

Trichostatin A attenuates oxidative stress-mediated myocardial injury through the FoxO3a signaling pathway

YUNHUI GUO^{1,2}, ZHIPING LI¹, CANXIA SHI¹, JIA LI¹, MENG YAO¹ and XIA CHEN¹

¹Department of Pharmacology, College of Basic Medical Sciences, Jilin University, Changchun, Jilin 130021; ²Department of Pharmacology, Jiamusi College, Heilongjiang University of Traditional Chinese Medicine, Jiamusi, Heilongjiang 150040, P.R. China

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Abstract. Trichostatin A (TSA), a histone deacetylase inhibitor, is widely used as an anticancer drug. Recently, TSA has been shown to exert a protective effect on ischemia/reperfusion (I/R) injury; however, the underlying mechanisms remain unclear. Forkhead box O3a (FoxO3a), a unique FoxO family member, has been shown to attenuate myocardial injury by increasing resistance to oxidative stress in mice. The present study aimed to investigate whether TSA exerts its cardioprotective effects through the FoxO3a signaling pathway. For this purpose, healthy male Wistar rats were pre-treated with TSA for 5 days before they were subjected to ligation/relaxation of the left anterior descending branch of the coronary artery and to 30 min of ischemia, followed by 24 h of reperfusion. The activities of creatine kinase (CK), lactate dehydrogenase (LDH), aspartate aminotransferase (AST) and superoxide dismutase (SOD), as well as the malondialdehyde (MDA) levels were examined. The H9c2 rat myocardial cell line was cultured in 10% FBS-containing DMEM for 24 h. The cells were incubated with/without TSA (50 nmol/l) for 1 h and then incubated with/without H₂O₂ (400 μ M) for 2 h. Reactive oxygen species (ROS) and mitochondrial membrane potential ($\Delta\psi_m$) were measured by probe staining in the H9c2 cells. The expression of FoxO3a, mitochondrial SOD2 and catalase was quantified by western blot analysis. The levels of H3 and H4 acetylation of the FoxO3a promoter region were examined by chromatin immunoprecipitation assay. TSA significantly reduced the myocardial infarct size and the activities of serum LDH, AST and CK in the rats. TSA also decreased the levels of MDA and

increased the activities of SOD in the myocardial tissue of the rats. Consistent with the reduced injury to the TSA-treated rats, TSA significantly reduced the H₂O₂-induced levels of ROS and increased $\Delta\psi_m$. In addition, TSA increased the expression of FoxO3a, SOD2 and catalase, which may be related to increasing the level of H4 acetylation of the FoxO3a promoter region. Our results thus revealed that TSA protected the myocardium from oxidative stress-mediated damage by increasing H4 acetylation of the FoxO3a promoter region, and the expression of FoxO3a, SOD2 and catalase.

Introduction

Acute myocardial infarction (MI) is a tremendous threat to the life and health of the population. However, accumulating evidence indicates that directly restoring the blood supply through reperfusion following MI in fact aggravates the damage. This additional damage, namely ischemia/reperfusion (I/R) injury (1-3) is a hotspot of current research. Recent studies have reported that the pathogenesis of myocardial I/R injury is multifactorial, involving the dysfunction of energy metabolism, robust reactive oxygen species (ROS) generation, cell calcium overload, inflammatory responses and apoptosis (4-8). Oxidative stress resulting from ROS may be a significant feature in the development of cardiac injury induced by I/R (9); however, the underlying regulatory mechanisms of cardiac oxidative stress resistance remain unclear. Forkhead box O3a (FoxO3a) plays an important role in metabolism, cell survival and resistance to oxidative stress in multiple cell types (10). However, its role in regulating tolerance to cardiac stress remains to be fully elucidated. FoxO3a interacts with the promoter of the superoxide dismutase 2 (SOD2) gene at a specific binding site in vascular smooth muscle cells (11); however, whether it influences SOD2 expression in cardiomyocytes has not yet been reported, at least to the best of our knowledge.

