

Original Paper

CD38 Deficiency Protects Heart from High Fat Diet-Induced Oxidative Stress Via Activating Sirt3/FOXO3 Pathway

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Key Words

Cd38 • Oxidative stress • Heart • Oleic acid • Sirt3/FOXO3 pathway

Abstract

Background/Aims: Previous studies showed that CD38 deficiency protected heart from ischemia/reperfusion injury and high fat diet (HFD)-induced obesity in mice. However, the role of CD38 in HFD-induced heart injury remains unclear. In the present study, we have investigated the effects and mechanisms of CD38 deficiency on HFD-induced heart injury. **Methods:** The metabolites in heart from wild type (WT) and CD38 knockout (CD38^{-/-}) mice were examined using metabolomics analysis. Cell viability, lactate hydrogenase (LDH) release, super oxide dismutase (SOD) activity, reactive oxygen species (ROS) production, triglyceride concentration and gene expression were examined by biochemical analysis and QPCR. **Results:** Our results revealed that CD38 deficiency significantly elevated the intracellular glutathione (GSH) concentration and GSH/GSSG ratio, decreased the contents of free fatty acids and increased intracellular NAD⁺ level in heart from CD38^{-/-} mice fed with HFD. In addition, *in vitro* knockdown of CD38 significantly attenuated OA-induced cellular injury, ROS production and lipid synthesis. Furthermore, the expression of mitochondrial deacetylase Sirt3 as well as its target genes FOXO3 and SOD2 were markedly upregulated in the H9C2 cell lines after OA stimulation. In contrast, the expressions of NOX2 and NOX4 were significantly decreased in the cells after OA stimulation. **Conclusion:** Our results demonstrated that CD38 deficiency protected heart from HFD-induced oxidative stress via activating Sirt3/FOXO3-mediated anti-oxidative stress pathway.

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Introduction

Over-nutrition leads to metabolic-related changes in hearts including lipid accumulation and mitochondrial disturbances [1]. Lipid accumulation in cardiomyocytes is greatly correlated with lipotoxic effects on the heart including diastolic dysfunction [2] and increased oxidative stress [3]. Oxidative stress causes impaired cardiac fatty acid metabolism and endoplasmic reticulum (ER) stress, and induces other severe adverse effects such as insulin resistance [4]. In addition, overweight and obesity result in excessive activation of cardiac renin-angiotensin system [5], which further facilitates NADPH activation and ROS production. Therefore, the altered myocardial lipid metabolism may have a causative role in metabolic heart diseases.

CD38 (also known as cyclic ADP ribose hydrolase) is a membrane glycoprotein that plays important roles in cell proliferation, muscle contraction, hormone secretion and immune responses [6]. Specifically, CD38 acts as a multifunctional ectoenzyme to catalyze the synthesis of cyclic ADP-ribose (cADPR) from NAD⁺ and hydrolysis of cADPR to ADP-ribose (ADPR) [7-9]. Cyclic ADPR plays an important role in Ca²⁺ release from Ca²⁺ stores [10], whereas its main product, ADPR, can be covalently attached to proteins such as TRPM2, a Ca²⁺-permeable cation channel, by which it also modifies Ca²⁺ signaling [11, 12]. Meanwhile, CD38 can also catalyze NADP⁺ to NAADP, which is a potent trigger for Ca²⁺ mobilization [13]. In addition, CD38 can modulate the activity of sirtuins, a family of the NAD-dependent deacetylases that had beneficial effects in life span, mitochondrial function and energy metabolism in mammalian cells [14, 15].

In mammals, the sirtuin family comprises seven proteins (Sirt1–Sirt7), which are categorized as class III histone deacetylases. Sirt3 is expressed in a variety of tissues, especially in mitochondria-rich tissues such as brain, liver and heart tissues [16]. Sirt3 regulates global protein acetylation in mitochondria to control oxidative metabolism via using mitochondrial NAD⁺ as a substrate [17]. Many studies showed that Sirt3 played important roles in many diseases including cardiac and neurodegenerative diseases [18, 19]. Dysfunction of mitochondria is associated with production of ROS [20]. It has been reported that high levels of free fatty acid can induce oxidative damage, whereas lowering the levels of free fatty acid can reduce oxidative stress in hepatocytes [21]. Moreover, elevated activity of NADPH Oxidase 2 (NOX2) increases palmitate-induced mitochondrial ROS production in cardiomyocytes [22]. In contrast, Sirt3 attenuates ROS production by activating the transcription factor FOXO3a, and then trans-activating antioxidant genes such as manganese superoxide dismutase (MnSOD) and catalase in NAD⁺-dependent manner [23-25]. Moreover, the upregulation of the expression and activity of Sirt3 increases SOD2 activity via deacetylation of the lysine residues, and then reduces mitochondrial oxidative stress and intracellular ROS production [26, 27].

