

The level of cAMP in the MCs was measured using an ELISA kit. The number of MCs was counted using a cytometer. The MCs were then centrifuged to remove the cell culture medium, and lysed by the addition of HCl (100 μ l, 0.1 M). The standard sample had a known concentration of cAMP. A total of 100 μ l of the standard and samples were added to a microplate that was coated with polyclonal goat anti-mouse antibody. A total of 50 μ l of cAMP-conjugated horseradish peroxidase and 50 μ l of anti-cAMP antibody were then added to the microplate, which was then shaken for 2 h. After washing, the substrate was added to develop color for 30 min. Immediately after the reaction was stopped, an OD value at the absorption wavelength of 450 nm was measured with a microplate reader, and the cAMP concentration of each sample was calculated.

Toluidine blue staining

Toluidine blue O is a commonly used synthetic basic dye. The cations in toluidine blue have a dyeing effect. The acidic substances of tissue cells are combined with the cations to be dyed, and the nucleus can be stained blue. Heparin and histamine in the MC cytoplasm can be stained purplish red when encountered with toluidine blue, and as such was used to detect MCs. The rat lung tissue sections were dewaxed and incubated with toluidine blue dye solution for 10 min. The excess staining solution was washed away and the color was separated with 95% alcohol for 5 min. Then, images were captured with a Leica inverted microscope. The number of toluidine blue-positive cells in the rat lung tissue sections was counted at $400 \times$ high power field (HFP).

Detection of cAMP and AAT1 expression by immunohistochemistry

The lung tissue slides were dewaxed by dimethylbenzene and then rehydrated by graded ethanol. After rinsing with PBS, the slides were incubated with 3% H₂O₂ for 10 min at room temperature in a humidified chamber. The slides were washed with PBS and incubated with goat serum for 30 min at 37 °C. The slide was incubated with the primary antibody (anti-cAMP, 1:100; anti-AAT1, 1:100) overnight at 4 °C. The slide was then rinsed with PBS and incubated with the secondary antibody for 1 h at 37 °C. 3,3-diaminobenzidine (DAB) was added to develop color, and the sections were stained with hematoxylin for 5 s. The slides were dehydrated by graded ethanol and made transparent in dimethylbenzene. Images were captured with a Leica inverted microscope.

Statistical analysis

The SPSS (20.0) software was used for statistical analysis. The data were presented as the mean \pm standard deviation. The difference between 2 groups was analyzed by an independent Student's *t*-test. A one-way analysis of variance (ANOVA) was used for the comparison of the differences among multiple groups following the Bonferroni post hoc test if the data were normally distributed, while the Dunnett T3 post hoc test followed ANOVA if the data were not normally distributed. Pearson's correlation analysis was used to identify the dose-dependence effect. P-values were set to 0.05, which was considered statistically significant.

Results

An endogenous SO₂/AAT pathway is present in mast cells

In this study, VECs and VSMCs were used as positive controls to detect an endogenous SO₂ pathway in MCs. The results demonstrated that endogenous SO₂ production was detected in the MC culture supernatant, and the SO₂ content was 2.75 times and 1.62 times higher than that of VECs and VSMCs, respectively (Fig. 1A). In mammals, the essential enzymes of endogenous SO₂ production are AAT1 and AAT2. With VECs and VSMCs as positive controls, the real time RT-PCR results showed that AAT1 and AAT2 mRNA were also present in MCs (Fig. 1B, C). Moreover, western blot analysis showed the presence of AAT1 (Fig. 1D) and AAT2 (Fig. 1E) proteins in MCs. The AAT activity was 10.77 ± 2.03 U/L in MCs, 20.73 ± 2.36 U/L in VECs, and 43.55 ± 18.86 U/L in VSMCs, respectively (Fig. 1F).

