Arachidonate 12/15-lipoxygenase-induced inflammation and oxidative stress are involved in the development of Diabetic Cardiomyopathy

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1 Abstract

2 Diabetes mellitus affects cardiac structure and function, and it has been suggested that diabetes leads to cardiomyopathy. Arachidonate 12/15-lipoxygenase (12/15-LOX) has 3 been suggested to play an important role in atherogenesis and heart failure. However, 4 $\mathbf{5}$ the role of 12/15-LOX in diabetic cardiomyopathy has not been examined. In this study, we investigated the effects of cardiac 12/15-LOX on diabetic cardiomyopathy. We 6 created streptozotocin (STZ)-induced diabetic mice and compared to those of Alox15 7 8 deficient (12/15-LOX KO) mice. Expression of 12/15-LOX as well as TNF-a and 9 NF- κ B were up-regulated in STZ-induced diabetic hearts. Disruption of 12/15-LOX 10 significantly improved STZ-induced cardiac dysfunction and fibrosis. Moreover 11 deletion of 12/15-LOX inhibited the increases of TNF- α and NF- κ B as well as the production of STZ-induced reactive oxygen species (ROS) in the heart. Administration 12of N-acetylcysteine (NAC) in diabetic mice prevented STZ-induced cardiac fibrosis. 13Neonatal cultured cardiomyocytes exposed to high glucose condition induced the 14 expression of 12/15-LOX as well as TNF- α , NF- κ B and collagen markers. These 15increases were inhibited by treatment of 12/15-LOX inhibitor. Our results suggest that 16cardiac 12/15-LOX-induced inflammation and oxidative stress are involved in the 17development of diabetic cardiomyopathy and that inhibition of 12/15-LOX could be a 18 19 novel treatment for this condition.

20

21 Keywords:

22 Inflammation, Oxidative stress, diabetic cardiomyopathy, 12/15-lipoxygenase

23

24 Non-standard Abbreviations and Acronyms:

25	12/15-LOX	12/15-lipoxygenase
26	12/15-HETE	12/15-Hydroxyeicosatetraenoic acid
27	CDC	Cinnamyl-3,4-dihydroxy-α-cyanocinnamate
28	FS	fractional shortening
29	LVDs	left ventricular systolic dimension
30	STZ	Streptozotocin
31	4-HNE	4-Hydroxy-2-nonenal
32	ROS	Reactive oxygen species
33	NADPH	nicotinamide adenine dinucleotide phosphate
34	Nox2	nicotinamide adenine dinucleotide phosphate
35		oxidase 2
36	Nox4	nicotinamide adenine dinucleotide phosphate

1		oxidase 4
2	NAC	N-acetylcysteine
3	NF-κB	nuclear factor -kappa B
4	TNF-α	Tumor necrosis factor-alpha
5	collagen1a2	collagen type I alpha2
6	collagen3a1	collagen type lll alpha1
7	Bip	immunoglobulin-heavy-chain-binding protein
8	CHOP	C/EBP-binding protein homologous protein
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10		

1 Introduction

Recently, the incidence of diabetes mellitus has increased rapidly, due to worldwide $\mathbf{2}$ changes in lifestyle. The epidemic of obesity together with sedentary lifestyles are 3 projected to result in over 300 million people having diabetes by 2025(1). According to 4 $\mathbf{5}$ the Framingham study, the relative rate of onset of heart failure in the diabetes is 6 significantly high, and onset of heart failure leads to a poor prognosis for diabetes patients(2). Diabetes is an independent risk factor for heart failure. In addition, the 7 8 number of cases of heart failure combined with diabetes continues to rise, and treatment will become a significant problem in the future(3). However, the detailed molecular 9 10 mechanism underlying development of this pathological condition has not been 11 elucidated yet(4; 5).

12 Arachidonic acid (AA) is a free fatty acid that is released from the cell membrane in response to various cytokines, peptides, and growth factors that become active under 13inflammatory conditions(6). There are three families of enzymes involved in the 14oxidative metabolism of AA. These include the LOXs, which produce 15hydroxyeicosatetraenoic acids (HETEs), the cyclooxygenases (COX-1 and COX-2) and 16cytochrome P-450 monooxygenases (7). The human LOX enzymes include 5-LOX, 1712-LOX and 15-LOX(8; 9). LOX enzymes are named according to the specific carbon 18 atoms of arachidonic acid that are oxidized. Thus, 12/15-LOX is a member of the LOX 19family that catalyzes the step from arachidonic acid to 12(S)-HETE and 2021 15(S)-HETE(10). 12/15-LOX was originally isolated from porcine leukocytes(11), but its tissue distribution is now known to be relatively wide, including blood vessels, the 2223brain, and the kidney(12).

Several lines of evidence have suggested that 12/15-LOX may play an important role in the development of atherosclerosis, nephropathy and neuropathy in diabetes mellitus(13-16). However, there is currently little evidence about a role played by 12/15-LOX in the development of diabetic cardiomyopathy. Therefore, we focused on the role of 12/15-LOX as a key molecule related to diabetic cardiomyopathy.

In the present study, we found that 12/15-LOX expression was significantly up-regulated in the heart of diabetic mice. We have shown that diabetes-induced activation of cardiac 12/15-LOX increased inflammation and oxidative stress in the diabetic heart. Conversely, disruption of 12/15-LOX reduces inflammation, oxidative stress and fibrosis in the diabetic heart, thereby improving systolic dysfunction. These findings suggest that inhibition of 12/15-LOX could be a novel treatment for diabetic cardiomyopathy.