It has been reported that the intracellular levels of NAD⁺ were increased in various tissues from CD38 deficient mice [8]. Chini et al. demonstrated that CD38 deficiency protected mice from HFD-induced obesity [28]. Our previous studies showed that mouse embryonic fibroblasts (MEFs) isolated from CD38 knockout mice were resistant to oxidative stress via inhibiting ROS production and Ca²⁺ overload [29] and CD38 deficiency significantly enhanced cardiac functions in male mice [30]. Recently we observed that CD38 deficiency protected heart from ischemia/reperfusion injury via activating anti-oxidative stress pathway and inhibiting Ca²⁺ overload [31]. Moreover, our previous study showed that CD38 deficiency moderately improved the cardiac function in Ang-II induced cardiac hypertrophy [32]. However, the role of CD38 in HFD-induced heart injury is not evaluated. In the present study, we observed the metabolic changes of CD38-deficient heart, which may be associated with CD38 deficiency-mediated resistance to HFD-induced oxidative stress. In addition, we observed the significant changes of redox homeostasis, cofactor generation and lipid metabolism in CD38-deficient heart compared with the heart from wild type mice. Knockdown of CD38 significantly reduced lipid-induced oxidative damages in H9C2 cardiomyocytes.

Materials and Methods

Animal experiments

Eight week-old, male CD38^{-/-} mice and the genetics- and age-matched wild-type male mice (C57BL/6 background) were used in the experiments. CD38^{-/-} mice were obtained from Dr. Frances E. Lund (Rochest). The mice were maintained under controlled temperature (22-24°C) and illumination (12-h dark/light cycle) with free access of water. All animals were treated in accordance with the Guide for the Care and Use of Laboratory Animals, and approved by the Ethics Committee of Nanchang University. Mice were fed with normal diet or high-fat diets (60%-HFD, D12492; Research Diets Inc.) for 12 weeks. Then, the mice were anesthetized by 4% chloral hydrate and hearts were dissected and rinsed with phosphate-buffered saline and then immediately stored at -80°C for further experiments.

Metabonomic analysis

Hearts isolated from WT and CD38^{-/-} mice were prepared for metabonomic analysis using the automated MicroLab STAR® system from Hamilton Company. After removing protein, the samples were loaded on Waters ACQUITY ultra-performance liquid chromatography (UPLC) and a Thermo-Finnigan LTQ mass spectrometer (LC-MS) or analyzed on a Thermo-Finnigan Trace DSQ fast-scanning single-quadrupole mass spectrometer.

Cell Culture and transfection

H9C2 cells (American Type Culture Collection, Manassas, USA) were cultured in DMEM (Thermo Fisher, Waltham, MA, USA) supplemented with 10% fetal bovine serum (FBS, Gibco Ltd., Gaithersburg, USA), 100 U/ml penicillin and 100 mg/ml streptomycin at 37°C in an atmosphere of 5% CO₂. H9C2 stable cell line with CD38 knocking-down was prepared in our laboratory as previously described [31].

Oleic acid stimulation

Oleic acid (Sigma-Aldrich) stock solution was prepared in PBS with 20% BSA. This solution was diluted to 0.5 mM Oleic acid solution with DMEM. The medium with only BSA was used as control. H9C2 cells were incubated with OA (0.5 mM) for 24 h before analysis.

Cell viability assay

H9C2 cells were seeded into 96-well plates and incubated at a density of 1×10⁴ cells/well for 24 h. The cells were treated with 0.5 mM Oleic acid (OA) or BSA for 6 h and 12 h, respectively. Cell viability was detected with cell counting kit-8 (CCK8) from Dojindo, Japan. H9C2 cells were treated with CCK8 at 37°C for 3 h, the absorbance of 450 nm was measured using a microplate reader to quantify the formazan products. The amount of the formazan dye generated by dehydrogenases in cells was directly proportional to the number of living cells.

Measurements of LDH and SOD activities

LDH and SOD activities were detected using Cytotoxicity LDH Assay Kit-WST and SOD Assay Kit-WST (Dojindo, Mashiki-machi, Japan) according to the manufacturer's protocols, respectively. Briefly, cells (2×10⁵/well) were seeded into 6-well plates and treated with 0.5 mM OA or BAS for 24 h. For LDH activity assay, culture supernatant was collected and centrifuged at 1000 rpm for 5 min. Supernatant was taken, mixed with Working Solution in 96-well plate at room temperature for 20 min, and then added Stop Solution to each well, then immediately measured the absorbance at 490 nm by a microplate reader. For SOD activity assay, the cells were washed with PBS. The cells were lysed by ultrasonication. The samples were centrifuged at 10000 ×g for 10 min at 4°C. The supernatant was used for SOD activity (relative activity) determination according to the instruction. Absorbance was measured at 450 nm using a microplate reader. All experiments were performed at least three times.

ROS detection

The intracellular ROS production was measured using H2DCF-DA (Sigma-Aldrich). Firstly, 2×10^5 /well H9C2 cells were incubated with 10 μ M H2DCF-DA for 30 min. Then the cells were washed twice with PBS and collected for detection. Then the fluorescent dichlorofluorescein was monitored with automatic microplate reader at 488 nm excitation wavelength and 525 nm emission wavelength. All procedures should be kept away from light.

