Diallyl sulfide protects against dilated cardiomyopathy via inhibition of oxidative stress and apoptosis in mice

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Abstract. Cytochrome P450 family 2 subfamily E member 1 (CYP2E1) is a member of the cytochrome P450 enzyme family and catalyzes the metabolism of various substrates. CYP2E1 is upregulated in multiple heart diseases and causes damage mainly via the production of reactive oxygen species (ROS). In mice, increased CYP2E1 expression induces cardiac myocyte apoptosis, and knockdown of endogenous CYP2E1 can attenuate the pathological development of dilated cardiomyopathy (DCM). Nevertheless, targeted inhibition of CYP2E1 via the administration of drugs for the treatment of DCM remains elusive. Therefore, the present study aimed to investigate whether diallyl sulfide (DAS), a competitive inhibitor of CYP2E1, can be used to inhibit the development of the pathological process of DCM and identify its possible mechanism. Here, cTnTR141W transgenic mice, which developed typical DCM phenotypes, were used. Following treatment with DAS for 6 weeks, echocardiography, histological analysis and molecular marker detection were conducted to investigate the DAS-induced improvement on myocardial function and morphology. Biochemical analysis, western blotting and TUNEL assays were used to detected ROS production and myocyte apoptosis. It was found that DAS improved the typical

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Professor Lianfeng Zhang, Key Laboratory of Human Disease Comparative Medicine, National Health Commission of China (NHC), Institute of Laboratory Animal Science, Peking Union Medical College, Chinese Academy of Medical Sciences, Building 5, Panjiayuan Nanli, Chaoyang, Beijing 100021, P.R. China E-mail: zhanglf@cnilas.org DCM phenotypes, including chamber dilation, wall thinning, fibrosis, poor myofibril organization and decreased ventricular blood ejection, as determined using echocardiographic and histopathological analyses. Furthermore, the regulatory mechanisms, including inhibition both of the oxidative stress levels and the mitochondria-dependent apoptosis pathways, were involved in the effects of DAS. In particular, DAS showed advantages in terms of improved chamber dilation and dysfunction in model mice, and the improvement occurred in the early stage of the treatment compared with enalaprilat, an angiotensin-converting enzyme inhibitor that has been widely used in the clinical treatment of DCM and HF. The current results demonstrated that DAS could protect against DCM via inhibition of oxidative stress and apoptosis. These findings also suggest that inhibition of CYP2E1 may be a valuable therapeutic strategy to control the development of heart diseases, especially those associated with CYP2E1 upregulation. Moreover, the development of DAS analogues with lower cytotoxicity and metabolic rate for CYP2E1 may be beneficial.

Introduction

Cytochrome P450 (CYP) family 2 subfamily E member 1 (CYP2E1) is a member of the CYP family of heme proteins and is expressed in the liver, brain and heart in humans and animals (1-4). The CYP2E1 protein is highly expressed on the mitochondria, Golgi apparatus, endoplasmic reticulum and plasma membrane (5,6). CYP2E1 is a key monooxygenase enzyme involved in the metabolism of several endogenous substrates, as well as hundreds of xenobiotics, obtained from diet and environmental contamination (2,4-6). Compared with other CYP family members, CYP2E1 possesses remarkably high NADPH oxidase activity. Thus, it can catalyze the reaction of NADPH and O₂ to generate reactive oxygen species (ROS) (7,8). CYP2E1 is the major enzyme for the production of cellular and mitochondrial ROS and reactive nitrogen species (RNS), which induce tissue damage mainly by damaging cellular and mitochondrial macromolecules, including mitochondrial DNA (9-11).

The expression level of CYP2E1 is upregulated in a wide variety of pathological states and diseases, including obesity, diabetes mellitus, alcoholic liver disease, heart diseases,

Key words: cytochrome P450 family 2 subfamily E member 1, inhibitor of CYP2E1, diallyl sulfide, dilated cardiomyopathy, oxidative stress

inflammation, Parkinson's disease and cancer (12-17). In view of the expression characteristic of CYP2E1, it has been used as a therapeutic target in drug discovery and already has been applied in preliminarily studies (18). Diallyl sulfide (DAS) is a competitive inhibitor of CYP2E1, which is isolated from garlic (18,19). It has been reported that DAS treatment can recover ethanol-induced pathological changes in the liver by inhibiting CYP2E1-mediated alcohol metabolism and subsequent oxidative stress (20,21). Furthermore, DAS treatment can prevent myocardial systolic dysfunction induced by chronic ethanol exposure (22). Studies of the effects of DAS suggested that CYP2E1 inhibition may have potential effects on the development of diseases characterized with high CYP2E1 expression.

Dilated cardiomyopathy (DCM), characterized by the dilation of the heart chambers and impaired systolic functions, is the most frequent cause of heart transplantation and it is the third most common reason for heart failure (HF) (23). Accumulating evidence has indicated the important role of oxidative stress in the pathophysiology of cardiac remodeling and HF (24,25). Furthermore, it has been shown that CYP2E1 is upregulated in multiple cardiovascular diseases, including DCM, hypertrophic cardiomyopathy (HCM) and ischemia (4,5,26-32). Our previous study revealed that the phenotype of CYP2E1 overexpressing-transgenic mice was similar to that of DCM, and knockdown of endogenous CYP2E1 expression markedly prevented the development of DCM in mice (4,27). In addition, since CYP2E1 is a major source of cellular and mitochondrial ROS/RNS, inhibition of CYP2E1 may improve the pathological development of DCM. The aforementioned studies suggest that inhibition of CYP2E1 may improve the pathological development of DCM and application of CYP2E1 inhibitors may become an effective therapeutic approach; however, to the best of our knowledge, no relevant studies have been reported so far.