The nucleosome is the basic structural unit of eukaryotic chromatin, which is made up of histones and DNA. Histone acetylation modification affects chromatin packaging and gene expression (12). Transcriptional activity is found in areas where the histone acetylation level is higher, and transcriptional silencing in areas where the histone acetylation level is usually lower (13). The acetylation of histone is a reversible process, which is regulated by histone acetyltransferases (HATs) and

Correspondence to: Professor Xia Chen, Department of Pharmacology, College of Basic Medical Sciences, Jilin University, 126 Xinmin Street, Changchun, Jilin 130021, P.R. China
E-mail: chenx@jlu.edu.cn

Abbreviations: TSA, trichostatin A; I/R, ischemia/reperfusion; MI, myocardial infarction; TTC, 2,3,5-triphenyltetrazolium chloride; AST, aspartate aminotransferase; CK, creatine kinase; LDH, lactate dehydrogenase; ROS, reactive oxygen species; $\Delta\psi_m$, mitochondrial membrane potential; SOD2, superoxide dismutase 2

Key words: trichostatin A, myocardial ischemia/reperfusion, forkhead box O3a, mitochondrial membrane potential, acetylation

histone deacetylases (HDACs) (14), and the dynamic balance between them plays a very important role in eukaryotic gene regulation (15). It has been demonstrated that HDAC inhibitors play a protective role in cardiovascular diseases, such as cardiac hypertrophy, inflammatory responses and heart failure (16).

Trichostatin A (TSA), a histone deacetylase inhibitor, is used as a promising anticancer agent (17,18). Our previous study demonstrated that TSA attenuated myocardial I/R injury (19); however, the underlying mechanisms remain to be elucidated. It has been shown that FoxO3a attenuates myocardial injury by increasing resistance to oxidative stress in mice (20) and that the endogenous HDAC inhibitor, β -hydroxybutyrate, can increase histone acetylation at the FoxO3a promoters in 293 cells (21). We therefore speculated that the mechanisms through which TSA attenuates myocardial injury may be associated with FoxO3a. In the present study, we demonstrate that TSA attenuates myocardial injury by increasing resistance to oxidative stress. We further investigated whether the mechanisms involved in these effects are mediated through the FoxO3a signaling pathway.

Materials and methods

Ethical approval of the study protocol. The study was approved by the Ethics Committee of Jilin University, and all experimental procedures conformed to the guidelines for the Animal Care and Use Committee of Jilin University, Changchun, China.

Animals and model of myocardial I/R in rats. A total of 80 healthy male Wistar rats (weighing, 180–220 g) were purchased from the Animal Centre of Jilin University (certificate number SCXK (JI) 2007-0003). In total 40 Wistar rats first were subjected to MI by ligating the left anterior descending branch of the coronary artery and to 30 min of ischemia, followed by 0, 6, 12 or 24 h reperfusion in order to determine the effects at different time points on FoxO3a expression; 10 rats were used at each time point. The other 40 rats were randomly divided into 5 groups as follows: i) the sham-operated (control), treated with saline (n=8); ii) the I/R group, in which rats were subjected to I/R and treated with saline (n=8); iii) the TSA 1 group, in which rats were subjected to I/R and treated with 0.2 mg/kg (I/R+T1) (n=8); iv) the TSA 2 group, in which rats were subjected to I/R and treated with 0.1 mg/kg (I/R+T2) (n=8); and v) the TSA 3 group, in which rats were subjected to I/R and treated with 0.05 mg/kg (I/R+T3) (n=8). The rats in the 5 groups were injected with TSA or saline by intraperitoneal administration once daily for 5 days. Standard rat chow and water were available *ad libitum*. On day 5, the animals in the 5 groups underwent 30 min left anterior descending (LAD) coronary artery ligation and 24 h of reperfusion according to the method previously described by Li *et al* (22). Briefly, the rats were anesthetized by an intraperitoneal injection of 10% chloral hydrate (0.35 g/kg) before undergoing a thoracotomy. After ensuring the appropriate depth of anesthesia, the rats were intubated without incision using a rodent ventilator. The thoracic cavity was opened and the heart was rapidly exposed. The left anterior descending coronary artery (LAD) was ligated by a 6-0 silk suture (2–3 mm below the left auricle) and the chest was immediately closed. Successful I/R injury was monitored by standard lead II electrocardiography (ECG). The sham-

operated rats only underwent the suture around the LAD, but were not ligated. Following reperfusion, the myocardial infarct size, serum myocardium enzyme activities and protein expression were evaluated.

Cell culture. The H9c2 rat myocardial cell line was obtained from the Cell Bank of Chinese Academy of Sciences (Shanghai, China). The cells were cultured in a 5% CO₂ humidified atmosphere at 37°C in 10% fetal bovine serum (FBS)-containing Dulbecco's modified Eagle's medium (DMEM). The cells were seeded in 6-well-plates for 24 h and were then incubated with/without TSA (50 nmol/l) for 1 h and then incubated with/without H₂O₂ (400 μ M) for 2 h.