Quantitative Real-Time PCR analysis

Cells (2×10^5 /well) were plated in 6-well plate and treated with 0.5 mM OA or BAS for 24 h. Total RNA from cells were isolated using the Trizol method (Thermo Fisher) followed by DNase treatment. RNA content was measured by Nano Drop 2000 (Thermo Fisher). Then RNA was reversely transcribed using the Takara high capacity cDNA synthesis kit (TaKaRa, Dalian, China) according to the instruction. Relative expression of mRNAs was determined after normalized to GAPDH using the Δ Ct method. Quantitative PCR was performed using the ABI-ViiA7 PCR machine. The sequences of the primers are listed below: CD38, forward 5'-CTGCCAGGATAACTACCGACCT-3' and reverse 5'-CTTTCCCGACAGTGTGCTTCT-3'; Sirt3, forward 5'-CCCAATGTCGCTCACTACTT-3' and reverse 5'-AGGGATACCAGATGCTCTCT-3'; SOD2, forward 5'-GGACAAACCTGAGCCCTAA-3' and reverse 5'-GCGACCTTGCTCCTTATTG-3'; FOXO3, forward 5'-CCTGTCCTACGCTGACCTGAT-3' and reverse 5'-AGTCCCTTCGTTCTGAACCCG-3'; NOX4, forward 5'-AGTCAAACAGATGGGATA-3' and reverse 5'-TGTCCCATATGAGTTGTT-3'; Bax, forward 5'-GGGTGGCAGCTGACATGTTT-3' and reverse 5'-GCCTTGAGCACCAGTTTGC-3'; Bcl2, forward 5'-GTTGCAGTACCGGATTCTCT-3' and reverse 5'-CGGA GGTGGTGTGAATCCA-3'; GAPDH, forward 5'-AGCCAAAAGGGTCATCATCT-3'; reverse 5'-GGGGCCATCCACAGTCTTCT-3'.

Western blot analysis

H9C2 cells were cultured in 100-mm culture dishes at a density of 1×10^6 cells/dish. After treated with 0.5 mM OA or BAS for 24 h, the cells were lysed with RIPA buffer (1 mM PMSF was added) on ice for 30 min, and centrifuged at 12000 \times g for 10 min at 4°C. Protein concentration was determined with Pierce BCA Protein Assay Kit (Thermo Fisher Scientific, Rockford, USA). Lysates then subjected to SDS-PAGE electrophoresis, transferred to PVDF membrane (Millipore), and probed with the primary antibodies against CD38 (Santa cruz), FASN (Abcam), Acetylated-Lysine (CST), Sirt3 (CST), SOD2 (CST), FOXO3 (Abcam), NOX2 (Abcam), NOX4 (Abcam), Bax (CST) and GAPDH (Abcam). The bound antibodies were detected by secondary HRP-conjugated antibodies and the bands were visualized using the ECL system (Thermo Fisher Scientific, USA). Band images were obtained by using Molecular Imager ChemiDoc XRS+ (Bio-Rad, Hercules, CA, USA) and the intensity was analyzed by Image Lab™ software version 5.1 (Bio-Rad).

Statistical analysis

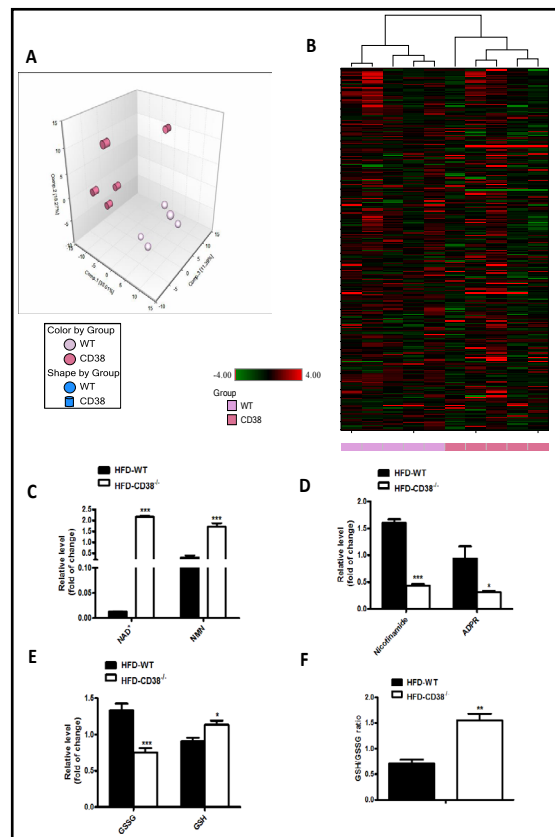
All data were presented as mean \pm SEM. The data were represented for three independent experiments. The statistical analyses were performed using SPSS19.0 software. The data were compared using Student's t test or one-way ANOVA. Differences were considered significant at $p < 0.05$.

Results

Deficiency of CD38 leads to significant changes of the global metabolomics of hearts

Although metabolic abnormality has been widely explored in many tissues with obesity, little is known about the global metabolic changes of the hearts in HFD condition. We used LC-MS and GC-MS to analyze metabolic fingerprints in hearts from CD38 deficient and wild type mice fed with HFD for 12 weeks. Principal component analysis and hierarchical clustering revealed a distinct difference in the two groups of samples, in which it may reflect significant metabolic shifts due to absence of CD38 (Fig. 1A-B). In addition, after peak detection, 271 known metabolites were acquired, and 87 of them were significantly altered ($p < 0.05$), including 27 metabolites were upregulated and 60 metabolites were downregulated. In addition, there were 17 compounds reaching statistically to a significant ($0.05 < p < 0.1$). An estimation of the false discovery rate (q-value) is calculated to take into account the multiple