Therefore, the present study aimed to investigate whether DAS can be used to inhibit the development of the pathological process of DCM and examined the possible underlying mechanism. For this purpose, cTnT^{R141W} transgenic mice, a mouse model with typical DCM phenotypes, such as decreased survival rate, dilated chambers, thin walls and cardiac dysfunction, which were successfully established in our previous studies (4,26,27), were used. Furthermore, the possible inhibitory mechanism of DAS on development of DCM was also investigated in the present study.

Materials and methods

Animals. The a-MHC-cTnT^{R141W} transgenic mice were generated in the Key Laboratory of Human Disease Comparative Medicine and maintained in a C57BL/6J genetic background in the Key Laboratory of Human Disease Comparative Medicine (Ministry of Health, Peking Union Medical College, Beijing, China), and exhibited DCM phenotypic characteristics consistent with those reported previously (4). A total of 58 4-month-old mice were used, including 40 cTnT^{R141W} transgenic mice and 18 wild-type mice. In total, there were 34 males weighing 28-32 g, and 26 females weighing 22-24 g. All mice were bred in an American Association for Accreditation of Laboratory Animal Care-accredited facility, housed at $21\pm2^{\circ}$ C with $50\pm5\%$ relative humidity under a 12-h day/night cycle and had free access to food and drinking water. The procedures were approved by the Animal Care and Use Committee at the Institute of Laboratory Animal Science, Peking Union Medical College (approval no. ZLF18004).

Groups and treatment. The 4-month-old male and female cTnT^{R141W} transgenic mice were randomly assigned to treatment groups. DAS (cat. no. A35801; Sigma-Aldrich; Merck KGaA) was diluted in corn oil to a final concentration of 80 mg/ml. In the treatment groups, cTnT^{R141W} transgenic mice were administered DAS at a dose of 200 (n=12) or 400 mg/kg (n=10) via intraperitoneal injection three times weekly for 6 weeks. A group of cTnT^{R141W} transgenic mice (n=9) and non-transgenic littermates (NTG; n=8) were treated with corn oil (0.1 ml per mouse) as the placebo control and wild-type normal control, respectively. As a commercially available drug control, a group of cTnT^{R141W} transgenic mice were treated with enalaprilat (cat. no. H20010498; Changzhou Pharmaceutical Factory) via intraperitoneal injection three times weekly for 6 weeks, an angiotensin-converting enzyme (ACE) inhibitor that has been widely used in the clinical treatment of DCM and HF (33,34), at a dose of 0.76 mg/kg (n=9). The dose of DAS was selected based on previous reports (35,36). Therefore, there were total five groups in this study, including NTG normal control, placebo model control, DAS treatment at high dose (400 mg/kg), DAS treatment at low dose (200 mg/kg) and enalaprilat. After the last echocardiograph examination, the animals were euthanasia via cervical dislocation.

Echocardiography. Heart function and structure were analyzed via echocardiography once every 2 weeks during treatment. Briefly, the mice were lightly anesthetized via an intraperitoneal injection of 216 mg/kg tribromoethanol and then subjected to 2-D guided M-mode echocardiography with a 30-MHz transducer for echocardiographic examination (Vevo770; FUJIFILM VisualSonics Inc.). Measurements of left ventricular (LV) fractional shortening (LVFS), LV ejection fraction (LVEF), LV diameter at end systole (LVESD), LV diameter at end diastole (LVEDD), LV posterior wall at end systole (LVPWD), LV anterior wall at end systole (LVAWD) were based on the analysis of \geq 10 separate cardiac cycles.

Histological analysis. For light microscopy, heart tissues were first fixed in 4% formaldehyde at room temperature for 24 h, embedded in paraffin and cut into 4- μ m thick sections (4). Heart tissue sections were imaged after H&E and Masson staining. For H&E staining, the sections were stained using a Hematoxylin and Eosin Staining kit (cat. no. G1121; Beijing Solarbio Science & Technology Co., Ltd.). Briefly, paraffin-embedded sections were dewaxed in xylene two times for 10 min each time at room temperature. The sections were rehydrated in a series of ethanol (100, 95, 85 and 75%) for 3 min per gradient, and for 2 min in distilled water. Subsequently, the sections were stained with hematoxylin stain for 3 min at room temperature and washed with distilled water to remove floating colors. The differentiation solution was differentiated for 3 min and washed with distilled water

twice, for 2 min each time. The sections were put into eosin dye solution for 2 min, then washed with distilled water for 2-3 sec at room temperature. Then, the sections were proceeded to the dehydrated step through soaking in a series of concentrations of alcohol at room temperature (for 2-3 sec at 75, 85 and 95%, for 1 min at 100%). The sections were then incubated with xylene twice at room temperature (for 1 min each time), followed by sealing in neutral gum and finally observed under the microscope.