AAT1 knockdown promoted mast cell degranulation

To investigate the impact of endogenous SO_2 on MC stabilization, MCs were infected with AAT1 knockdown lentivirus. Com-



Fig. 1. The presence of endogenous SO₂/AAT pathway in MCs. (A) The SO₂ content in the supernatant of MCs, VECs and VSMCs detected by HPLC-FD; (B) Relative AAT1 mRNA levels in MC, VEC and VSMC detected by real time RT-PCR; (C) Relative AAT2 mRNA levels in MC, VEC and VSMC detected by real time RT-PCR; (C) Relative AAT2 protein expression in MC, VEC and VSMC detected by western blot; (E) Relative AAT2 protein expression in MC, VEC and VSMC detected by western blot; (E) Relative AAT2 protein expression in MC, VEC and VSMC detected by western blot; (F) AAT activity in MC, VEC and VSMC detected by colorimetric analysis. *P < 0.05, **P < 0.01. The data are presented as the mean \pm SD. The results in Fig. 1B-1C were normalized with those in MCs. All the research was carried out for three times independently.

pared with the scramble group, the SO₂ content in the supernatant and the cAMP protein expression in the AAT1-knockdown MCs were significantly reduced (Fig. 2A and B), while MC degranulation, represented by β -hexosaminidase release rate, increased by approximately 2.18 fold (Fig. 2C). Exogenous SO₂ donor supplementation successfully rescued the effect of AAT1 knockdown, demonstrated by the upregulation of cAMP protein expression and reduction of MC degranulation (Fig. 2B, C). To further investigate whether the effects of SO₂ are dose-dependent, we examined the β -hexosaminidase release rate (Fig. 2D) and cAMP (Fig. 2E) level after supplementing with different dose of SO2 (50-200 μ M). The data demonstrate that SO₂ upregulated cAMP and downregulated MC degranulation in a dose-dependent manner. Moreover, AAT2 knockdown (Figure Supplementary Fig. S1. A) had no significant effect on the SO₂ content in the supernatant (Fig. S1. B), the cAMP level in the MCs (Fig. S1. C) and MC degranulation (Fig. S1. D) compared with the scramble group. These results suggest that SO₂ produced by AAT1 mainly upregulates the cAMP level and inhibits mast cell degranulation.

cAMP level elevation blocked SO₂ reduction-driven MC degranulation

To further investigate whether cAMP mediates the inhibitory effect of endogenous SO_2 on MC degranulation, AAT1-depleted MCs were treated with the PDE inhibitor IBMX or adenylate cyclase (AC) activator forskolin. Compared with the scramble group, both the SO_2 content and cAMP expression in the AAT1-depleted MCs were significantly decreased (Fig. 3A, B), and MC degranulation was significantly increased (Fig. 3C); however, the treatment of MCs with the AC activator forskolin or the PDE inhibitor IBMX to

elevate cAMP content successfully abolished AAT1 knockdowninduced MC degranulation (Fig. 3B, C).

SO_2 protected against hypoxia-induced mast cell degranulation in vitro

Hypoxic challenge downregulated AAT1 protein expression but did not affect AAT2 protein expression in the MCs *in vitro* (Fig. 4A, B). Accordingly, the SO₂ content in the MC supernatant was significantly reduced (Fig. 4C). Furthermore, AAT1 overexpression markedly blunted the effect of hypoxia on the intracellular cAMP level (Fig. 4D) and MC degranulation *in vitro* (Fig. 4E). Importantly, treatment with SQ22536 to inhibit cAMP synthesis abolished AAT1 overexpression-inhibited MC degranulation (Fig. 4E).