Chemicals and antibodies. TSA was purchased from TCI Development Company Ltd. (Shanghai, China). Polyclonal rabbit antibody against catalase was purchased from Abcam (ab76110; Cambridge, CA, USA). Polyclonal rabbit antibodies to FoxO3a (12829) and SOD2 (13141) were purchased from Cell Signaling Technology (Danvers, MA, USA). Glyceraldehyde 3-phosphate dehydrogenase (GAPDH; 60004-1-Ig) antibody was from the ProteinTech Group, Inc. (Chicago, IL, USA). Goat anti-rabbit IgG horse-radish peroxidase (HRP)-conjugated secondary antibody (A0208), goat anti-mouse IgG HRP-labeled secondary antibody (A0216), the cell counting kit-8 (CCK-8), ROS assay kit and mitochondrial membrane potential detection kit were from the Beyotime Institute of Biotechnology (Jiangsu, China). The EZ ChIP™ Chromatin Immunoprecipitation kit was purchased from Millipore Corp. (Billerica, MA, USA). Enhanced chemiluminescence (ECL) detection reagents were from Thermo Fisher Scientific (Waltham, MA, USA). Other chemical reagents were commercially available and analytically pure.

Cell viability. The viability of the H9c2 cells (in 96-well plates) was measured by CCK-8 assay according to the manufacturer's instructions. Briefly, the cells were exposed to H₂O₂ at various concentrations for different durations. CCK-8 solution (10 μ l) was added followed by further incubation for 2 h and the absorbance were measured at 450 nm using a microplate reader (Infinite M200 PRO; Tecan Corp., Mannedorf, Switzerland).

Measurement of myocardial infarct size (MIS). Following 0, 6, 12 and 24 h of reperfusion, the animals were euthanized and the hearts were rapidly removed. The MIS size was assessed by the 2,3,5-triphenyltetrazolium chloride (TTC) method as previously described (23). Briefly, the excised hearts were washed using 0.9% saline. The right ventricular tissues were removed and left ventricles were frozen and stored at -20°C for 5 min and then were sliced into 1-mm-thick sections, giving 5 slices of equal thickness along the long axis from the apex to the base. The slices were incubated for 15–30 min in TTC at pH 7.4 at 37°C to identify the viable myocardium stained red, while the infarct tissue remained white. The slices were fixed in 10% formaldehyde solution and photographed with a digital camera. Areas stained in red and white were measured by a BI-2000 Medical Image Analysis System (Chengdu TME Technology, Chengdu, China). The infarct size was calculated using the following equation: %Infarct volume = infarct volume/total left ventricle volume \times 100%.

Measurement of myocardial enzyme [creatine kinase (CK), lactate dehydrogenase (LDH) and aspartate aminotransferase (AST)] activities in serum. At the end of the experiment, blood samples were obtained from the abdominal aorta, incubated at 37°C for 30 min and then centrifuged at 3,000 rpm for 10 min. The supernatant from each group was collected and frozen at -80°C for further analysis. The activities of CK, LDH and AST were measured using CK, LDH and AST kits (Jiancheng Bioengineering Institute, Nanjing, China) with an automatic biochemistry analyzer (EOS880; Hospitex Diagnostics, Italy) according to the manufacturer's instructions.

Measurement of malondialdehyde (MDA) levels and SOD activities. At the end of the experiment, the myocardial tissue was homogenated for 5 min in the ice and then centrifuged at 2,000 rpm for 5 min. The supernatant from each group was collected for measuring the levels of MDA and the activities of SOD using respective kits (Jiancheng Bioengineering Institute).

Western blot analysis. Proteins were separated by 5-18% sodium dodecylsulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto polyvinylidene fluoride membranes. The membranes were blocked with 5% skimmed dry milk for 2 h at room temperature and washed 3 times in Tris-buffered saline with Tween (TBST); the membranes were then incubated with the primary antibodies to FoxO3a, SOD2, catalase and GAPDH overnight at 4°C. After washing with TBST again, HRP-labeled secondary antibodies goat anti-rabbit (1:1,000) or goat anti-mouse (1:2,000) were added followed by incubation for 1.5 h at room temperature. Immunoblots were developed by ECL detection kits and the bands were quantified by Quantity One software.