For Masson trichrome staining, the sections were incubated in celestine blue solution (cat. no. G1345; Beijing Solarbio Science & Technology Co., Ltd.) for 5 min and briefly washed with H₂O. Then, sections were incubated in hemalun solution for 5 min and in fuchsine acid/ponceau xylidine (0.5% fuchsine acid, 1.5% ponceau xylidine, 1,75% glacial acetic acid) for 5 min. Subsequent, sections were briefly washed with H₂O and incubated in phosphomolybdic acid (1%) for 10 min and in aniline blue solution (2.5% anilin blue, 2.5% glacial acetic acid) for 5 min. Next, they were briefly washed with H₂O and incubated in acetic acid (1%) for 1 min. Finally, the sections were briefly incubated in an ascending ethanol series followed by xylol, before they were embedded.

Transmission electron microscopy (TEM). Heart tissues (~2 mm³) from the free wall of the left ventricle were fixed in 2.5% glutaraldehyde (preparation for 100 ml solution: 50 ml of 0.2 M phosphate buffer, 10 ml of 25% glutaraldehyde and 40 ml double distilled water) at 4°C for 12 h (4). After washing three times with phosphate solution (pH=7.2), the samples were fixed in 1% osmium acid at 4°C for 2 h. Dehydration was carried out sequentially with four graded concentrations of ethanol between 50-100% at room temperature. After epichlorohydrin replacement and epoxy embedding at room temperature, tissue sections were cut on an ultramicrotome. The semi-thin sections (thickness, 900 nm) were further stained with an equal mixture of 4% uranyl acetate and acetone for 30 sec at room temperature, and then in lead citrate for 2 min. The final ultra-thin sections (90 nm) were analyzed under a JEM-1230 transmission electron microscope (JEOL Ltd.).

Measurement of ROS. After treatment, the mice were sacrificed and total lysates were prepared as previously reported (4). H_2O_2 , malondialdehyde (MDA; lipid peroxidation) and glutathione (GSH) in heart tissues were measured with relevant assay kits (H_2O_2 Assay kit, cat. no. ab102500; MDA Assay kit, cat. no. ab118970; GSH/GSSG Ratio Detection Assay kit, cat. no. ab138881; all from Abcam) following the manufacturer's procedures.

For the H_2O_2 assay, 10 mg tissue was first washed in cold PBS and homogenized in 500 μ l assay buffer with a Dounce homogenizer sitting on ice. The sample was centrifuged 13,000 x g for 5 min at 4°C at top speed using a microcentrifuge to remove any insoluble material. Finally, the content of H_2O_2 in the heart tissue was measured with a microplate reader at excitation/emission (Ex/Em)=535/587 nm.

For the lipid peroxidation assay, 10 mg tissue was first washed in cold PBS and homogenized in 303 μ l lysis solution with a Dounce homogenizer sitting on ice. The sample was centrifuged at 13,000 x g for 10 min at 4°C to remove insoluble material. Then, thiobarbituric acid (TBA) reagent was added

into the supernatant to generate MDA-TBA adduct. Finally, the absorbance was measured immediately at Ex/Em=532/553 nm.

For the glutathione assay, 10 mg tissue was first washed in cold PBS, then was resuspended in 400 μ l cold lysis buffer. The samples were homogenized and centrifuged at 13,000 x g for 15 min at 4°C to remove any insoluble material. Next, 50 μ l GSH assay mixture was added into each GSH standard or collected sample supernatant. Finally, the absorbance was measured at Ex/Em=490/520 nm with a fluorescence microplate reader.

TUNEL assay. For apoptotic cell staining, paraffin sections of heart tissues were stained using an ApopTag Plus Peroxidase In Situ Apoptosis kit (cat. no. S7101; MilliporeSigma) following the manufacturer's procedures. In briefly, heart tissues were first fixed in 4% formaldehyde at room temperature for 24 h, embedded in paraffin and cut into $4-\mu m$ thick sections. The tissue slides were rehydrated and were treated with 25 μ g/ml proteinase K at 37°C for 8 min at room temperature. After washing and incubation with equilibration buffer for 5 min at room temperature, Tdt was diluted at 1:3.9 with reaction buffer and added to heart sections for 1 h at 37°C. After applying stop solution for 10 min at room temperature, the samples were incubated with anti-digoxigenin peroxidase conjugate at 37°C for 30 min. Slides were first treated with a 1:20 dilution of diaminobenzidine (3.3'-diaminobenzidine) substrate for 1 min at room temperature, then counterstained with hematoxylin (cat. no. G1121; Beijing Solarbio Science & Technology Co., Ltd.) for 30 sec at room temperature. The sections were proceeded to the dehydrated step through soaking in a series of concentrations of alcohol at room temperature (for 2-3 sec at 75, 85 and 95%, for 1 min at 100%). Then, the sections were incubated with xylene twice at room temperature (for 1 min each time), followed by sealing in neutral gum and finally observed under the microscope. In total, nine visual fields were randomly selected in each group to observe the apoptosis under a light microscope (BX53; Olympus Corporation; magnification, x400). The results are expressed as the percentage of apoptotic cells among the total cell population.