Fig. 1. CD38 deficiency leads to significant changes of the metabolites in hearts. The metabolite levels in heart tissue from wild type and CD38 knockout mice were determined by Principal component analysis (PCA) (A) and unsupervised hierarchical clustering analysis (B) of metabolomics assay (n=5). The red represents the elevated metabolites and the green represents the depressed metabolites. The relative concentrations of nicotinamide adenine dinucleotide (NAD) and NAD⁺ precursor nicotinamide mononucleotide (NMN) in hearts from CD38 knockout and wild type mice (C). The relative levels of CD38 metabolites including nicotinamide and ADP-ribose in hearts from CD38 knockout and wild type mice (D). The concentrations of GSH and GSSG were measured in heart tissues from wild type and CD38 knockout mice (E). The GSH/GSSG ratios were determined in hearts from CD38 knockout and wild type mice (F). Data are shown as means ± SEM, **p<0.01, n = 5 per group.



comparisons that normally occur in metabolomic-based studies. A q-value threshold of <0.05 was used to remove any p-values (up to a 95% confidence) that could have been false positives.

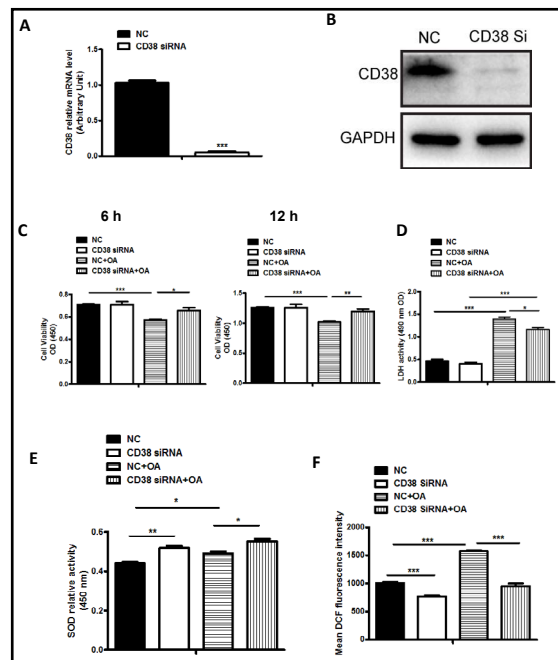
CD38 deficiency leads to marked changes of NAD⁺ metabolism including the increased intracellular GSH concentration and GSH/GSSG ratio in the heart of mice fed with HFD

Among the altered metabolites, we observed that there were significant changes in NAD⁺ metabolism in CD38 deficient hearts. The CD38 substrate NAD⁺ level was elevated in CD38 deficient heart (Fig. 1C). In addition, NAD⁺ precursor nicotinamide mononucleotide (NMN) was also increased (Fig.1C). However, the metabolites of CD38 including nicotinamide and ADP-ribose were significantly reduced in CD38 deficient hearts compared with those in WT group (Fig.1D). Furthermore, we found that two substances glutathione (GSH) and oxidized glutathione (GSSG) that were closely related to the oxidative stress were significantly changed in hearts of CD38 deficient mice compared to WT counterparts. The level of GSH in CD38 deficient heart was remarkably increased compared with those in WT, while GSSG was significantly decreased (Fig. 1E). Therefore, the GSH/GSSG ratio in CD38 deficient heart (1.55 ± 0.29) was remarkably increased compared with those (0.70 ± 0.18) in WT mice fed with HFD (p<0.01; Fig. 1F). These results indicated that CD38 deficiency protected hearts from HFD-induced oxidative stress.

Knockdown of CD38 protects from Oleic acid-induced cell death and oxidative stress

In order to investigate the protective effects of CD38 deficiency on lipid-induced oxidative stress *in vitro*, we observed the effects of CD38 deficiency on oleic acid-induced oxidative stress in cardiomyocytes using CD38 knockdown H9C2 stable cell lines. First, we examined CD38 expression in CD38 knockdown H9C2 stable cell lines. As expected, CD38 mRNA level was approximately decreased by 97% in the knocking-down stable cell lines (Fig.

Fig. 2. CD38 knockdown protects cardiomyocytes from oleic acid-induced cell death and oxidative stress. The CD38 mRNA (A) and protein (B) expressions were determined by q-PCR and western blot in the CD38 knockdown stable H9C2 cell lines and normal H9C2 cells (cardiomyocytes). Survival rates of cardiomyocytes were detected by CCK8 assay in CD38 knockdown H9C2 cells or control cells after incubation with 0.5 mM oleic acid for 6 h (left) and 12 (right) hours (C). The LDH (D) and SOD (E) activities in the supernatant of the cell culture after oleic acid stimulation were determined by measuring the absorbance (OD) at 490 nm and 450 nm, respectively, in CD38 knockdown H9C2 cells or normal cells. The mean fluorescence intensities of the OA-induced ROS production were quantitatively analyzed with different groups of cells (F). Data are shown as means \pm SEM, * p <0.05, ** p <0.01 and *** p <0.001, n = 3 per group.