Immunofluorescence staining. The cultured H9c2 cells were fixed with 4% paraformaldehyde for 30 min and then washed 3 times in phosphate-buffered saline (PBS). Blocking solutions (0.2% Triton X-100 and sheep serum in PBS) were added to the fixed cells followed by incubation for 30 min at 37°C. The cells were then incubated with primary antibody against FoxO3a overnight at 4°C. After washing with PBS, the appropriate secondary antibody was added followed by incubation for 1 h in the dark at room temperature. The nuclei were then stained with Hoechst 33342 for 20 min. After washing with PBS, immunofluorescence was examined under a fluorescence microscope.

Measurement of ROS production. Intracellular ROS of H9c2 was measured by 2',7'-dichlorodihydrofluorescein diacetate (DCFH-DA) as a fluorescent probe according to the manufacturer's instructions. Briefly, H9c2 cells were loaded with DCFH-DA (10 μ M) for 30 min at 37°C and washed 3 times with serum-free medium. The fluorescence image was observed under a fluorescence microscope.

Detection of mitochondrial membrane potential ($\Delta\psi$ m). The dye, JC-1, was used as a fluorescent probe to detect $\Delta\psi$ m as previously described (24). CCCP (10 mM) provided in the kit was added to the cell culture medium at 1:1,000 for 20 min at room temperature, then the H9c2 cells were incubated with JC-1 dye (10 μ M) for 30 min at 37°C and washed twice with

JC-1 dyeing buffer (1X). The red and green fluorescence were observed under a fluorescence microscope.

Chromatin immunoprecipitation assay. Chromatin immunoprecipitation assay (25) was performed in accordance with the manufacturer's instructions with slight modifications. In brief, protein/DNA complexes of H9c2 cells (5×10^6) were cross-linked with formaldehyde and were sonicated to 200-1000 bp of DNA fragments. The crosslinked protein/DNA was then immunoselected with specific antibodies and the protein/DNA complexes were then eluted. Protein-DNA crosslinks were reversed during incubation at 65°C and DNA was purified to subject to quantitative PCR using the following primers: 5'-AAAGGCATCCCAAGGTAT-3' (forward) and 5'-CCA GCTCTACAGCTCGTAC-3' (reverse).

Statistical analyses. Data are shown as the means \pm SD. Analysis of variance (ANOVA) was used to compare quantitative variables between more than 2 groups. Intergroup differences were assessed by the Chi-square test and Student's t-test for quantitative variables. A p-value <0.05 (two-sided) was considered statistically significant. Data were analyzed with GraphPad Prism 5.0 software.

Results

Oxidative stress induces myocardial injury. In order to observe the effects of oxidative stress on H9c2 rat myocardial cells, H_2O_2 was used as an inducer. The results from CCK-8 assay revealed that cell viability decreased following exposure to H_2O_2 in a dose- and time-dependent manner. The cell survival rate at H_2O_2 400 μ M for 2 h was $53.96 \pm 5.6\%$ (Fig. 1A), compared with the control group, and the cell viability was significantly decreased ($p < 0.01$). Subsequently, intracellular ROS levels were detected using the DCFH-DA probe; the strength of the green fluorescence directly reflects intracellular ROS. The levels of ROS increased gradually following exposure to H_2O_2 compared with the untreated group, and the fluorescence intensity was significantly increased at 2 h (Fig. 1B).

FoxO3a is downregulated in oxidative stress-induced myocardial injury. To investigate the role of FoxO3a under myocardial oxidative stress conditions, we performed the experiments *in vitro* and *in vivo*. We examined the expression of FoxO3a in the H9c2 cells exposed to H_2O_2 at 400 μ M for different periods of time. A significant downregulation in FoxO3a expression was observed following exposure of the cells to H_2O_2 at 400 μ M for 2 h (Fig. 2A). We also detected the expression of FoxO3a in rats subjected to MI and 30 min of ischemia, followed by 0, 6, 12 or 24 h of reperfusion. The expression of FoxO3a was also decreased in the myocardial tissue in the rats with I/R injury (Fig. 2B).