Reverse transcription (RT)-PCR. Total RNA was isolated from heart tissues using TRIzol[®] reagent (cat. no. 15596018; Invitrogen; Thermo Fisher Scientific, Inc.) and were used to synthesize cDNA with a RT kit (cat. no. RR820A; Takara Bio, Inc.) following the manufacturer's procedures. mRNA expression levels of procollagen type III α 1 (Col3 α 1) was detected via RT-PCR using GAPDH for normalization under standard conditions. The primers were as follows: Col3 α 1 forward, 5'-CTCAAGAGCGGAGAATACTGG-3' and reverse, 5'-CAA TGTCATAGGGTGCGATA-3'; and GAPDH forward, 5'-CAA GGTCATCCATGACAACTTTG-3' and reverse, 5'-GTCCAC CACCCTGTTGCTGTAG-3'.

Western blotting. Heart tissues were homogenized and extracted using lysis buffer [tissue protein extraction reagent (cat. no. 78510); protease inhibitor cocktail (cat. no. 87785); phosphatase inhibitor cocktail (cat. no. 78420); PMSF (100 μ M; cat. no. 36978; all from Thermo Fisher Scientific, Inc.); 100:1:1:1 ratio for configuration], followed by fractionation using a Mitochondrial/Cytosol Fractionation kit (cat. no. ab65320; Abcam) to obtain the cytosolic and mitochondrial fractions. An enhanced BCA protein assay kit (cat. no. P0010; Beyotime Institute of Biotechnology) was used to quantify the protein concentration. The total lysates, cytosolic and mitochondrial fractions (50 μ g) were separated via 15% SDS-PAGE and transferred to a nitrocellulose membrane (Immobilon NC; MilliporeSigma). Following blocking in 5% fat-free milk in TBS-0.1% Tween-20 (TBST) for 1 h at room temperature, the membranes were incubated at 4°C overnight with primary antibodies targeted against: CYP2E1 (1:500; cat. no. ab28146), cardiac troponin T (1:500; cTnT; cat. no. ab8295), cytochrome c (1:500; cat. no. ab13575), procaspase 3 (1:1,000; cat. no. ab13847), procaspase 9 (1:1,000; cat. no. ab47537; all from Abcam), cleaved (active) caspase 3 (1:500; cat. no. 9507S) and cleaved caspase 9 (1:500; cat. no. 9664S; both from Cell Signaling Technology, Inc.), anti-\beta-tubulin (1:1,000; cat. no. ab21058) and anti-voltage dependent anion channel 1 (1:1,000; cat. no. ab14734; both from Abcam) were used. Following washing with TBST, the membranes were incubated with a HRP-conjugated secondary antibody (1:10,000; anti-rabbit IgG, cat. no. ZB-2301 or anti-mouse IgG, cat. no. ZB-2305; ZSGB-BIO, Inc.) at room temperature for 1 h. Primary antibody binding was visualized using a chemiluminescent detection system (Western Blotting Luminal Reagent; Santa Cruz Biotechnology, Inc.) and analyzed using the densitometry function of Quantity One software (version 3.0; Bio-Rad Laboratories, Inc.).

Cell lines and culture. H9c2 cells (National Laboratory Cell Resource Sharing Service Platform) were cultured in DMEM (Thermo Fisher Scientific, Inc.), supplemented with 10% FBS (Thermo Fisher Scientific, Inc.) and 100 U/ml penicillin-100 μ g/ml streptomycin (Invitrogen; Thermo Fisher Scientific, Inc.) at 37°C with 5% CO₂. Cells were treated with DAS (12 h; 100 μ mol/l; cat. no. A35801; Sigma-Aldrich; Merck KGaA) or isoprenaline hydrochloride (ISO; 12 h; 50 μ mol/l; cat. no. I5627; Sigma-Aldrich; Merck KGaA) at 37°C with 5% CO₂ as required.

Calcein-AM/PI double staining. Cell viability was evaluated using calcein-AM/PI double staining following the manufacturer's instruction (cat. no. C542; Dojindo Laboratories, Inc.). Briefly, H9c2 cells were seeded in 96-well plates in triplicate at a density of 1×10^4 cells/100 µl/well with 100 μ l culture medium and cultured for 24 h. The cells were treated with ISO (50 μ mol/l) or DAS (100 μ mol/l) for 12 h at 37°C with 5% CO₂ and divided into four groups, including control, ISO, control + DAS and ISO + DAS groups. The cells were then washed twice with PBS and the working solution (PBS: calcein-AM: PI=1,000:1:1) and incubated for 15 min at 37°C. Live cells with yellow-green fluorescence and dead cells with red fluorescence (Leica Microsystems GmbH) were observed using a fluorescence microscope (magnification, x100) at 490±10 nm and 545 nm excitation wavelengths. In total, three images of each well were captured for a total of three replicate wells for counting.

Statistical analysis. All experiments were performed ≥3 times. Statistical analyses were performed using SPSS software (version 19.0; IBM Corp.). The data were analyzed using

two-tailed unpaired t-tests and one-way ANOVA followed by Tukey's post hoc analysis. Data are presented as the mean \pm SD. P<0.05 was considered to indicate a statistically significant difference.

Results

DAS improves cardiac morphology breakage and dysfunction in cTnT^{R141W} DCM model mice. Compared with the NTG group, the cTnT^{R141W} transgenic mice showed typical DCM phenotypes, such as dilated chambers, thin walls and cardiac dysfunction. The DCM phenotypes were evidenced by increases in LVESD and LVEDD, and decreases in LVPWS, LVPWD, LVAWS, LVAWD, LVEF and LVFS (Fig. 1A-I; Tables SI-SIII).