2A). In addition, the CD38 protein level was also significantly downregulated in the CD38 knockdown H9C2 cells (Fig. 2B). After incubation with 0.5 mM oleic acid for 6 and 12 hours, the cells viability was determined using CCK8. As shown in Fig. 2C, the cell viability was significantly decreased in CD38 siRNA group compared with control siRNA group after OA stimulation for 6 and 12 hours, suggesting that CD38 knockdown might protect cells against cytotoxicity from oleic acid. Moreover, the results also showed that OA remarkably increased LDH activity and knockdown of CD38 significantly reduced LDH release in cardiomyocytes (Fig. 2D). In addition, the results showed that CD38 knockdown cells were resistant to OA-induced impairment of SOD activity in cardiomyocytes (Fig. 2E). Importantly, we found OA significantly increased ROS production in control cells, while CD38 knockdown reduced ROS generation (Fig. 2F). Taken together, these results indicated that CD38 deficiency protected cardiomyocytes from OA-induced oxidative stress *in vitro*.

CD38 deficiency significantly decreases lipid synthesis in vivo and in vitro

Besides GSH and GSSG, we also found that metabolites involved in lipid synthesis were changed in CD38 deficient heart compared with WT under HFD. Lipid overload was one of causes resulting in oxidative damage. In CD38 deficient heart, we observed that most of free fatty acids were significantly decreased compared with WT mice. As shown in Fig. 3A, all of the long chain fatty acids including stearate (18:0), nonadecanoate (19:0), eicosenoate (20:1) and arachidate (20:0) were reduced in CD38 deficiency group. Moreover, palmitate (16:0) and oleate (18:1n9) were also decreased although there was no statistical difference. And the polyunsaturated fatty acids (n3 and n6), such as linoleate (18:2n6), linolenate [α or γ ; (18:3n3 or 6)], docosadienoate (22:2n6) and dihomo-linoleate (20:2n6) were also decreased in CD38 deficient heart tissue (Fig. 3B). Moreover, we also observed that there was a significant decrease in lipid deposition in CD38 knockdown H9C2 cells treated with oleic acid through Oil Red O staining (Fig. 3C). And we also found that knockdown of CD38 significantly decreased intracellular TG level in H9C2 cells induced by oleic acid (Fig. 3D). Importantly, CD38 knockdown decreased the expression of key lipogenic gene FASN in H9C2 cells treated with oleic acid (Fig. 3E-F). These results suggested that decreased lipid synthesis in CD38 deficient heart and CD38 knockdown cardiomyocytes may reduce lipid-induced oxidative stress.

Fig. 3. CD38 deficiency leads to significant decreases of lipid synthesis in vivo and in vitro. The free fatty acids including long chain fatty acid (A) and polyunsaturated fatty acid (B) were quantitatively determined in hearts from CD38 knockout and wild type mice. The lipids of cardiomyocytes were detected by Oil red O staining in CD38 knockdown H9C2 cells treated with or without oleic acid (C). The intracellular TG levels were quantitatively measured in CD38 knockdown H9C2 cells under OA stimulation (D). The western blot images (E) and quantitative analysis (F) of the protein level of the lipid synthetic gene FASN were determined in H9C2 cells. Data are shown as means \pm SEM, * p <0.05, ** p <0.01 and *** p <0.001, n = 3-5 per group.

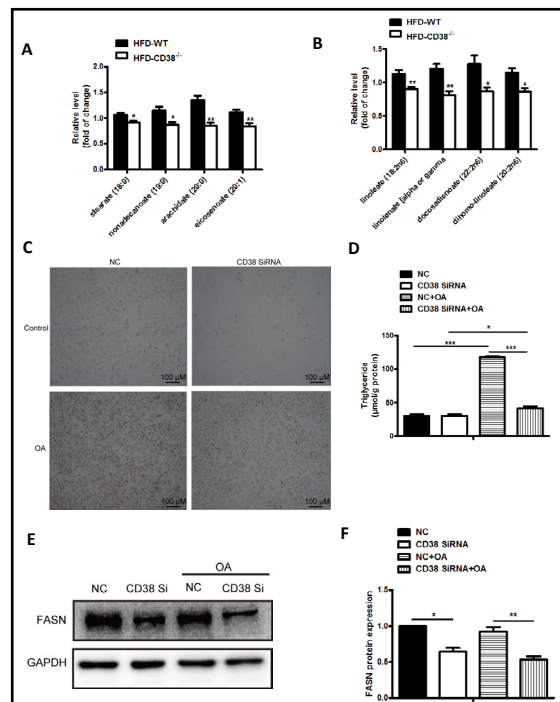
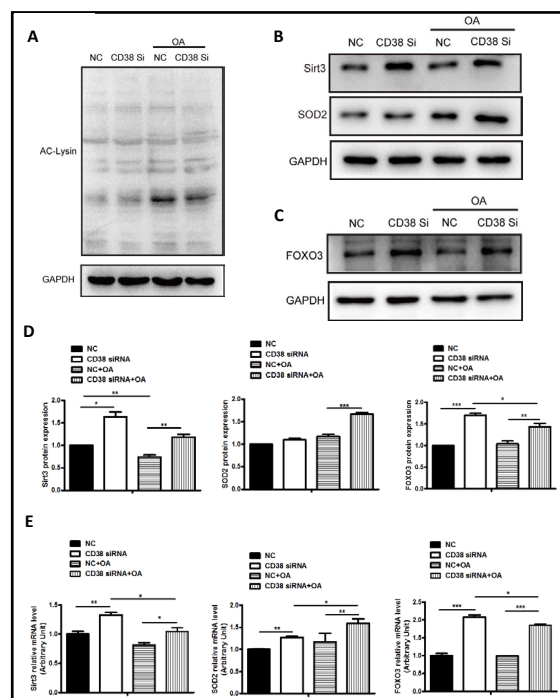