TSA attenuates myocardial injury mediated by oxidative stress. To examine the effects of TSA on myocardial injury mediated by oxidative stress, we performed the experiments *in vitro* and *in vivo*. We measured the MIS by the TTC method. Compared with the I/R group, the MIS was significantly reduced in the TSA 0.2 mg/kg (T1)- and 0.1 mg/kg (T2)-treated groups, from 53.36 to 15.89% ($p < 0.01$) and 20.06% ($p < 0.001$) respec-

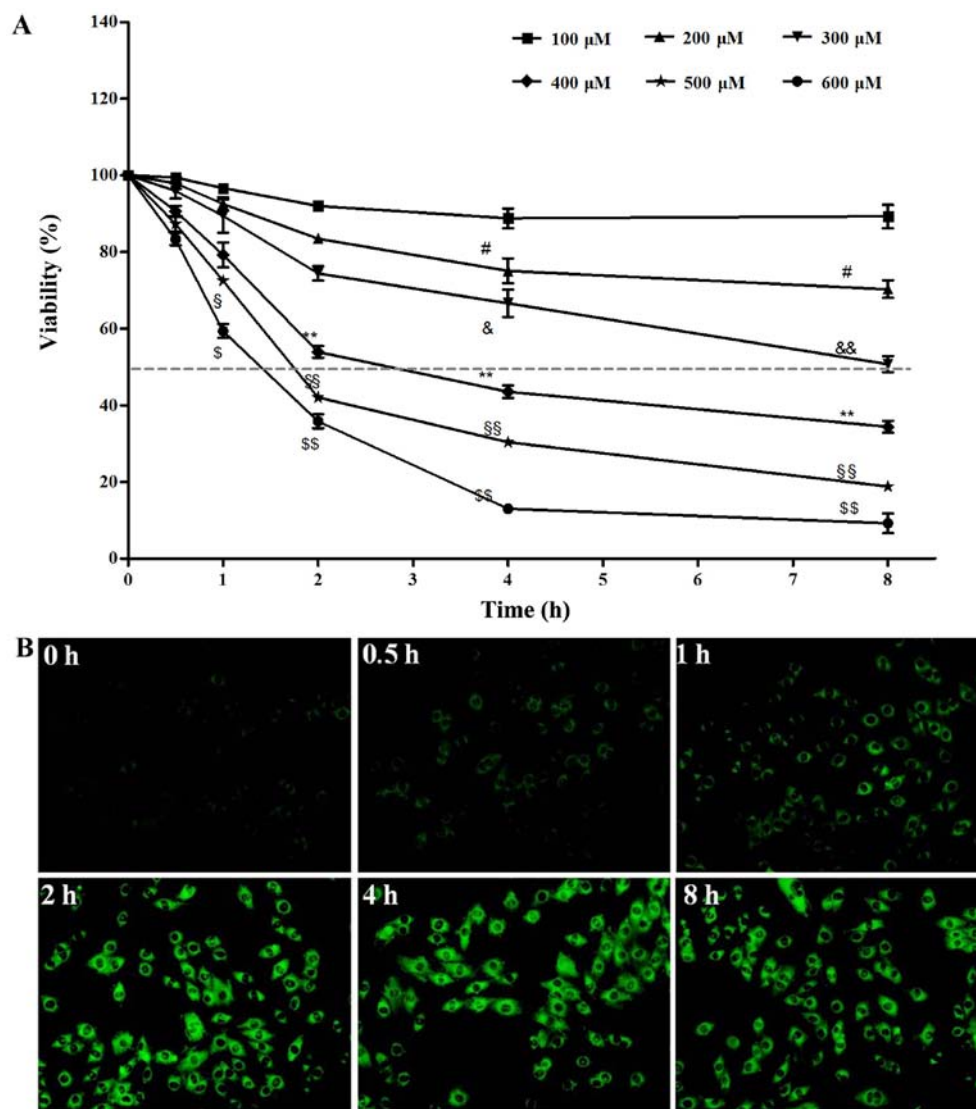


Figure 1. Oxidative stress induces myocardial injury. (A) Viability of H9c2 cells exposed to H_2O_2 at various concentrations for different periods of time are shown. Data are the means \pm SD; #, &, §, $p < 0.05$ and &&, **§§, $p < 0.01$ vs. untreated cells. (B) Images of reactive oxygen species (ROS) production by H9c2 cells exposed to H_2O_2 (400 μ M) are shown (magnification, $\times 200$).

tively; however, the MIS between the I/R and TSA 0.05 mg/kg (T3)-treated groups was not significantly altered (Fig. 3A). Subsequently, in order to evaluate the extent of myocardial injury in the rats, we measured the activities of CK, LDH and AST in serum after 24 h of reperfusion. In accordance with MIS, TSA significantly decreased the activities of CK, LDH and AST in serum (Fig. 3B). In addition, the MDA level was decreased and the SOD activity was increased in the TSA-treated groups compared with those in the sham-operated group (Fig. 3C).