SO_2 inhibited the hypoxia-driven downregulation of cAMP and increase in MC degranulation in vivo

Chronic hypoxic challenge downregulated the protein and mRNA expression of AAT1 (Fig. 5A, B, C) and the level of cAMP (Fig. 5D), but increased MC accumulation and degranulation in the rat lung tissues *in vivo* (Fig. 5E). The mRNA expression of AAT2 was also downregulated (Figure Supplementary Fig. S2. A), but hypoxia exposure did not affect AAT2 protein expression (Fig. S2. B, C) in the rat lung tissues *in vivo*. Importantly, exogenous SO₂ donor supplementation successfully rescued the downregulation of cAMP protein expression *in vivo* (Fig. 5D). Moreover, the supplementation with a SO₂ donor decreased the hypoxia-induced perivascular MC accumulation and degranulation in the lung tissues of rats exposed to a hypoxic environment *in vivo* (Fig. 5E).



Fig. 2. Knockdown of AAT1 promoted the degranulation of MCs. (A) The SO₂ content in the MC supernatant detected by HPLC-FD. NaHSO₃/Na₂SO₃ mixture (1:3 M/M) at a dose of 100 μ M was used as a SO₂ donor; (B) cAMP level in MCs detected by immunofluorescence. NaHSO₃/Na₂SO₃ mixture (1:3 M/M) at a dose of 100 μ M was used as a SO₂ donor; (C) The release rate of β -hexosaminidase in MCs detected by colorimetric analysis. NaHSO₃/Na₂SO₃ mixture (1:3 M/M) at a dose of 100 μ M was used as a SO₂ donor; (D) The release rate of β -hexosaminidase in MCs treated with different concentration of SO₂ donor NaHSO₃/Na₂SO₃ mixture (50–200 μ M) detected by colorimetric analysis; (E) The cAMP content in MCs treated with different concentration of SO₂ donor NaHSO₃/Na₂SO₃ mixture (50–200 μ M) detected by colorimetric analysis; (E) The cAMP content in MCs treated with different concentration of SO₂ donor NaHSO₃/Na₂SO₃ mixture (50–200 μ M) detected by ELISA. *P < 0.05 and the data are presented as the mean ± SD. All the research was carried out for three times independently.

Discussion

SO₂ is regarded as a toxic gas and environmental pollutant; however, SO₂ can be endogenously generated in mammals, dissolved in plasma and derived into sulfite and generate bisulfite to maintain the stability of the internal environment [27,28,34,35,36]. In mammals, sulfur-containing amino acids, such as L-cysteine, are converted to L-cysteine sulfinate by the catalysis of cysteine dioxygenase, which is converted to β -thionylpyruvate through AAT1/2 and eventually spontaneously decomposes into pyruvate and SO₂ [27,34,36]. In this study, the presence of endogenous SO₂ was first detected in the cell culture supernatant of MCs. The mRNA level and the protein expression levels of AAT1 and AAT2, which are key enzymes for the production of endogenous SO_2 , as well as AAT activity were detected in the MCs. Therefore, we first suggest that there might exist an endogenous SO_2 /AAT pathway in MCs.

In the present study, VSMCs and ECs, two kinds of endogenous SO_2 source cells [27,28], were used as positive reference to demonstrate the existence of endogenous SO_2 pathway in MCs. Interestingly, the AAT expression and activity in the certain cell seemed not parallel to SO_2 content released from the cell. For example, VSMC showed the highest AAT1/2 mRNA levels, AAT2 protein and AAT activity, and second highest AAT1 protein level. However, the levels of SO_2 in VSMC was lower than that of MC. We speculated that there might be other impactors involved in the regula-



Fig. 3. The elevation of cAMP levels blocked AAT1 knockdown-induced MC degranulation. (A) The SO₂ content in MC supernatant detected by HPLC-FD; (B) The cAMP level in MCs detected by immunofluorescence; (C) The release rate of β -hexosaminidase in MCs detected by colorimetric analysis. *P < 0.05 and the data are presented as the mean ± SD. All the research was carried out for three times independently.

tion of endogenous SO_2 content such as non-enzymatic SO_2 generation, SO_2 consumption and clearance according to a comprehensive review about SO_2 metabolism and SO_2 donor [37], which merits to be investigated in the future.