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1 **Research Design and Methods**

 $\mathbf{2}$ Animal models. All experimental protocols were approved by Jikei University review board. Male C57BL/6 and Alox15-deficient (12/15-LOX KO) mice aged 7 weeks were 3 used in this study. Alox15-deficient mice on a C57BL/6 background were purchased 4 $\mathbf{5}$ from The Jackson Laboratory. For the diabetic model, 7-week-old male C57BL/6 and 6 Alox15-deficient mice were intraperitoneally injected with a single dose of streptozotosin (STZ) at 150mg/kg body weight (Wako 545-00283). One week after 7 8 induction of diabetes, the antioxidant N-acetylcysteine (NAC) (SIGMA 9165) was 9 administered to the three groups in the drinking water for 15 weeks (average 1.44 10 g/kg/day). 12/15-LOX inhibitor cinnamyl-3,4-dihydroxy-cyanocinnamate (CDC, 11 BIOMOL International, LP)(17), 8mg/kg once a day, was administered by subcutaneous 12injection for 4 weeks after induction of diabetes. We purchased the male db/db mice aged 11 weeks from CLEA Japan, Inc and used in this study. 13

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Physiological and histological analysis. Echocardiography was performed with a Vevo 15770 High Resolution Imaging System (Visual Sonics Inc.). To minimize variability of 16the data, heart rate was ~500-600 beats per minute when cardiac function was assessed. 17Average systolic pressure and heart rate were recorded by a photoelectric pulse device 18 (Softron BP-98A, Softron Co) placed on the tail of conscious mice. 4-µm frozen cross 19sections of the heart were fixed in 4% paraformaldehyde and subjected to 2021immunofluorescence staining for 12/15-LOX (Abcam ab87353), 4-Hydroxy-2-nonenal (4-HNE) (abcam ab20953) and Wheat germ agglutinin. Nuclei were stained with 2223Hoechst (Life technologies Japan H3570). Pictures were taken on a Biorevo microscope (Keyence, BZ-9000). Cardiac tissue was fixed by perfusion with 20% Formalin Neutral 24Buffer Solution (Wako 136-10041), sectioned at 4-µm thickness and stained. For 2526measurement of the percent area of cardiac fibrosis (% cardiac fibrosis), we selected 5 27fields at random and calculated the ratio of Masson-stained fibrosis area to total myocardium area with the software "Photoshop CS5" (Adobe Systems Inc.) as 28described previously(18). 29

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RNA analysis. Total RNA was isolated from the hearts of mice with RNA-Bee Reagent
(Cosmo Bio Inc.). In brief, after preparing total RNA, first-strand cDNA was
synthesized with QuantiTect® Reverse Transcription (QIAGEN Inc.). Real-time PCR
was performed using Thermal Cycler Dice TP800 (TAKARA Bio Inc.) with SYBR®
Premix Ex TaqTM II (TAKARA Bio Inc.) according to the manufacturer's instructions.
The specific oligonucleotide primers for GAPDH, TNF-α, collagen1α2 and collagen3α1

1 were selected using Primer3 (v. 0.4.0) (http://frodo.wi.mit.edu/).

 $\mathbf{2}$

Western blot analysis. Whole cell lysates were prepared in RIPA buffer 3 (Sigma-Aldrich Japan Ltd. R0278). Lysates (30 µg) were resolved by SDS-PAGE. 4 $\mathbf{5}$ Proteins were transferred to a PVDF membrane (Life Technologies Co. Ltd.), which 6 was incubated with the primary antibody, followed by anti-mouse, anti-rabbit IgG, light chain specific (The Jackson Laboratory). Specific proteins were detected using 7 8 enhanced chemiluminescence (ECL Prime) (GE Healthcare). The primary antibodies used for Western blotting were as follows: 4-HNE (abcam ab20953), α-Tubulin (Santa 9 10 Cruze Biotechnology, Inc sc-5286) and actin (Sigma-Aldrich Japan Ltd. A4700), 11 NF-κB p50, phosphor-NF-κB p65 (Cell Signaling Technologies #3035, #3039), Histone H3 (Cell Signaling Technologies #3638), Bip, CHOP (Cell Signaling Technologies 12 #3183, #2895). ELISA was performed according to the manufacturer's instructions to 13examine the levels of 12(S)-HETE and 15(S)-HETE (Assay Designs). We used the 14NE-PER® (Nuclear and Cytoplasmic Extraction kit, Thermo Fisher Scientific Inc.) for 15nuclear extraction of proteins. NIH 16the ImageJ software was used (http://rsbweb.nih.gov/ij/) to analyze intensity of Western blot bands. 17

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19 **Cell culture.** Cardiomyocytes and cardiac fibroblasts were prepared from neonatal rats 20 and cultured as described previously(19). Cardiomyocytes and cardiac fibroblasts were 21 exposed to a high glucose concentration (25 mmol/L glucose; HG) and harvested at the 22 indicated time points. Those were treated with normal glucose concentration (5.5 23 mmol/L glucose; LG) and 19.5mM mannitol to control for osmolarity.

Cardiomyocytes were cultured in media with HG with or without 10μM CDC and 1mM
N-acetyl-L-cysteine (NAC, Sigma-Aldrich Japan Ltd.) or with LG for 24-72 hours at
37°C. Palmitic acids (100uM) were treated after 48 hours after split of cardiomyocyte
and incubated for 24 hours.