In order to investigate whether TSA reduces the ROS levels following exposure of the H9c2 cells to H_2O_2 , intracellular ROS production was detected by the DCFH-DA probe. Compared with the H_2O_2 -exposed group, treatment with TSA reduced the intracellular ROS level (Fig. 3D). It has been reported that oxidative stress induces mitochondrial dysfunction (26). Thus, in this study, to investigate the protective effect of TSA against oxidative stress-induced mitochondrial damage, $\Delta\psi_m$ in the H9c2 cells was assessed using the JC-1 probe. The $\Delta\psi_m$ of the H9c2 cells declined markedly following exposure to H_2O_2 ,

whereas treatment with TSA increased the $\Delta\psi_m$ in the H9c2 cells (Fig. 3E).

Effect of TSA on the expression of FoxO3a. FoxO3a is an important regulator of the resistance to oxidative stress and the downregulation of FoxO3a increases cell death in response to oxidative stress in human chondrocytes (27). Resveratrol, a potent antioxidant, has previously been demonstrated to protect photoreceptor cells from apoptosis by upregulating the FoxO family in experimental retinal detachment (28). Our experimental results revealed that the expression of FoxO3a in the H9c2 cells was markedly reduced in the presence of H_2O_2 compared with that of the control group ($p < 0.01$). However, TSA elicited a significant increase in FoxO3a expression compared with the H_2O_2 alone-treated group ($p < 0.01$) (Fig. 4A and B). In line with the *in vitro* experimental results, TSA increased the expression of FoxO3a following 24 h of reperfusion in rats (Fig. 4C; $p < 0.01$ vs. sham-operated group; $p < 0.01$ vs. I/R group).

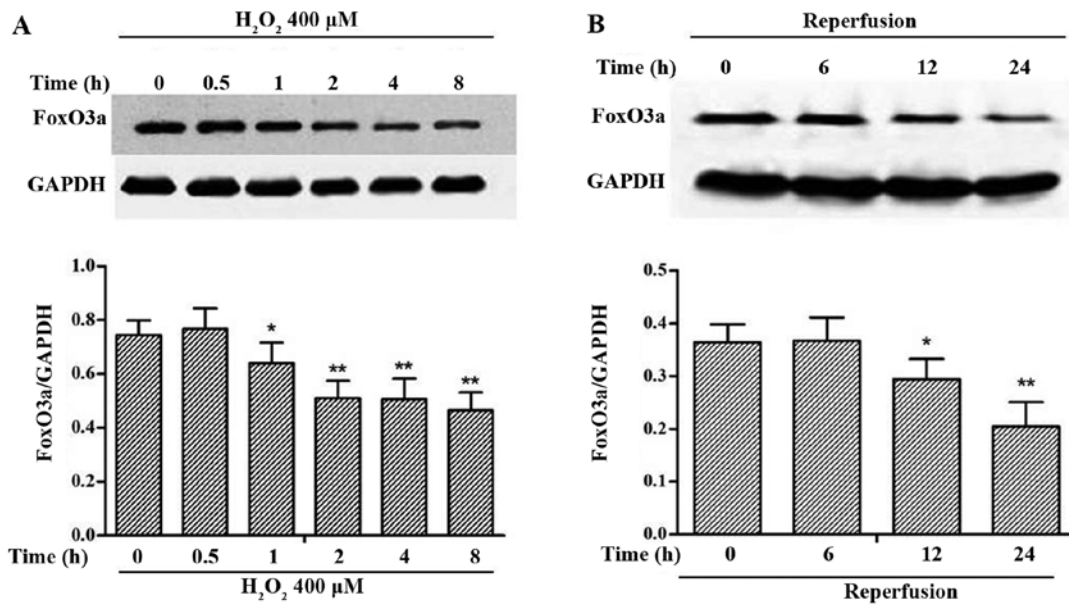


Figure 2. Forkhead box O3a (FoxO3a) is downregulated in oxidative stress-induced myocardial injury. (A) Effects of exposure to 400 μ M H_2O_2 for 0, 0.5, 1, 2, 4 and 8 h on the expression of FoxO3a in H9c2 cells. (B) FoxO3a expression after 30 min of ischemia followed by 0, 6, 12 and 24 h of reperfusion in the rats. Data are the means \pm SD; * p <0.05 and ** p <0.01 vs. 0 h.