Numerous studies have demonstrated that MCs are abundantly present in exposed areas of the body (such as airways, skin and intestines) and are usually located near blood vessels and nerves. In this location, MCs can take on a sentinel role in early host defense and play an important role in many inflammatory diseases such as allergic rhinitis, conjunctivitis, asthma and arthritis [38,39]. The local microenvironment can directly affect the maturity, phenotype and function of MCs. These mediators released from MC enable MC to initially respond in harmful situations and respond to the altered environment by communicating with cells involved in physiological and immune responses. After myocardial ischemia-reperfusion injury, MCs activate and degranulate. Renin is released, which in turn activates the local renin-angiotensin system and ultimately leads to ventricular fibrillation [40]. MCs are also involved in the pathogenesis of allergic asthma. Activated MCs release histamine and activate smooth muscular contraction, bronchial secretion and airway mucosal edema [41]. However, the endogenous regulatory mechanism underlying degranulation following MC activation is not fully understood. Here, we showed that a spontaneous MC degranulation accompanied with a decrease in SO₂ content in AAT1 knockdown MCs, which was concurrently rescued by SO₂ donor. Especially, SO₂ donor rescued the AAT1 knockdown-induced MC degranulation in a dose-dependent manner. On the contrary, AAT2 knockdown did not affect the SO₂ content in the supernatant and MC degranulation. Therefore, these findings suggest that the endogenous SO₂/AAT1 pathway might have an important inhibitory effect on MC degranulation.

Subsequently, investigating the possible mechanisms by which endogenous SO₂ inhibits MC degranulation is an important issue. Previous studies found that the interventions of elevating intracellular cAMP level including cAMP-synthesis agonists Forskolin and isoprenaline, and cAMP-degradation inhibitor IBMX correlated well with the inhibition of histamine, leukotriene C4, and PGD2 release in MCs [42], suggesting that cAMP is an important controller for MC degranulation. It was a coincidence that VSMCderived SO₂ inhibited cell proliferation via increasing the cAMP level and activating PKA pathway [24]. The abovementioned findings give us a hint that cAMP level might bridge endogenous SO₂ and MC degranulation. As we expected, the data showed that SO₂ donor recovered the AAT1 knockdown-reduced cAMP level in a dose-dependent manner. Furthermore, the AC activator Forskolin and the PDE inhibitor IBMX both increased cellular cAMP levels and simultaneously blocked AAT1 knockdown-induced excessive degranulation of MCs, suggesting that endogenous SO₂/AAT1 pathway inhibits the degranulation of MCs by increasing intracellular cAMP levels under physiological conditions.

The potential role of endogenous SO₂ in regulating MC degranulation under pathophysiologic conditions, as well as the underlying mechanisms, was unclear. Previous studies showed that hypoxia significantly stimulates MC degranulation, causing the release of cytokines such as angiogenic factors to induce retinal neovascularization, resulting in eye diseases and even blindness [43]. MCs are infiltrated in hypoxic microenvironments caused by tumors, and MC activation promotes angiogenesis and tumor invasion [44]. Hypoxia is a significant feature of inflammatory tissue, including the lung of asthmatic patients, which is closely related to mast cell infiltration and degranulation [45]. In the pathology of fatal asthma, inflammatory cells infiltrate in the air-



Fig. 4. AAT1 overexpression significantly inhibited the hypoxia-induced degranulation of MCs *in vitro*. (A-B) Western blot analysis of protein expression of AAT1 (A) and AAT2 (B) in MCs in normoxic and hypoxic groups *in vitro*; (C) The SO₂ content in supernatant of MCs was detected by HPLC *in vitro*; (D) The cAMP content in MCs detected by ELISA *in vitro*; (E) Colorimetric analysis of the release rate of β -hexosaminidase in MCs *in vitro*; *P < 0.05, **P < 0.01 and data are presented as the mean ± SD. All the research was carried out for three times independently.