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Measurement of intracellular ROS in cardiomyocytes. We assayed ROS production by using chloromethyl-2,7-dichlorodihydro-fluorescein diacetate (CM-H2DCFDA; Invitrogen C6827). Cardiomyocytes were cultured with 5μM CM-H2DCFDA at 37°C in the dark for 30 min and fixed the formation. CDC (10μM) was added 30min before treatment of high glucose. Pictures were taken on a Biorevo microscope (Keyence, BZ-9000). Fluorescence intensity was assessed for image analysis of the histogram by the software "Photoshop CS5".

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Assessment of the mitochondrial membrane potential. The mitochondrial membrane potential ($\Delta \Psi$ m) was assessed using Mito Tracker red. Briefly, cardiomyocytes were loaded with 0.5µM MitoTracker[®] Orange CMTMRos (Invitrogen M7510) at 37°C for 30 min, washed and fixed the formation. Fluorescence intensity was assessed by the "Photoshop CS5".

6

7 **Statistical analysis.** Data are shown as the mean s.e.m. Multiple group comparisons

8 were performed by one-way ANOVA, followed by Bonferroni's test for comparison of

9 means. Comparisons between two groups were made using the two-tailed unpaired

10 Student's *t* test or two-way ANOVA. In all analyses, P < 0.05 was considered

11 statistically significant.

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1 Results

2 Expression of 12/15-LOX pathway is up-regulated in the diabetic heart.

To determine whether 12/15-LOX is a key molecule in the development of diabetic 3 cardiomyopathy, we examined the mRNA level of 12/15-LOX in the diabetic heart 4 $\mathbf{5}$ using PT-PCR. We created STZ-induced diabetic mice (WT-STZ) and compared them 6 with control (saline) mice (WT). Cardiac expression of 12/15-LOX was significantly up-regulated with a peak at 4 weeks after STZ treatment in WT-STZ (Figure 1A). We 7 8 next examined a production of 12(S)-HETE and 15(S)-HETE, a major metabolite of 9 12/15-LOX. Production of 12(S)-HETE and 15(S)-HETE were also significantly 10 increased in the hearts of WT-STZ compared them with WT through the protocol until 11 16 weeks after STZ treatment (Figure 1B). Immunohistochemistry showed that 12expression of 12/15-LOX was specifically up-regulated in cardiomyocytes of diabetic heart, but not in vascular cell and fibroblast cell (Figure 1C). 13

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15 Animal characteristics of Wild and Alox15-deficient mice after STZ treatment

To investigate the relationship between up-regulation of 12/15-LOX and STZ-induced 16diabetic cardiomyopathy, we created STZ-induced diabetic mice using Alox15-deficient 17mice (KO) and compared them to WT-STZ mice. Animals were divided into 4 groups 18(WT, KO, WT-STZ, KO-STZ). Blood pressure (systolic blood pressure; SBP, diastolic 19blood pressure; DPB), heart rate, plasma glucose levels and body weight, in each animal 2021were measured at 0, 8 and 16 weeks after saline (WT, KO) or STZ (WT-STZ, KO-STZ) treatment. Single injection of STZ (150 mg/kg ip) to adult male mice resulted in 2223strongly high plasma glucose levels for a week after STZ treatment. Same dose of saline 24was injected as a control. High dose STZ-induced model has been established as a diabetes model and recognized as a useful model of diabetic cardiomyopathy(13; 20). 2526There was no difference in blood pressure and heart rate among the 4 groups under normal or STZ-induced high glucose conditions. Plasma glucose levels remained 2728around 10mmol/L in WT and KO mice without STZ treatment. No difference in plasma glucose level was observed between WT and KO, WT-STZ and KO-STZ at every 29weeks of age. Although body weight was not a tendency to increase at advancing age in 30 31mice after STZ treatment than control mice, there was no difference between WT and KO, WT-STZ and KO-STZ (Table). 32

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34 Disruption of 12/15-LOX improves cardiac dysfunction and fibrosis induced by 35 hyperglycemia.

36 Echocardiography was performed at 0, 8, 16 weeks after saline or STZ treatment and

1 compared between 4 groups. Fractional shortening (FS) was gradually impaired and left ventricular systolic dimension (LVDs) was increased in WT-STZ from 8 weeks after $\mathbf{2}$ induction of diabetes compared with those of WT. In contrast, systolic dysfunction and 3 left ventricular dilatation were not observed in KO and KO-STZ (Figure 2A). 4 $\mathbf{5}$ Histological examination revealed that 16 weeks after induction of diabetes, the fibrotic 6 area progressively extended to the peri-vascular and interstitial areas in WT-STZ heart 7 compared to WT heart. This increase was significantly inhibited in KO-STZ heart 8 compared to that of WT-STZ (Figure 2B). Quantitative analysis revealed that % cardiac fibrosis was significantly decreased in KO-STZ compared to those of WT-STZ (Figure 9 2C). These results suggest that 12/15-LOX affects cardiac dysfunction of STZ-induced 10 11 diabetic cardiomyopathy and the increased expression of 12/15-LOX-induced by 12hyperglycemia might cause cardiac dysfunction and fibrosis.

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14 Disruption of 12/15-LOX decreases cardiac inflammation and oxidative stress 15 induced by hyperglycemia.

We next examined the expression of inflammatory cytokine genes by RT-PCR. We 16found that cardiac expression of tumor necrosis factor- α (TNF- α) was up-regulated in 17WT-STZ heart compared to WT heart (Figure 3A). Cardiac collagen markers such as 1819 collagen1a2 and collagen3a1 were also increased in WT-STZ heart compared to WT 20heart at every four-week after STZ treatment, these increase were canceled by 21disruption of 12/15-LOX (Figure 3B). Moreover, we examined the relationship between 12/15-LOX expression and activation of NF-KB in WT-STZ heart and KO-STZ heart by 2223western blot analysis. Cardiac activation of NF- κ B was up-regulated in WT-STZ heart compared to WT heart. This up-regulation of NF-kB was also inhibited by disruption of 2412/15-LOX (Figure 3C, 3D), indicating that 12/15-LOX may induce cardiac 2526inflammation, which was involved in the development of STZ-induced diabetic 27cardiomyopathy.