Firstly, after 2 weeks of treatment, the early stage of drug intervention, it was found that improvement was already observed in both DAS groups, and echocardiographic parameters in both DAS groups showed an improved inhibition trend compared with that of the enalaprilat group. For example, LVESD decreased by 26.5 and 25.9% in the DAS high and low dose group, respectively, compared with the placebo model control group (P<0.001; Fig. 1B), while LVESD decreased by 18.9% in the enalaprilat group compared with the placebo model control group (P<0.01; Fig. 1B). Moreover, LVEF increased by 57.8 and 43.1% in the DAS high and low dose group, respectively, compared with the placebo model control group (P<0.001; Fig. 1H), while LVEF increased by 35.2% in the enalaprilat group compared with the placebo model control group (P<0.05; Fig. 1H).

After 6 weeks of treatment, LVESD decreased by 22.4% in the DAS high dose group (P<0.001; Fig. 1B), while it decreased by 11.1% in the enalaprilat group (P<0.05; Fig. 1B) compared with the placebo model control group. There was a significant increase in LVEF in the DAS high dose group compared with the placebo model control group (P<0.001; Fig. 1H), while there was no difference between the enalaprilat group and the placebo model control group (P>0.05; Fig. 1H). Furthermore, there was a significant increase in LVAWD in the enalaprilat group compared with the DAS high dose group (P<0.001; Fig. 1G). LVPWS increased in both the DAS high dose group and the enalaprilat group compared with the placebo model group (P<0.05; Fig. 1D), while it showed no significant difference between these two groups (P>0.05).

In summary, both DAS groups and the enalaprilat group exhibited a significant improvement on DCM phenotypes, as evidenced by changes in cardiac function, chamber size and wall thickness. While each factor has its advantages, DAS exhibited a stronger effect on control of chamber dilation and dysfunction, and the enalaprilat group was more effective at increasing wall thickness. Furthermore, this improvement on morphology and function of model mice occurred earlier in both DAS groups compared with that of the enalaprilat group.

DAS inhibits cardiac pathological development in cTnT^{R141W} DCM model mice. After drug treatment, hearts from all five groups were sampled for gross morphology and pathological examinations.

First of all, the protein expression level of CYP2E1 was determined via western blotting after treatment with



Figure 1. Echocardiographic analysis of cardiac morphology and function. Mice in the five groups were analyzed after 6 weeks of treatment: NTG (wild-type control), cTnT^{R141W} (placebo control), treatment with 400 mg/kg DAS, treatment with 200 mg/kg DAS and treatment with enalaprilat (commercially available drug control). (A) Representative M-mode echocardiographic images of the LV long axis. (B) LVESD. (C) LVEDD. (D) LVPWS. (E) LVPWD. (F) LVAWS. (G) LVAWD. (H) LVEF. (I) LVFS. n=7. *P<0.05, ***P<0.001 vs. NTG group; #P<0.05, ##P<0.001 vs. cTnT^{R141W} group; \$P<0.05 vs. 400 mg/kg DAS group. NTG, non-transgenic; LV, left ventricular; LVESD, left ventricular end-systole diameter; LVEDD, left ventricular end-diastole diameter; LVPWS, left ventricular anterior wall at end systole; LVPWD, left ventricular posterior wall at end diastole; LVAWS, left ventricular anterior wall at end systole; LVEF, left ventricular ejection fraction; LVFS, left ventricular fractional shortening; DAS, diallyl sulfide.

different doses of DAS. Compared with the placebo model control group, the protein expression level of CYP2E1

in 200 and 400 mg/kg DAS groups decreased by 35.5% (n=3; P<0.01) and 51.8% (n=3; P<0.001; Fig. 2A and B), respectively.

Moreover, CYP2E1 protein expression decreased by 35.9% in the enalaprilat group (n=3; P<0.01; Fig. 2A and B). The expression level of cTnT protein was also analyzed. The mutant form of cTnT is the human mutant form that was introduced by the transgenic method, which is the cTnT band at the higher molecular weight in Fig. 2A, and the location of endogenous cTnT proteins in mouse myocardium was slightly lower than that of exogenous mutants. The expression level of the cTnT showed no difference between the three therapeutic groups and the placebo model group (Fig. 2A and C).

The increased heart to body weight ratio of $\text{cTnT}^{\text{R141W}}$ mice was reversed by DAS treatment to almost the normal level both in the 400 and 200 mg/kg group (P<0.05; Fig. 2D). DAS treatment significantly improved the chamber dilation, wall thinning and myocyte disarray in $\text{cTnT}^{\text{R141W}}$ DCM model mice, as determined via H&E staining (Fig. 2E and F). Through ultrastructure observation using TEM, it was found that DAS treatment improved the poor myofibril organization, including diffusion, damage and lysis, in $\text{cTnT}^{\text{R141W}}$ mice (Fig. 2H). Quantitative analysis of the Masson staining and the mRNA expression level of Col3 α 1 showed that collagen deposition in the interstitial space of $\text{cTnT}^{\text{R141W}}$ mice was significantly reduced in both DAS and enalaprilat groups compared with the placebo model control group (Fig. 2G and I-K).

Both DAS and enalaprilat treatment exhibited a favorable therapeutic effect on improving of the destroyed microstructure and ultrastructure of the myocardium in DCM model mice. Moreover, the DAS group exhibited a greater advantage in reducing index of heart to body weight ratio.