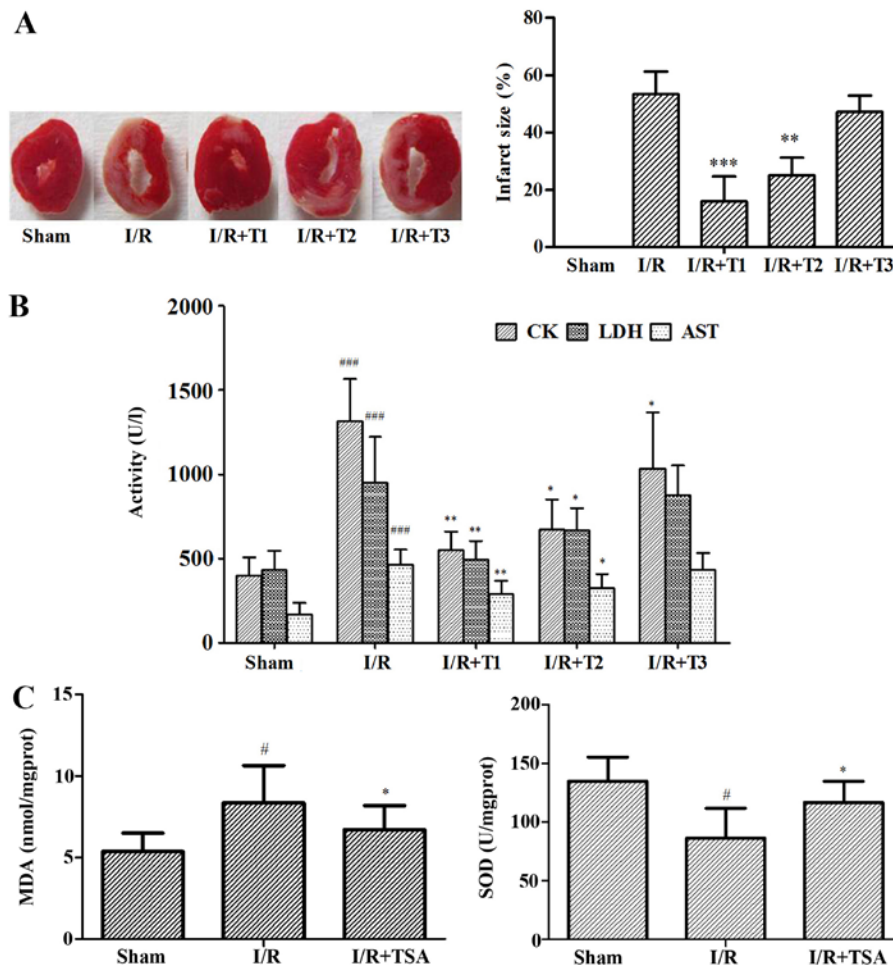


Figure 3. Trichostatin A (TSA) attenuates myocardial injury mediated by oxidative stress. (A) The myocardial infarct size was assessed by TTC. Rat hearts were subjected to 30 min of ischemia followed by 24 h of reperfusion (n=8) and graphs show the myocardial infarct size (infarct volume/left ventricular volume). ** p <0.01 and *** p <0.001, vs. I/R group. (B) The activities of CK, LDH and AST in the serum of rats whose hearts were subjected to 30 min of ischemia followed by 24 h of reperfusion were measured (n=8). T1, TSA 0.2 mg/kg; T2, TSA 0.1 mg/kg; and T3, TSA 0.05 mg/kg. Data are the means \pm SD; * p <0.05 and ** p <0.01 vs. ischemia/reperfusion (I/R) group; *** p <0.001 v. sham-operated (sham) group. (C) The levels of malondialdehyde (MDA) and the activities of superoxide dismutase (SOD) of rats whose hearts were subjected to 30 min of ischemia followed by 24 h of reperfusion were measured (n=8). Data are the means \pm SD; * p <0.05 and vs. I/R group; # p <0.05, vs. sham group.