way wall and surrounding parenchyma. MCs can be located in the airway wall and are activated after being stimulated by hypoxia. Inflammatory mediators are then released by MCs, which then recruit more inflammatory cells, ultimately causing intense bronchospasm and sudden death [46]. Moreover, MC degranulation during hypoxic conditions plays an important role in the pathogenesis of pulmonary vascular remodeling [47]. Therefore, hypoxia stimulation was used as a model to activate MC degranulation in the present study. The data showed that AAT1 protein expression in the MCs was decreased while AAT2 protein expression remained unchanged under a hypoxic environment. Simultaneously, the decrease in the supernatant SO₂ content, the reduction of cAMP level and the cell degranulation occurred in the hypoxic MCs. However, MCs with AAT1 overexpression no longer responded to hypoxic stimulation representing by the fact that there were no differences in SO₂ content, cAMP level and cell degranulation in the AAT1-overexpressing MCs between normoxic and hypoxic groups. While, an AC inhibitor, SQ22536, restored the hypoxiadriven cell degranulation in the AAT1-overexpressing MCs. The abovementioned findings provided in vitro experiment evidence for the hypothesis that the downregulation of endogenous SO₂/ AAT1 and subsequent decreased cAMP might be involved in the hypoxia-driven MC degranulation.

Furthermore, in a rat model of chronic hypoxic stimulation, we found that hypoxic exposure significantly inhibited AAT1 protein expression but not AAT2 protein expression in the lung tissue, pulmonary vessel and perivascular tissue. Also, the SO₂ content in the lung tissue was decreased with perivascular MC accumulation and degranulation and reduction of cAMP expression in the rats with hypoxic exposure. In accordance with the data obtained from

in vitro experiments, the supplementation of SO₂ donor recovered SO₂ content in the lung tissue and blocked the hypoxia-inhibited cAMP expression and hypoxia-induced perivascular MC accumulation and degranulation, which further supported the relationship among endogenous SO₂/AAT1 pathway, cAMP level and MC mediator release. Combined with the significance of MC degranulation in the vascular inflammation and the findings that SO₂ donor inhibited hypoxic pulmonary vascular inflammation reported in our previous study [29], the abovementioned results in the present study imply that hypoxia-driven endogenous SO₂/AAT1 downregulation might contribute to MC degranulation and be involved in inflammation of the pulmonary blood vessels *in vivo*.

Regarding the effect of SO₂ on the MC degranulation, Melendez et al. reported an opposite result and described that 0.5 mM and 5 mM of Na₂SO₃ induced cardiac MC degranulation due to oxidative stress [48]. We speculated that reasons contributing to the discrepancy might include the followings: (1) The aim of the two studies is different. Melendez et al. focused on the toxic effect of SO_2 as an air pollutant on the MC, while we aimed to explore the physiological and pathophysiological significance of endogenous SO₂ generated from an AAT1-catalyzed transamination reaction in the MC. The difference might affect the experimental design and results of the two studies. (2) The experimental design is different. Based on the findings of the existence of endogenous SO₂/ AAT pathway in the MCs, we designed to explore the effect of endogenous SO₂ on the MCs using AAT1/2 knockdown intervention and SO_2 donor rescue. (3) The dose and composition of reagent is different. In the study of Melendez et al, 0.5 mM and 5 mM of Na₂-SO₃ was used, while we mainly used 100 µmol/L NaHSO₃/Na₂SO₃ (1:3 M/M) as a SO₂ donor. As described in the material and meth-



Fig. 5. SO_2 donor inhibited the hypoxia-driven downregulation of cAMP and increase in MC degranulation *in vivo*. (A) Western blot analysis of AAT1 protein expression in rat lung tissue; (B) Real time RT-PCR analysis of AAT1 mRNA levels in rat lung tissue; (C) Immunohistochemistry analysis of AAT1 protein expression in rat lung tissue; (D) Immunohistochemistry analysis of cAMP protein expression in rat lung tissue; (E) Toluidine blue staining was used to detect MC accumulation and degranulation (red arrows) around rat pulmonary vascular tissue. The rats in the hypoxic + SO₂ group were injected with NaHSO₃/Na₂SO₃ mixture (0.18 mmol and 0.54 mmol per kilogram body weight) before hypoxia exposure each day. The rats of the normoxic and hypoxic groups were injected with the same volume of physiological saline. **P < 0.01 and data are presented as the mean ± SD, n = 6 each group.