To verify the ameliorative effect of DAS on the pathological phenotype of cardiomyopathy *in vitro*. Cell experiments were performed, and it was found that DAS could significantly inhibit the increased expression of CYP2E1 induced by ISO treatment in H9c2 cells, as well as significantly improve the cell death induced by ISO (Fig. S1), which were all consistent with the results *in vivo*.

DAS reduces oxidative stress level in $CTnT^{R141W}$ DCM model mice. CYP2E1 catalyzes the production of ROS even in the absence of substrate, leading to oxidative stress (37). Therefore, H₂O₂, MDA and GSH levels were measured as indicators of oxidative stress in all five groups.

 H_2O_2 and MDA levels were significantly decreased (P<0.001; Fig. 3A and B), while GSH levels were significantly increased (P<0.01 and P<0.001; Fig. 3C) in both 400 and 200 mg/kg DAS treatment groups compared with those of the placebo model group. Inhibition of the expression of CYP2E1 by DAS resulted in a reversion of the levels of H_2O_2 , MDA and GSH in the myocardium to almost the normal levels in both the 400 and 200 mg/kg DAS groups. By contrast, no significant differences in those three indicators were observed in the enalaprilat group compared with the placebo model group (P<0.05; Fig. 3A-C).

Therefore, inhibition of oxidative stress level is one of the possible underlying mechanisms of DAS, rather than enalaprilat, that is involved in the protection against the pathological process of DCM.

DAS inhibits mitochondrial pathways of apoptosis in *cTnT*^{R141W} DCM model mice. The induction of CYP2E1 causes cytochrome c release and activation of the mitochondrial apoptosis pathway in the heart in $cTnT^{R141W}$ DCM model mice (4).

In both DAS groups, the increased release of cytochrome *c* in cTnT^{R141W} DCM model mice almost reached those of normal levels (P<0.05; Fig. 4A and B). The activation of caspase 9 was decreased by 66.6% (P<0.001) and 66.9% (P<0.001; Fig. 4A and C) in the 400 and 200 mg/kg DAS groups, respectively, compared with the placebo model group. The activation of caspase 3 was decreased by 65.1% (P<0.001) and 62.7% (P<0.001; Fig. 4A and D) in the 400 and 200 mg/kg DAS groups, respectively, compared with the placebo model group.

The release of cytochrome *c* from the mitochondria to the cytoplasm triggers the apoptosis of cardiac myocytes in $cTnT^{R141W}$ model mice (38,39). It was determined that there were 69.3 and 64.7% decreases in proportion of apoptotic cells in the 400 (P<0.05) and 200 mg/kg groups (P<0.05; Fig. 4E and F), respectively, compared with the placebo model group. Furthermore, DAS and enalaprilat exhibited no difference in apoptosis inhibition.

Therefore, inhibition of the mitochondrial apoptosis pathway is one of the possible mechanisms of DAS that is involved in the protection against the pathological process of DCM.

Discussion

CYP2E1 is an inducible gene that is upregulated under multiple conditions, such as fasting and nutrition intake, as well as in a wide variety of pathophysiological states (2,12-17). In addition to metabolizing endogenous substrates and xenobiotics, CYP2E1 is the main source of cellular ROS, and its NADPH oxidase activity is higher than that of other CYP family members (9-11).

Our previous studies revealed that the expression level of CYP2E1 was increased in multiple mouse models of heart disease, including those of DCM and HCM, and the possible mechanism of upregulation of CYP2E1 expression were examined, in which it was found that Myc was upregulated under pathological stimulus and binds to the CYP2E1 promoter to activate its transcription (26,27). Furthermore, it was determined that the increase of CYP2E1 expression induced cardiac myocyte apoptosis via mitochondrial pathways, and knockdown of endogenous of CYP2E1 expression using small interfering RNA could attenuate the pathological development of DCM in CTnT^{R141W} mice (4).

Therefore, in the present study, DAS, a selective inhibitor of CYP2E1, was used to treat cTnT^{R141W} DCM model mice, and to observe whether DAS can be used to inhibit the development of the pathological process of DCM and its possible mechanism.

DAS is an organosulfur compound derived from the metabolism of allicin and has anti-cancer properties (40). DAS inhibits the activity of CYP2E1 and thus has attracted attention as a potential therapeutic or prophylactic agent (41,42). DAS treatment has been applied in several diseases. For instance, DAS treatment can attenuate the pathogenesis of diseases associated with CYP2E1 upregulation in animal models, such as alcoholic liver disease, nonalcoholic steatohepatitis, diabetes and alcoholic cardiomyopathy (18,43-46). At present, to the best of our knowledge, the use of DAS in treating DCM animal models to observe its therapeutic effects had not been previously reported.