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29 12/15-LOX induces cardiac oxidative stress in the diabetic heart.

We examined the relationship between 12/15-LOX and cardiac oxidative stress in the diabetic heart. Immunohistological staining and western blotting showed that expression of cardiac 4-HNE, a major marker of oxidative stress, was up-regulated in myocardium in WT-STZ heart compared to WT heart, and this enhancement was significantly inhibited by disruption of 12/15-LOX (**Figure 4A, 4B, 4C**). We next examined the nicotinamide adenine dinucleotide phosphate (NADPH) oxidase isoforms by RT-PCR at 4 weeks after induction of diabetes. Among the NADPH isoforms, expression of 1 nicotinamide adenine dinucleotide phosphate oxidase 4 (Nox4) but not nicotinamide

2 adenine dinucleotide phosphate oxidase 2 (Nox2) was up-regulated in WT-STZ heart

3 compared to WT heart and KO heart. This increase was also inhibited in KO-STZ heart

4 (Figure 4D).

 $\mathbf{5}$ To investigate the subcellular mechanism of the increase in reactive oxygen species 6 (ROS) in the diabetic heart, intracellular ROS levels in cardiomyocytes were estimated under HG and LG by fluorimetry in vitro. Cardiomyocytes under HG showed 7 8 enhancement of fluorescence intensity of DCF-DA by 7 fold at 24 hours compared to findings for those under LG (Figure 4E, 4F). Treatment with 12/15-LOX inhibitor 9 10 (CDC) inhibited the enhancement of DCF-DA fluorescence under HG condition (Figure 4E, 4F). To further investigate the subcellular origins of ROS production in the diabetic 11 heart, we used MitoTracker® Red to evaluate the effect of HG on the mitochondrial 12 membrane potential in cardiomyocytes. The result showed that there was loss of 13 mitochondrial membrane potential ($\Delta \Psi$ m) as indicated by a decrease in the 14fluorescence intensity of MitoTracker[®] Red in cardiomyocytes under HG. This decrease 15in the fluorescence intensity was improved by treatment with CDC (Figure 4E, 4F). 16

17 These *in vivo* and *in vitro* results suggest that activation of NADPH oxidase and 18 mitochondrial membrane abnormalities were occur in the diabetic heart and 12/15-LOX 19 pathway is involved in the process of production of ROS and oxidative stress in the 20 diabetic heart.

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Cardiac fibrosis and inflammation are ameliorated by treatment with antioxidant in the diabetic heart

24To investigate the relationship between cardiac oxidative stress and cardiac inflammation, we examined the hearts of WT-STZ mice who were administered the 2526antioxidant N-acetylcysteine (NAC) in their drinking water (WT-STZ NAC) throughout 27the experimental period and compared them to the heart of WT, KO, WT-STZ and 28KO-STZ mice. NAC was administered to diabetic mice for 15 weeks and histological analysis and cardiac expressions of TNF- α , collagen1 α 2, and collagen3 α 1 were 29examined. Histological examination revealed that cardiac peri-vascular fibrosis was 30 31improved in WT-STZ NAC heart as well as KO-STZ heart compare to findings in the 32WT-STZ heart (Figure 5A). Quantitative analysis revealed that the percentage of cardiac fibrosis was significantly decreased in KO-STZ and WT-STZ NAC hearts 33 compared to those of WT-STZ hearts (Figure 5B). Moreover, RT-PCR showed that 34increased expression of TNF- α , collagen1 α 2 and collagen3 α 1 in WT-STZ NAC and 3536 KO-STZ hearts were significantly inhibited compare with the expressions in WT-STZ

hearts (Figure 5C, 5D). These results indicate that cardiac oxidative stress has a major
 role in promoting cardiac inflammation in the STZ-induced diabetic heart.

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4 Cardiac 12/15-LOX pathway induced by high glucose condition increases the 5 expression of cardiac inflammation in vitro.

To further investigate the role of 12/15-LOX in cardiomyocytes under HG, we cultured neonatal cardiomyocytes under LG and HG using mannitol to adjust osmotic pressure for 24, 48 and 72 hours. RT-PCR showed that hyperglycemia up-regulated the expression of 12/15-LOX in cardiomyocytes (**Figure 6A**). Production of 12(s)-HETE and 15(s)-HETE were also increased in cardiomyocytes under HG condition (**Figure 6B**).

12Furthermore, RT-PCR demonstrated the up-regulation of TNF- α , collagen1 α 2 and collagen 3α 1 in cardiomyocytes under HG after 24 hours (Figure 6C, 6D). To determine 13 whether increased expression of cardiac 12/15-LOX affects the up-regulation of TNF- α , 14collagen1a2 and collagen3a1 in cardiomyocyte, we cultured neonatal cardiomyocytes 15with CDC under HG and examined the expression of each markers. Increased 16expression of TNF- α , collagen1 α 2 and collagen3 α 1 under HG were inhibited by 17treatment with CDC (Figure 6C, 6D). These results suggest that, cardiac 12/15-LOX 18 induced by hyperglycemia implicated the up-regulation of inflammatory cytokines such 19 as TNF- α , collagen1 α 2 and collagen3 α 1. 20

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22 Treatment of antioxidant inhibits cardiac inflammation induced by high glucose 23 condition in vitro

24We next investigated the relationship between the oxidative stress and inflammation induced by hyperglycemia in cardiomyocytes. Treatment with NAC under HG, RT-PCR 2526showed that up-regulation of TNF- α , collagen1 α 2 and collagen3 α 1 in cardiomyocytes 27was inhibited (Figure 7A, 7B). Moreover, western blotting revealed that nucleus cardiac 28activation of NF-KB using anti-NF-KB p50 antibody and anti phospho-NF-KB p65 (Ser468) antibody was up-regulated under HG compared to under LG after 24 hours. 29This activation of NF-κB was significantly inhibited by treatment with NAC in cultured 30 31cardiomyocytes (Figure 7C, 7D).