Fig. 4. CD38 knockdown increases the expressions of Sirt3, FOXO3 and SOD2 in oleic acid-treated cardiomyocytes. The protein acetylation levels of CD38 knockdown H9C2 cells with or without OA were analyzed by western blotting with antibody against acetylated-lysine (A). The expressions of Sirt3 (B), SOD2 (B) and FOXO3 (C) protein were determined by western blot analysis in CD38 knockdown H9C2 stable cells with or without oleic acid stimulation. The protein levels of Sirt3, SOD2 and FOXO3 were quantified from three different experiments (D). The mRNA expressions of Sirt3, SOD2 and FOXO3 were determined by qPCR (E). The values represent the means \pm SEM, * p <0.05, ** p <0.01 and *** p <0.001, n = 3 per group.



Knockdown of CD38 promotes the expressions of Sirt3, FOXO3 and SOD2 in Oleic acid-treated cardiomyocytes

As shown in Fig. 1C, the level of NAD⁺ was increased in CD38 deficient heart tissue. Sirt3 is a NAD⁺-dependent deacetylase which is mainly localized in mitochondria, and it deacetylates a number of mitochondrial proteins including manganese superoxide dismutase, which is responsible for limiting the accumulation of reactive oxygen species (ROS)[26]. In order to assess whether the mechanism of the protection form OA-induced oxidative stress was involved in the CD38/Sirt3 pathway, we first examined the total protein

acetylation levels. The results showed that OA increased the level of protein acetylation and CD38 knockdown decreased it compared with control group under OA stimulation (Fig. 4A). Moreover, the mRNA and protein level of Sirt3 was also examined by RT-qPCR and Western blot analyses, respectively. CD38 knockdown dramatically increased the expression of Sirt3 at both protein (Fig. 4B and 4D) and mRNA levels (Fig. 4E) in OA stimulated cells. In addition, the results showed that the expression of SOD2, one of the Sirt3 target genes, was also upregulated in both OA stimulated and vehicle treated CD38 knockdown H9C2 cells, especially under the OA stimulation (Fig. 4B, 4D and 4E). In addition, our results showed that the expression of FOXO3, another target gene of Sirt3 which also regulates SOD2 expression, was consistent with Sirt3 in CD38 knockdown cells (Fig. 4C-E). These findings suggested that CD38 deficiency protected cardiomyocytes from OA-induced oxidative stress through Sirt3-mediated activation of FOXO3/SOD2 signaling pathway.

Knockdown of CD38 attenuates the expressions of NOX2, NOX4 and Bax but increases Bcl2 expression in Oleic acid-treated cardiomyocytes

As NADPH oxidase was associated with ROS production, we next detected the effects of CD38 knockdown on the expression of NOX2 and NOX4 after OA stimulation. We observed that the expressions of NOX2 and NOX4 were decreased in CD38 knockdown cells at both protein and mRNA levels, especially the NOX4 (Fig. 5A, B). Furthermore, the expressions of the apoptotic-related proteins were markedly altered when the CD38 was knockdown, seen as the Bax expression was significantly decreased, whereas the Bcl-2 expression was markedly increased in response to OA stimulation (Fig. 5C-E). As showed in Fig. 5F, the ratio of Bax/Bcl2 was significantly decreased in CD38 knockdown H9C2 cells with or without oleic acid stimulation. The working model of protective effects of CD38 deficiency on HFD-induced heart injury was presented in Fig. 6.

Fig. 5. CD38 knockdown decreases the expressions of NOX2, NOX4 and Bax but increases Bcl2 expression in oleic acid-treated cardiomyocytes. The expressions of NOX2 and NOX4 protein were determined by western blot analysis with or without oleic acid stimulation in CD38 knockdown or normal H9C2 cells (A). The expressions of NOX4 mRNA were determined by qPCR with or without oleic acid stimulation (B). The expressions of Bax protein were determined by western blot analysis in CD38 knockdown or normal H9C2 cells with or without oleic acids (C). The mRNA expressions of Bax (D) and Bcl2 (E), and the ratio of Bax/Bcl2 (F) were quantitatively determined by qPCR in CD38 knockdown or normal H9C2 cells with or without oleic acid stimulation. The values represent the means \pm SEM, * p <0.05, ** p <0.01 and *** p <0.001, n = 3 per group.