Figure 2. Pathological histology observation. (A) CYP2E1 and cTnT expression in the heart tissues of mice in the NTG, cTnT^{R141W}, 400, 200 mg/kg DAS and enalaprilat groups was detected via western blotting. (B) CYP2E1 and (C) cTnT were semi-quantitatively analyzed, using GAPDH for normalization (n=3). (D) Ratio of HW to BW (n=6). (E) H&E staining patterns of whole-heart longitudinal sections. (F) Magnification of H&E-stained sections of the left ventricle (magnification, x400; scale bar, 20 μ m). (G) Magnification of Masson's trichrome-stained left ventricle sections (magnification, x400, scale bar, 20 μ m). (G) Magnification of Masson's trichrome-stained left ventricle sections (magnification, x400, scale bar, 20 μ m). (J) Quantitative analysis of Masson staining (n=3). (J) Col3 α 1 expression was detected and (K) analyzed (n=3). *P<0.05, ***P<0.001 vs. NTG group; #P<0.05, ##P<0.01, ###P<0.001 vs. cTnT^{R141W} mice. cTnT, cardiac troponin T; HW, heart weight; BW, body weight; Col3 α 1, procollagen type III α 1; NTG, non-transgenic; DAS, diallyl sulfide; CYP2E1, cytochrome P450 family 2 subfamily E member 1.

DAS, as an active ingredient in garlic, it is less toxic than other garlic sulfides, such as diallyl disulfide and diallyl trisulfide. DAS can be rapidly metabolized into diallyl sulfoxide (DASO), diallyl sulfone (DASO2) and allyl mercaptan. Allyl mercaptan is the decomposition product of DAS, and DASO and DASO2 are sulfur oxidation products of DAS.



Figure 3. Determination of oxidative stress levels. Levels of (A) H₂O₂, (B) MDA and (C) GSH in the heart tissues of mice in the NTG, cTnT^{R141W}, 400, 200 mg/kg DAS and enalaprilat groups were determined using colorimetric assays (n=3). *P<0.05, ***P<0.001 vs. NTG group; #*P<0.01, ##*P<0.001 vs. cTnT^{R141W} mice. MDA, malondialdehyde; GSH, glutathione; NTG, non-transgenic; DAS, diallyl sulfide.

DASO and DASO2 are the main metabolites among DAS metabolites, and their formation is mediated by the CYP2E1 enzyme. DAS is both a selective inhibitor and a substrate of CYP2E1. Since sulfur is more nucleophilic than DAS terminal carbon, sulfur strongly binds to heme of CYP2E1, leading to competitive inhibition of CYP2E1 by DAS or metabolism into DASO (18,47,48). DAS inhibits the metabolism of P-450 2E1 substrates via competitive inhibition mechanisms and by inactivating P-450 2E1 via a suicide-inhibitory action of DASO2 (18,47,48).

cTnT^{R141W} mice develop typical DCM phenotypes from 4 months of age, with death from HF after 8 months of age (27,49). cTnT^{R141W} transgenic mice displayed typical familial dilated cardiomyopathy phenotypes with dilated chambers, thin walls and cardiac dysfunction, as well as pathological phenotypes with myocytes disarray and fibrosis (4,26,27).

In the present study, it was identified that DAS treatment improved the DCM phenotypes of chamber dilation, wall thinning, myocyte disarray, fibrosis, poor myofibril organization and decreased ventricular blood ejection in cTnT^{R141W} DCM model mice. Furthermore, DAS treatment inhibited ROS production and decreased cytochrome *c* release, caspase 9-dependent caspase 3 activation, and thus, apoptosis of myocytes in cTnT^{R141W} DCM model mice.

The current results are in line with others. DAS analogues reduce the cytotoxicity associated with metabolism of alcohol, analgesics and xenobiotic by inhibiting CYP2E1 activity. In pathological conditions associated with CYP2E1 upregulation or CYP2E1-mediated adverse reactions (including diabetes/hyperlipidemia, Parkinson's disease, AIDS and cancer), DAS can reduce the damage caused by the pathological stimulus (18). DAS has been reported as a potentially effective intervention. The experimental results from previous studies have shown that DAS could not only inhibit ethanol and drug-mediated cytotoxicity, but also inhibit HIV proteinand diabetes-mediated toxicities by selectively inhibiting CYP2E1 in various cell types (18,50-53). In addition, the antioxidant/anti-inflammatory effects of DAS further support its use as dietary supplements (18). The current research, along with that of others, indicated that DAS could improve the symptoms of related diseases by targeting CYP2E1. In addition, the present study examined a NTG normal group with DAS treatment, and the heart structure and function of NTG group mice treated with DAS (400 mg/kg) did not shown abnormality in the echocardiography analysis (Table SIV). However, DAS, as a result of its rapid metabolism to DASO and DASO2, also causes cellular toxicity (18,54). Therefore, there is a need to modify the parent DAS into an analogue, which is a stronger inhibitor, but a weaker substrate of CYP2E1 (18,54).

Enalaprilat was selected in the present study to compare it with the therapeutic effect and characteristics of DAS. Enalaprilat is an ACE inhibitor that has been widely used in the clinical treatment of HF, including in patients with DCM (33,34,55-57). Enalaprilat is characterized by difficulties in direct absorption in the gastrointestinal tract and low oral absorption, and in terms of renal clearance, there is a barrier to entry of enalaprilat into the kidneys (58-60). DAS, as an active ingredient of garlic, is less toxic than other sulfides in garlic. The characteristic of DAS includes its action specificity, rapid metabolic clearance and cytotoxicity; therefore, DAS can be chemically modified to adjust its cytotoxicity and metabolic clearance rate (18,47,48).