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1 Discussion

In the present study we demonstrate that the arachidonate 12/15-LOX is an important molecule involved not only in the onset and development of diabetic cardiomyopathy, but also in increased inflammation and oxidative stress in the heart.

It has been known for a long time that after STZ administration, 6 $\overline{7}$ alternations occur in the enzymatic activity of cardiac contractile proteins, 8 such as myosin and actomyosin, in rats and mice, and that cardiac contractile function gradually deteriorates (4; 21). This finding has provided 9 a basis for the well-known diabetic cardiomyopathy model, where STZ 10 11 administration quickly increases plasma glucose levels, and high plasma glucose levels are maintained for a long time, as shown in our experiments 12as well. In the hearts of wild-type mice exposed to increased plasma glucose 13levels, cardiac 12/15-LOX expression peaked at 4 weeks and its up-regulation 14persisted until 16 weeks after administration. Immunohistochemistry 15showed that the expression of 12/15-LOX in the heart was specifically 16 up-regulated in cardiomyocytes but not in vascular and fibroblast cells. 17Cardiac function gradually deteriorated from 8 weeks onward after STZ 18 19 administration, whereas this deterioration was suppressed in the Alox15-deficient mice. These results suggests that 12/15-LOX is involved in 2021the onset and development of STZ-induced diabetic cardiomyopathy.

Arachidonic acid (AA) metabolites are known to be involved in the 22development of tissue inflammation and fibrosis in non-cardiovascular 2324diseases such as pulmonary fibrosis (22; 23), suggesting an important role 25for AA as a potential pathway in the pathogenesis of cardiac fibrosis. A 26number of papers have demonstrated that the 12/15-LOX pathway induces 27inflammation in a variety of tissues. For example, overexpressed 12/15-LOX 28in macrophage-like J774.1 cells and HVSMC increased production of inflammatory cytokines such as interleukin-6 and TNF- α (24; 25). In an 29asthma model, 12/15-LOX inhibition reduced airway 30 experimental 31inflammation and attenuated cell injury (26). Furthermore, 12/15-LOX caused cell growth of cardiac fibroblasts and matrix production in the heart 32(27) and mice with cardiac overexpression of 12/15-LOX showed increased 33 the expressions of MCP-1 and TNF-a thereby inducing cardiac fibrosis and 34dysfunction (28). TNF-a is an important molecule that triggers inflammation 35 36 and cell injury in the heart (29), which is known to induce cardiac fibrosis as

a consequence (30; 31). Its suppression has been shown to improve diabetic 1 $\mathbf{2}$ cardiomyopathy, reduce cardiac fibrosis, and recover cardiac function (21; 32; 33). In agreement with these findings, our results showed that the 3 expressions of TNF-a and collagen markers were elevated accompanied by 4 an increase in the expression of 12/15-LOX in the STZ-induced diabetic heart, $\mathbf{5}$ 6 whereas these increases were suppressed in the Alox15-deficient mice $\overline{7}$ leading to cardiac function being preserved. Moreover, administration of a 12/15-LOX inhibitor (CDC) suppressed the up-regulation of TNF-a 8 associated with high blood glucose levels in vitro. Our findings suggest that 9 12/15-LOX-induced TNF- α may have a major role in the development of 10 11 cardiac fibrosis in STZ-induced diabetic cardiomyopathy.

Oxidative stress has been well-known to be involved in the 12development of diabetic cardiomyopathy, and increased ROS production is 13shown to induces various cardiovascular complications including cardiac 14dysfunction(32). Accumulating evidence suggests that mitochondria and 1516 NADPH oxidase are an important source of ROS production in the diabetic heart(34-36) where the crosstalk between mitochondria and NADPH oxidase 17are an important represents a feed-forward vicious cycle of ROS 1819 production(37; 38). It is important to elucidate the cellular mechanisms of ROS increase in the diabetic heart. Our histological examination revealed 2021that cardiac production of 4-HNE was increased in the diabetic heart and DCF fluorescence showed that intracellular ROS levels in cardiomyocytes 22were increased under high glucose conditions. We also showed the cardiac 2324expression of NOX4, one of the NADPH homologues, is elevated in the diabetic heart and the mitochondrial membrane potential is decreased in 2526cardiomyocytes exposed to high glucose levels in vitro. Moreover, disruption 27of 12/15-LOX decreased the production of 4-HNE and expression of NOX4 in the diabetic heart, while a 12/15-LOX inhibitor (CDC) 28decreased 29intracellular ROS levels in cardiomyocytes and restored the mitochondrial membrane potential. These results suggest that oxidative stress is increased 30 31 in the diabetic heart and that cardiomyocyte ROS production by mitochondria and NADPH oxidase plays a pivotal role in the onset and 32development of diabetic cardiomyopathy. Moreover, our results suggest that 33 34 12/15-LOX is an important molecule involved in the process of production of ROS and oxidative stress in the diabetic heart. 35