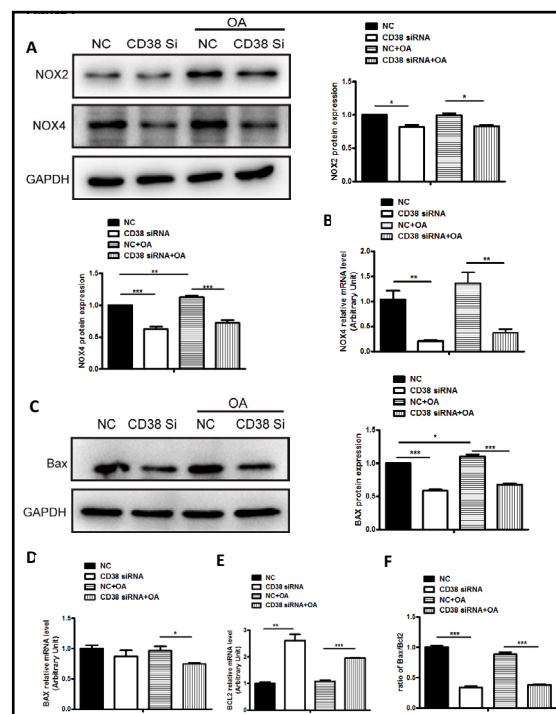
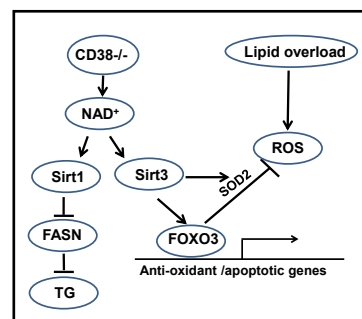


Fig. 6. The working model for the protection of CD38 deficiency on the lipid-induced oxidative stress in cardiomyocytes. CD38 deficiency increases the intracellular NAD⁺, and the increased NAD⁺ as a substrate activates Sirt1 signaling pathway, then inhibiting FASN. Finally, the content of triglycerides in heart was decreased. On the one hand, the increased NAD⁺ also activates Sirt3 which is located in the mitochondria and then promoted the nuclear translocation of FOXO3, resulting in the increases of the expressions of antioxidant/anti-apoptotic genes including SOD2 and Bcl2. On the other hand, Sirt3 inhibits the productions of the lipid-overloading induced ROS through directly activating SOD2.



Discussion

Overweight or obesity, especially abdominal obesity, has become an independent risk factor for many diseases such as cardiovascular diseases, diabetes, multiple types of cancer and renal diseases [33, 34]. It has been reported that mice consuming a Western diet manifested increased heart and vascular fibrosis which further resulted in cardiac hypertrophy and dysfunction [35]. Individuals with a BMI greater than 30 had more signs of early vascular pathological changes than non-obese people such as coronary arteries dysfunction. Under normal conditions, triglycerides are mainly stored in adipocytes and only a limited triglycerides are stored in the non-fat cells [36]. When adipose tissue fails to store all excess lipids, they will accumulate in other tissues such as muscle, liver and heart. Lipid accumulation in heart was positively correlated with the degree of obesity. In many obese persons, excessive lipid accumulation in and around heart was observed, and the epicardial and intramyocardial lipids were detrimental to cardiac function [37].

It has been reported that CD38 deficiency mice were resistant to high fat diet induced obesity [28]. Our previous data also showed that CD38 deficiency alleviated cardiac function and ROS production in different models. However, whether CD38 could confer protective effects on heart after high fat diet has not been reported. Based on our previous study, we further examined the effects of CD38 deficiency on heart under high fat diet. In the present study, we investigated the metabolic changes in the hearts of CD38 deficiency mice after high fat diet using global metabolomics, a promising qualitative and quantitative analysis approach for all small molecules in biological samples under various stimuli [38]. Our results showed that CD38 deficient hearts exhibited numerous metabolic changes especially in redox homeostasis, nicotinate and nicotinamide metabolism and lipid metabolism compared with wild type. Many studies demonstrated that the imbalance of redox state and oxidative stress was associated with HFD [39, 40]. In fact, the increased production of reactive oxygen species, oxidative stress and mitochondrial dysfunctions in heart and other tissues were observed in obesity and obesity-associated metabolic diseases [41, 42]. In the present study, we found that CD38 deficiency significantly decreased oxidative stress under HFD, evidenced by the increased reduced glutathione (GSH) and decreased oxidized glutathione (GSSG) levels, in which the GSH/GSSG ratio was significantly increased in CD38 deficient hearts compared with WT. GSH is an important antioxidant as its thiol group can be oxidized to GSSG by reactive oxygen species. The intracellular GSH/GSSG ratio is considered to be a valid index of oxidative stress. Our results indicated that the increase of the GSH/GSSG ratio in CD38 deficient hearts was one of the mechanisms for decreasing oxidative stress induced by HFD.

It is well-established that high concentrations of free fatty acids or cholesterol in blood are risk indicators for some diseases such as coronary heart disease and diabetes [43, 44]. Furthermore, it has been reported that an impaired ability of the heart to oxidize excess lipids resulted in an accumulation of intra-myocardial fatty acid metabolites including triglyceride, long chain acyl CoA, ceramide and diacylglycerol (DAG), nevertheless, lowering these harmful

lipid metabolites can improve cardiac dysfunction [45, 46]. Our study using metabolomic analysis also showed that most of the free fatty acids were reduced while kinds of carnitine were elevated, as well as the key metabolic intermediate acetyl coenzyme A was increased almost 15 folds in heart tissue from CD38 deficient mice, highlighting the importance of lipid metabolism in CD38-mediated resistance to lipid-induced oxidative stress in hearts.