ods part [27–28], the dose and composition used in our study is close to physiological status of SO_2 . The comparison and analysis of the contrary results from the two studies may benefit the researchers to pay attention on the dose range of SO_2 and avoid the toxic effect of SO_2 at high dose in the future.

SO₂ gas and NaHSO₃/Na₂SO₃ (1:3 M/M) are the two common widely used SO₂ donors. SO₂ gas dissolved in water forms a hydrated SO₂ complex (SO₂·H₂O) that dissociates twice to generate HSO₃⁻ and SO₃²⁻. The formation of SO₂ from a NaHSO₃/Na₂SO₃ (1:3 M/M) solution depends on the complex kinetics of the dynamic equilibrium between gaseous SO₂ and HSO₃⁻/SO₃²⁻ [37]. Therefore, whether a SO₂ gas solution or NaHSO₃/Na₂SO₃ (1:3 M/M) solution is administrated to cells or animals, they may exist in the form of

 SO_2 and HSO_3^-/SO_3^{2-} . However, there exist the differences in the amount of SO_2 releases, release rate, delivery rate to tissues or cells, action efficiency and clearance rate between the SO_2 gas solution and NaHSO₃/Na₂SO₃ mixtures [37]. For instance, a previous study demonstrated that gaseous SO_2 might have a stronger vasodilator effect compared with a NaHSO₃/Na₂SO₃ mixture [28]. In future studies, we would compare the difference in the usage of SO_2 donor between the SO_2 gas solution and NaHSO₃/Na₂SO₃ mixture in different animal models and cell types.

SO₂, as an endogenous new gas signaling molecule, has a significant inhibitory effect on MC degranulation induced by hypoxia. The findings reported in this work may provide novel ideas to aid in the clinical prevention and treatment of many MC-related

diseases, such as allergic rhinitis, asthma, conjunctivitis, psoriasis, mastocytosis, various cancers, cirrhosis and cardiovascular diseases (including idiopathic cardiomyopathy, atherosclerosis, myocarditis, ischemic heart disease and pulmonary hypertension). In addition, this study helps to further clarify the biological and pharmacological roles of SO₂ at physiological concentrations, and may attract additional researchers to develop clinically applicable SO₂ donors.

Conclusions

In summary, this work demonstrates that an endogenous SO₂/ AAT pathway exists in MCs and importantly, that SO₂ acts as a novel MC stabilizer. Endogenous SO₂ deficiency can promote MC degranulation. Conversely, supplementation with SO₂ prevents hypoxia-induced MC degranulation. SO₂ inhibits MC degranulation by enhancing the cAMP pathway. The present data provide a theoretical basis for exploring the mechanisms for MC-mediated pathological processes, such as inflammatory reactions and allergies. The data presented here also provide basis for novel potential treatment strategies for pulmonary vascular inflammation.

Compliance with Ethics Requirements

All institutional and National Guidelines for the care and use of animals (fisheries) were followed. This study was approved by the Animals Care and Use Committee of Peking University First Hospital, Beijing, China.

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Declaration of Competing Interest

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

Author contributions

L. Zhang, H. Jin, Y. Song, J. Du and Y. Huang designed the experiments, performed the experiments, acquired the data, analyzed the data and wrote the manuscript; Y. Wang and Y. Sun performed the experiments; C. Tang designed the experiments; and S. Y. Chen revised the manuscript.

Appendix A. Supplementary material

Supplementary data to this article can be found online at https://doi.org/10.1016/j.jare.2020.08.017.

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