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Recently, diabetic patients with hypertension have increased in

numbers with the risk of cardiovascular disease thought to be increased in 1 $\mathbf{2}$ these patients. Research into molecules involved in both diseases is of critical importance, given that any such molecule will likely represent a viable 3 therapeutic target in the future. Diabetes not only causes hyperglycemia and 4 increases oxidative stress, but also exhibits a complicated disease condition $\mathbf{5}$ 6 associated with concomitant hypertension eventually leading to $\overline{7}$ atherosclerosis and organ injury(39). Persistent hyperglycemia associated with diabetes, insulin resistance, and postprandial hyperglycemia induces 8 inflammation and oxidative stress and causes myocardial injury(40; 41). We 9 previously confirmed by microarray analysis that 12/15-LOX 10 was 11 up-regulated in the heart of a hypertensive heart failure model and demonstrated that it is involved in the inflammation of the heart and 12development of cardiac fibrosis(28). Our present *in vivo* and *in vitro* studies 13demonstrate that 12/15-LOX is involved in the development of disease not 14only in the hypertensive heart failure model but also in the STZ-induced 15diabetic cardiomyopathy model, suggesting that 12/15-LOX is an important 16 molecule affecting the heart in both diseases. 17

It has been shown that 12/15-LOX is involved in the onset of a 18variety of diseases associated with type 1 and type 2 diabetes, such as 19 cardiovascular disease, hypertension, renal disease, and neurodegenerative 2021disorders. Furthermore, the role of 12/15-LOX has been suggested in humans as well. Indeed, the expression of 12/15-LOX was shown to be an 22independent risk factor for atherosclerosis in 828 patients with diabetes(42), 2324and 12/15-LOX inhibition is shown to lead to reductions in the size of 25cerebral infarction in humans(43-46).

26We acknowledge several limitations of our study. First, not all 27groups were evaluated over the entire course of cardiac fibrosis development in the study. Second, we could not entirely exclude the effects of the 28pharmacological agents used, such as CDC and NAC, especially on cardiac 29fibrosis, due to lack of several control groups in the study. Furthermore, 30 31while we demonstrated that the active involvement of ROS in high glucose leads to the upregulation of 12/15-LOX, which, in turn, results in cardiac 32fibrosis, the precise mechanisms linking 12/15-LOX to profibrotic signaling 33 34still remain largely unknown. Despite these limitations, however, we believe we have provided important new insights into diabetic cardiomyopathy. 35

In conclusion, this study demonstrated the mechanisms through

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which the expression of 12/15-LOX in the heart, associated with persistent hyperglycemia, leads to the development of diabetic cardiomyopathy via cardiac inflammation and oxidative stress. Moreover, it was suggested that inhibition of 12/15-LOX could potentially be a useful treatment for not only

- 5 diabetic cardiomyopathy but also for diabetic complications.
- 6

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- 9 conducted experiments, performed data analyses and wrote the manuscript. Y.K. and
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- 25 The authors declare no competing financial interests.
- 26
- 27 Disclosures
- 28 None
- 29
- 30

1 Table Animal characteristics of Wild and Alox15-deficient mice after STZ treatment

Wild type mice (WT) and *Alox15-deficient mice* (KO) were used to create a streptozotocin (STZ) –induced diabetic mce. Animals were divided into 4 groups (WT, KO, WT-STZ, KO-STZ). Blood pressure (Systolic blood pressure; SBP, Diastolic blood pressure ; DPB), heart rate, plasma glucose levels and body weight, in each animal were

6 measured at 0, 8 and 16 weeks after saline (WT, KO) or STZ (WT-STZ, KO-STZ)

7 treatment (Table 2). *P<0.05 vs. WT (0w), **P<0.01 vs. WT (0w), **P<0.05 vs. KO

8 (0w), ##P<0.01 vs. KO (0w), † P<0.05 vs. WT (16w), ¶ P<0.05 vs. KO (16w). Results

- 9 represent mean s.e.m. n=10-15.
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- 11

	Wild (WT)			Alox15-deficient (KO)				
Parameters		0w	8w	16w		0w	8w	16w
Blood pressure (mmHg)								
(SBP)	WT	105.7(3.4)	100.2(2.1)	116.3(4.8)	ко	105.1(4.7)	100.8(4.7)	112.3(3.8)
	WT-STZ	108.1(5.0)	108.6(4.9)	120.7(3.9)	KO-STZ	105.7(4.7)	102.6(5.4)	117.7(2.6)
(DBP)	WT	58.2(3.1)	70.3(0.8)	63.7(4.1)	ко	62.1(3.2)	59.9(3.2)	68.7(4.4)
	WT-STZ	60.2(4.0)	65.9(3.7)	65.7(3.9)	KO-STZ	58.1(3.8)	57.5(4.4)	64.4(2.3)
Heart rate (bpm)	WT	627(19)	655(9)	665(22)	ко	631(19)	618(38)	646(14)
	WT-STZ	615(25)	583(35)	601(22)	KO-STZ	595(28)	593(32)	590(16)
Plasma glucose (mol/l)	WT	10.1(0.4)	11.9(1.0)	10.3(0.7)	КО	10.7(0.6)	11.8(0.7)	11.0(0.7)
	WT-STZ	10.3(0.6)	31.2(0.5)**	33.3(0.2)**	KO-STZ	10.1(1.1)	31.1(1.7)##	30.9(0.6)##
Body weight (g)	WT	22.6(0.4)	26.8(1.0)*	29.9(1.4)**	ко	19.6(0.8)	21.2(1.9)	26.1(0.9)#
	WT-STZ	22.4(1.0)	20.9(0.5)* †	23.4(1.0)*	KO-STZ	22.4(0.8)	20.8(0.6)	22.5(1.3)#,¶