CD38 is a major NAD⁺ hydrolase in mammals which can use NAD⁺ as a substrate to generate nicotinamide and an ADP-ribosyl product. Microarray data revealed that CD38 was highly expressed in adipose tissues in obese people [47, 48]. In addition, many studies have also shown that CD38 expression is a reliable negative prognostic marker for chronic lymphocytic leukemia (CLL) patients [49-51]. Therefore, the overexpression of CD38 is correlated with many human diseases. It has been reported that CD38 deficiency increased autophagy [52], while autophagy is closely related to cardiac diseases. In the present study, we found that the intracellular concentration of NAD⁺ in hearts from CD38 deficient mice was 177 folds higher than wild type mice, which was consistent with our previous study in CD38 knockout or knockdown models [7, 53]. In addition, the intermediate nicotinamide mononucleotide was increased by approximately 6 folds while the nicotinamide and ADPR, the metabolites of CD38, were significantly reduced in CD38 deficient heart under treated with HFD. The canonical role of NAD⁺ is to facilitate hydrogen transfer in key metabolic pathways. Conversion of NAD⁺ to NADH is also important for oxidation of fatty acids and amino acids in mitochondria. Therefore, the increased NAD⁺ promoted the utilization of NAD⁺ for lipid metabolism in CD38 deficient heart, attenuating HFD-induced oxidative stress. In addition, increased NAD⁺ also activates the Sirtuins family, such as Sirt1 and Sirt3, the NAD⁺-dependent deacetylases [14, 54], which deacetylate kinds of target genes involved in anti-oxidative stress pathway. To confirm the protective effects of CD38 deficiency on lipid-induced injury in cardiomyocytes, a CD38 knockdown H9C2 stable cell line was used to elucidate the mechanisms. Our results showed that CD38 knockdown significantly increased cell survival after the cells treated with OA. Moreover, CD38 knockdown also improved OA-induced oxidative stress in H9C2 cells, manifested by decreased LDH activity and increased SOD activity. In addition, we also observed that ROS production was significantly increased when cells were treated with oleic acid, while CD38 knockdown reduced its generation compared with control group. These findings indicated that CD38 deficiency alleviated lipid-induced oxidative stress *in vivo* and *in vitro*.

Sirt3 pathway was one of the most important pathways for protecting cells against oxidative stress [26, 55]. It has been reported that Sirt3-deficient cardiomyocytes had high levels of ROS and Sirt3 reduced doxorubicin-induced ROS production mainly via regulating the key antioxidant SOD2 [55]. In addition, a recent study indicated that Sirt3/SOD2 signaling also played a protective role in mitochondrial oxidative stress induced by bile acid in hepatocytes [27]. The reduced oxidative stress by Sirt3 not only relied on the deacetylation of SOD, but also the deacetylated FOXO3 to protect mitochondria against oxidative damage [56]. Sundaresan et al. reported that Sirt3 induced FOXO3a translocation to the nucleus and augmented FOXO3a-dependent antioxidant genes expression, mainly through upregulation the activities of MnSOD and Catalase [23]. In the present study, we detected the acetylation level of total protein which indirectly reflected the activity of Sirtuins. And the results showed that oleic acid increased acetylation level while CD38 knockdown reduced it compared with control group, indicating that CD38 knockdown increased Sirtuins activity under OA treatment. Moreover, we also examined the expressions of oxidative stress relative signaling molecules Sirt3, SOD and FOXO3. The results also showed that the expressions of Sirt3 and its target genes such as SOD and FOXO3 were upregulated in CD38 knockdown cells with or without OA stimulation. Taken together, these experimental results indicated that CD38 knockdown reduced lipid overload-induced oxidative stress in cardiomyocytes possibly through activating Sirt3/FOXO3 signaling pathway. _ENREF_37

NAPDH oxidases (NOXs) are responsible for ROS generation, in which the NOX2 and NOX4 are significantly associated with cardiovascular diseases [57]. Studies showed that NOX2 inhibition could prevent oxidative stress induced by saturated fatty acids in

cardiomyocytes [22] and NOX4 regulated basal and pathological ROS production [58, 59]. It has been reported that the NOX4 deficiency mice were sensitive to HFD-induced obesity, suggesting its anti-adipogenic role [60]. In the present study, we observed that NOX2 and NOX4 expressions were significantly down-regulated in CD38 knockdown cells after 24 hr OA stimulation, especially NOX4. It was well known that oxidative stress was often accompanied by apoptosis. Our results showed that the anti-apoptotic factor Bcl-2 expression was increased, whereas the proapoptotic factor Bax expression was decreased in CD38 knockdown cells treated with OA, suggesting that the reduced NOX expressions and oxidative stress-induced apoptosis may also partially contribute to the protection of CD38 deficiency on the lipid overload injury in cardiomyocytes.

Conclusion

Our results from *in vivo* and *in vitro* studies demonstrated that CD38 deficiency significantly protected hearts from lipid-induced injury via regulating redox homeostasis, lipid metabolism and apoptosis in myocardial cells through activating Sirt3/FOXO3 signaling pathway. Our findings provide new insights into the elucidation of the mechanisms of lipid-induced oxidative stress and heart injury.

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Disclosure Statement

The authors declare no competing financial interests.

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