In the present study, both of DAS and enalaprilat exhibited satisfactory inhibition on the pathological development of DCM in cTnT^{R141W} mice; however, each has its own characteristics. DAS treatment showed advantages in improving cardiac function and chamber dilation *in vivo*, as evidenced by the changes in echocardiographic parameters of LVFS, LVEF, LVESD and LVEDD. Furthermore, this improvement was more pronounced in the early stage of the treatment in both DAS groups than in the enalaprilat group, according to the significance findings after 2, 4 and 6 weeks of treatment. However, enalaprilat treatment was more effective in improving the microstructure and ultrastructure



Figure 4. Analysis of the mitochondria-dependent apoptosis pathways. (A) Cyt-*c* release and activation of caspase 9 and caspase 3 in the heart tissues of mice in the NTG, $cTnT^{R141W}$, 400, 200 mg/kg DAS and enalaprilat groups were detected via western blotting. (B) Cyt-*c* in the cyto and mito was semi-quantitatively analyzed using β -tubulin or VDAC1 for normalization. (C) Procaspase 9 and (D) procaspase 3 and active caspases 3 and 9 were semi-quantitatively analyzed using β -tubulin for normalization (n=3). (E) Cardiac myocyte apoptosis was detected using a TUNEL assay, and the arrows indicate TUNEL-positive cells (magnification, x400). (F) Number of positive cells was counted, and the proportion of apoptotic cells among the total cells in each image was calculated (n=3). *P<0.05, **P<0.01, ***P<0.001 vs. NTG group; #P<0.05, ##P<0.001 vs. cTnTR^{141W} mice. Cyt-*c*, Cytochrome *c*; cyto, cytoplasm; mito, mitochondria; VDAC1, voltage dependent anion channel 1; NTG, non-transgenic; DAS, diallyl sulfide.

of myocardium. In terms of the mechanism of action, DAS, rather than enalaprilat, exhibited an improved inhibitory effect on controlling oxidative stress levels, using H_2O_2 , MDA and GSH as indicators. Moreover, both of DAS and enalaprilat showed an excellent control effect on the mitochondrial apoptosis pathway.

The results of enalapril in improving oxidative stress are not satisfactory, and the exact mechanism remains unknown. However, it has been reported that MDA levels were not reduced in the enalapril-treated endothelial cells, while this treatment improved those of glutathione peroxidase and superoxide dismutase (61). This suggested that the specific indicators of enalaprilat in controlling oxidative stress are different from those detected in the present study, which may partly explain why the enalaprilat does not show a favorable control ability on improving MDA and other indicators of oxidative stress.

In addition, the current study selected a widely used myocardial cell line, H9c2, to verify the ameliorative effect of DAS on the pathological phenotype of cardiomyopathy *in vitro*. DAS could significantly inhibit the increased expression of CYP2E1 induced by ISO in H9c2 cells, and it also significantly improve the cell death induced by ISO, which were all consistent with the results *in vivo*.

According to previous research, underlying cellular processes, including inflammation and hypertrophy, myocyte apoptosis and necrosis, and deposition of extracellular matrix, have a direct association with the oxidative state of cardiac cells (54,62). Garlic and its active ingredients (such as phenols and saponins) have antioxidant effect (63). DAS can improve the oxidative stress induced by thallium acetate (TI), reduce the increase of MDA and nitric oxide levels in serum and liver, the decrease of GSH and catalase activities in liver and the decrease of total antioxidant capacity caused by TI (64). DAS can oxidize and glycosylate low-density lipoprotein to prevent additional oxidation or deterioration of glycosylation, which may benefit patients with diabetes-related vascular disease (65). Garlic oil is a source of DAS, and studies reported that garlic oil could be recommended as an option for treating hypertrophic cardiovascular disease (66). Cardiovascular diseases involve serum total cholesterol, low-density lipoprotein oxidation, platelet aggregation and hypertension (66). In vitro studies have shown garlic-induced inhibition of lipid synthesis, platelet aggregation, erythrocyte lipid peroxidation and low-density lipoprotein oxidation (66).

A limitation of the current study was that the improvement of DAS occurred in the early stage of the treatment, while the effect did not last a sufficient duration in the DAS low dose group. Therefore, future work will examine the range of diseases that DAS could be applied, as well as develop structural analogues with a lower cytotoxicity and metabolic rate.

In conclusion, the present study demonstrated that DAS had advantages in terms of improved chamber dilation and increased ventricular blood ejection. Furthermore, the regulatory mechanisms underlying DAS involvement in the DCM pathological development, included inhibition of both the oxidative stress levels and the mitochondria-dependent apoptosis pathways.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

LZ conceived the experiments and wrote the paper. DL analyzed the data and wrote the paper. SP performed most of the experiments and analyzed the data, including echocardiography parameters analysis, pathological analysis, western blotting, biochemical analysis and immunohistochemical analysis. WD contributed to the echocardiography procedure. NL and SG contributed to the genotyping and animal breeding and management. JL contributed to the TEM analysis. XZ contributed to animal administration procedures and RT-PCR analysis. LZ and DL confirmed the authenticity of all the raw data. All authors read and approved the final version of the manuscript.

Ethics approval and consent to participate

All animal procedures were approved by the Animal Care and Use Committee at the Institute of Laboratory Animal Science, Peking Union Medical College (approval no. ZLF18004). All mice were bred in an AAALAC-accredited facility and animal experiments were carried out in accordance with animal welfare requirements and the Guidelines for the Care and Use of Laboratory Animals published by the Chinese Ministry of Health.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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