12

13

14

1 Figure Legends

2 Figure 1 Expression of 12/15-LOX pathway is up-regulated in the diabetic heart.

(A) Expression of 12/15-LOX in the hearts of wild-type (WT), wild-STZ (WT-STZ) 3 mice by RT-PCR. Cardiac expression of 12/15-LOX was up-regulated in WT-STZ 4 $\mathbf{5}$ compared to WT at every weeks of age. (B) 12/15(S)-HETE levels in the hearts of WT and WT-STZ by ELISA. Production of 12/15(S)-HETE was increased in the hearts of 6 WT-STZ compared to WT. *P<0.05 **P<0.01 WT-STZ vs. WT. Error bars indicate $\overline{7}$ 8 s.e.m. n= 10-15 in each group. (C) Immunofluorescence staining of 12/15-LOX in the hearts of WT and WT-STZ at 4 weeks after induction of diabetes. Expression of 9 10 12/15-LOX (red) was specifically up-regulated in cardiomyocytes of diabetic hearts. Wheat germ agglutinin (WGA) was used to label cardiomyocytes membranes (green). 11 12Upper panel is the staining of cardiomyocytes membranes (green). Middle panel is the staining of 12/15-LOX (red). Lower panel is the merged image of 12/15-LOX, 13cardiomyocytes membranes and hoechst staining. Scale bar, 60µm. 14

15

Figure 2 Disruption of 12/15-LOX improves cardiac dysfunction and fibrosis induced by hyperglycemia.

(A) Echocardiographic findings in wild-type (WT), wild-STZ (WT-STZ), *Alox15-deficient* (KO) and *Alox15-deficient-STZ* (KO-STZ) mice. Left ventricular FS
was gradually decreased and the LVDd was increased in WT-STZ compared with WT.
These changes observed in WT-STZ were further exacerbated by aging. FS and LVDs
were improved in KO-STZ compared to WT-STZ. *P<0.05 vs. WT, *P<0.05 vs.
KO-STZ (16w). Error bars indicate s.e.m. n= 10-15 in each group

(B) Masson Trichrome staining in the hearts of WT, WT-STZ, KO-STZ at 16 weeks
after induction of diabetes. An increase in peri-vascular and interstitial fibrosis was
observed in WT-STZ; this fibrosis decreased in KO-STZ mice. Scale bar, 60μm. The
percentage area of fibrosis in the hearts of WT, WT-STZ and KO-STZ. *P<0.05 vs. WT,
*P<0.05 vs WT-STZ. Error bars indicate s.e.m. n=7-10.

29

Figure 3 Disruption of 12/15-LOX decreases cardiac inflammation and oxidative stress induced by hyperglycemia.

32 (A, B) RT-PCR analysis for tumor necrosis factor- α (TNF- α), collagen markers 33 collagen1 α 2 and collagen3 α 1 in the hearts of wild-type (WT), wild-STZ (WT-STZ) and 34 *Alox15-deficient-*STZ (KO-STZ) mice at 4-16 weeks after induction of diabetes.

35 Cardiac expression of TNF- α , collagen1 α 2 and collagen3 α 1 was up-regulated in

WT-STZ compared to WT hearts. These increases were also inhibited by disruption of 1 12/15-LOX. *P<0.05 **P<0.01 vs. WT, *P<0.05 **P<0.01 vs. WT-STZ. Error bars 2 indicate s.e.m. n=7-10. (C) Western blot analysis of NF-kB activation in the hearts of 3 WT, WT-STZ and KO-STZ using anti-histone antibody and anti-NF-κB p50 antibody. 4 $\mathbf{5}$ We extracted the nuclear proteins of NF-kB p50 by using Nuclear and Cytoplasmic 6 Extraction kit. (D) Analyzing intensity of Western blot bands showed that cardiac activation of NF-kB was up-regulated in WT-STZ heart compared to WT heart. The 7 8 increased expression of NF-κB was also inhibited by disruption of 12/15-LOX. *P<0.05 9 vs. WT, [#]P<0.05 vs. WT-STZ. Data are shown as mean+s.e.m of duplicates and are 10 representative of 1 experiment out of 5.

11

12 Figure 4 12/15-LOX induces the cardiac oxidative stress in the diabetic heart.

(A) Immunohistological staining (brown) of 4-Hydroxy-2-nonenal (4-HNE) in the 13hearts of wild-type (WT), Alox15-deficient (KO), wild-STZ (WT-STZ) and 1415Alox15-deficient-STZ (KO-STZ) mice. Upper panel is x20, lower panel is x400, Scale bar, 1mm and 30µm. (B) Western blot analysis of 4-HNE expression in the hearts of WT, 16 17KO, WT-STZ and KO-STZ using anti-4-HNE antibody and anti-actin antibody. (C) Analyzing intensity of Western blot bands showed that cardiac 4-HNE level (38kD) was 18 significantly increased in WT-STZ compared to WT hearts and this increase was 19significantly inhibited by disruption of 12/15-LOX. *P<0.05 vs. WT, [#]P<0.05 vs. 20KO-STZ. Data are shown as mean+s.e.m of duplicates and are representative of 1 21experiment out of 5. (D) RT-PCR analysis for NADPH oxidase 2 (Nox2) and 4 (Nox4) 22in the hearts of WT, KO, WT-STZ and KO-STZ was examined at 4 weeks after 23induction of diabetes. *P<0.05 vs. WT, [#]P<0.05 vs. WT-STZ. Error bars indicate s.e.m. 24n=4-6. (E) Representative images of CM-H2DCFDA and MitoTracker[®] Orange 2526CMTMRos fluorescence in cultured neonatal cardiomyocytes under normal glucose (LG), high glucose (HG) and high glucose+12/15-LOX inhibitor (CDC 10µmol/L) 27(HG+CDC) groups. (F) Quantitative results of analysis. Treatment of CDC significantly 2829attenuated the increase of ROS caused by high glucose condition. *P<0.05 vs. LG, 30 [#]P<0.05 vs. HG. Error bars indicate s.e.m. n=4-6.

31

32 **Figure 5** *Cardiac fibrosis and inflammation are ameliorated by treatment with* 33 *antioxidant in the diabetic heart.*

- 34 (A) Masson Trichrome staining in the hearts of wild-type (WT), *Alox15-deficient* (KO),
- 35 wild-STZ (WT-STZ), Alox15-deficient-STZ (KO-STZ) and wild-STZ with NAC
- 36 (WT-STZ+NAC) at 16 weeks after induction of diabetes. An increase in peri-vascular

and interstitial fibrosis was observed in WT-STZ; this fibrosis decreased in 1 WT-STZ+NAC. Scale bar, 60µm. (B) The percentage area of fibrosis in the hearts of $\mathbf{2}$ WT, KO, WT-STZ and WT-STZ+NAC. *P<0.05 vs. WT, [#]P<0.05 vs WT-STZ+NAC. 3 Error bars indicate s.e.m. n=7-10. (C, D) RT-PCR analysis for TNF-α, collagen1α2 and 4 $\mathbf{5}$ collagen3a1 in the hearts of WT, KO, WT-STZ and WT-STZ+NAC at 4-16 weeks after 6 induction of diabetes. Cardiac expression of TNF- α , collagen1 α 2 and collagen3 α 1 were up-regulated in WT-STZ compared with WT hearts. These increases were also inhibited 7 by NAC. *P<0.05 **P<0.01 vs. WT, *P<0.05 **P<0.01 vs. WT-STZ+NAC. Error bars 8 9 indicate s.e.m. n=5-10.

10

11 Figure 6 Cardiac 12/15-LOX pathway induced by hyperglycemia increases the 12 expression of cardiac inflammation in vitro.

Cardiomyocytes were treated with high glucose condition (25 mmol/L; HG) or normal glucose condition (5.5 mmol/L; LG) for the indicated times (0-72h). (A) Expression of 12/15-LOX in cardiomyocytes by RT-PCR. (B) Production of 12/15(s)-HETE in cardiomyocytes by ELISA. *P<0.05 vs. LG, $^{#}P<0.05$ vs. HG. Error bars indicate s.e.m. n=4-7.

18 Cardiomyocytes were treated with 12/15-LOX inhibitor (CDC 10 μ mol/L) for the 19 indicated time (24-72 hours). (C, D) Expression of TNF- α , collagen1 α 2 and 20 collagen3 α 1 in cardiomyocytes treated with LG, HG and HG with CDC by RT-PCR.

21 Up-regulation of TNF- α , collagen1 α 2 and collagen3 α 1 under HG were inhibited by 22 treatment with CDC. *P<0.05 vs. LG, [#]P<0.05 vs. HG. Error bars indicate s.e.m. n=4-7.

23

Figure 7 Treatment of antioxidant inhibit cardiac inflammation induced by high glucose condition in vitro.

26Cardiomyocytes were treated with high glucose condition (25 mmol/L; HG) or normal glucose condition (5.5 mmol/L; LG) for the indicated times (0-72h) (A, B) Expression 2728of TNF- α , collagen1 α 2 and collagen3 α 1 in cardiomyocyte by RT-PCR. Up-regulation of TNF- α , collagen1 α 2 and collagen3 α 1 in cardiomyocytes under HG were inhibited by 29treatment with NAC. *P<0.05 vs. LG, [#]P<0.05 vs. HG. Error bars indicate s.e.m. n=4-7. 30 31(C) Western blot analysis of NF-κB activation in cardiomyocytes using anti-NF-κB p50 antibody, anti-phospho-NF-kB p65 (Ser468) antibody and anti-histone antibody. (D) 32Analyzing intensity of Western blotting showed that cardiac activation of NF-KB was 33 up-regulated in HG compared to LG after 24 hours. The activation of NF-KB was also 34inhibited by treatment with NAC. *P<0.05 vs. LG, [#]P<0.05 vs. HG. Data are shown as 35

36 mean+s.e.m of duplicates and are representative of 1 experiment out of 5.

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- 36

Figure1B



Figure1C

WGA



Merge

Bar.60µm





N=10-15, *: P<0.05, #: P<0.05

Figure 2B



Masson stain Bar.60µm x200



N=7-10, *: P<0.05, #: P<0.05

Figure 3A



Figure 3B



N=7-10, *P<0.05, **P<0.01 #P<0.05 ##P<0.01







Bar.30µm



N=4, *P<0.05, #P<0.05





Figure 4E



Figure 4F



Figure 5A

16 weeks



Masson stain Bar.60µm x200

Figure 5B



N=7-10, *: P<0.05, #: P<0.05

Figure 5C



Figure 5D







N=4-7, *P<0.05





N=4-7, *P<0.05



N=4-7, *P<0.05, #P<0.05



Figure 7B





N=4-7, *P<0.05, #P<0.05







N=5, *P<0.05, #P<0.05









Bar.60µm

0

WΤ

db/db

WT-STZ

Palmitic acid (100uM) N=4

N=4-6, *P<0.05, #P<0.05