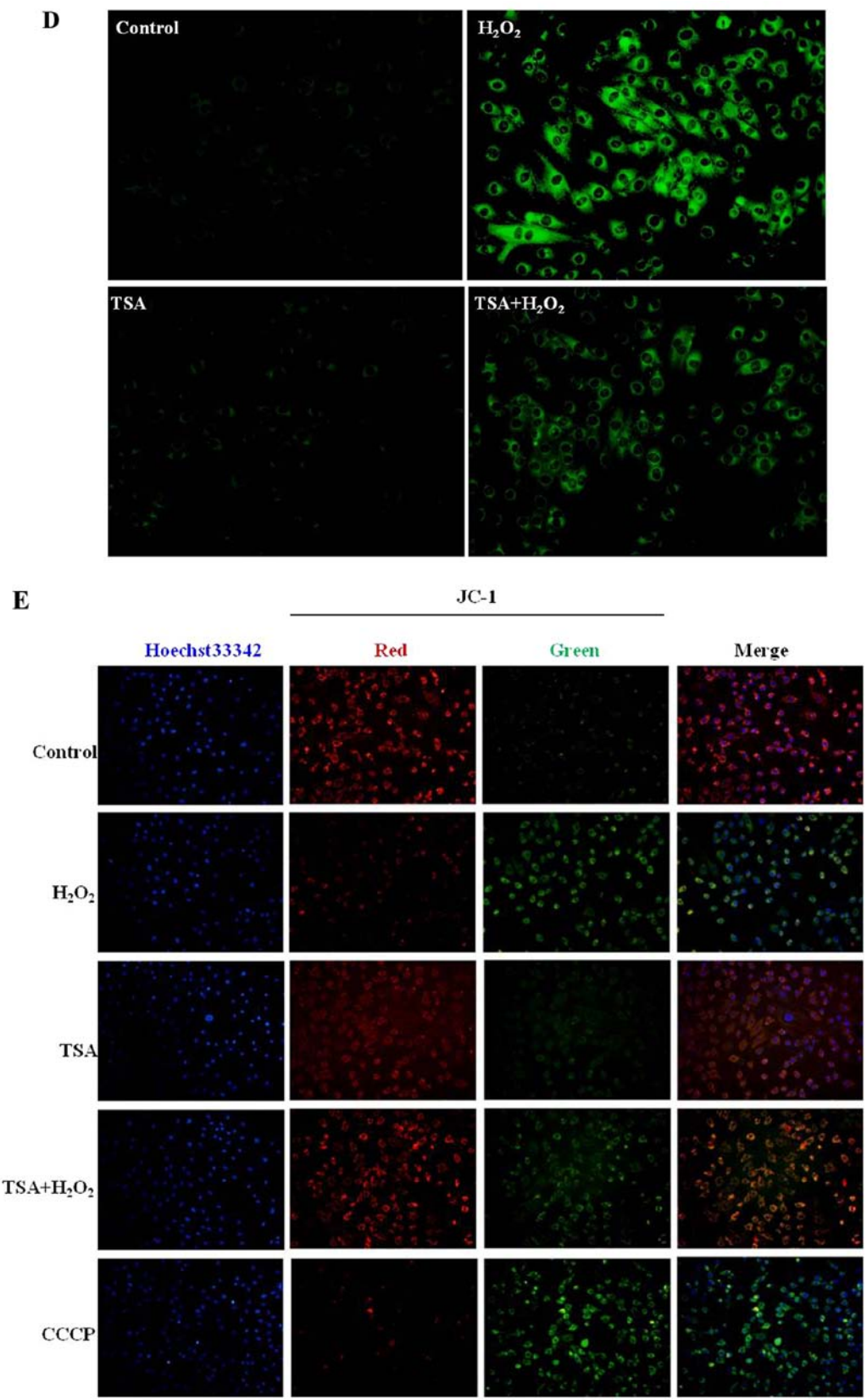


Figure 3. Continued. (D) Reactive oxygen species (ROS) production was evaluated by the DCFH-DA probe following exposure or not of the H9c2 cells to 400 μ M H₂O₂ for 2 h in the absence or presence of 50 nmol/l TSA in (magnification, x200). (E) $\Delta\psi_m$ was assessed by using JC-1 probe in H9c2 cells using JC-1 staining treated with/without 400 μ M H₂O₂ for 2 h in the absence or presence of 50 nmol/l TSA in H9c2 cells are shown (magnification, x100), CCCP was used as a positive control.

Effect of TSA on SOD2 and catalase levels. Mitochondrial SOD and catalase as endogenous enzymes are known to be the target proteins of FoxO3a that contributes to their protective effects against oxidative stress (29,30). The levels of SOD2 and

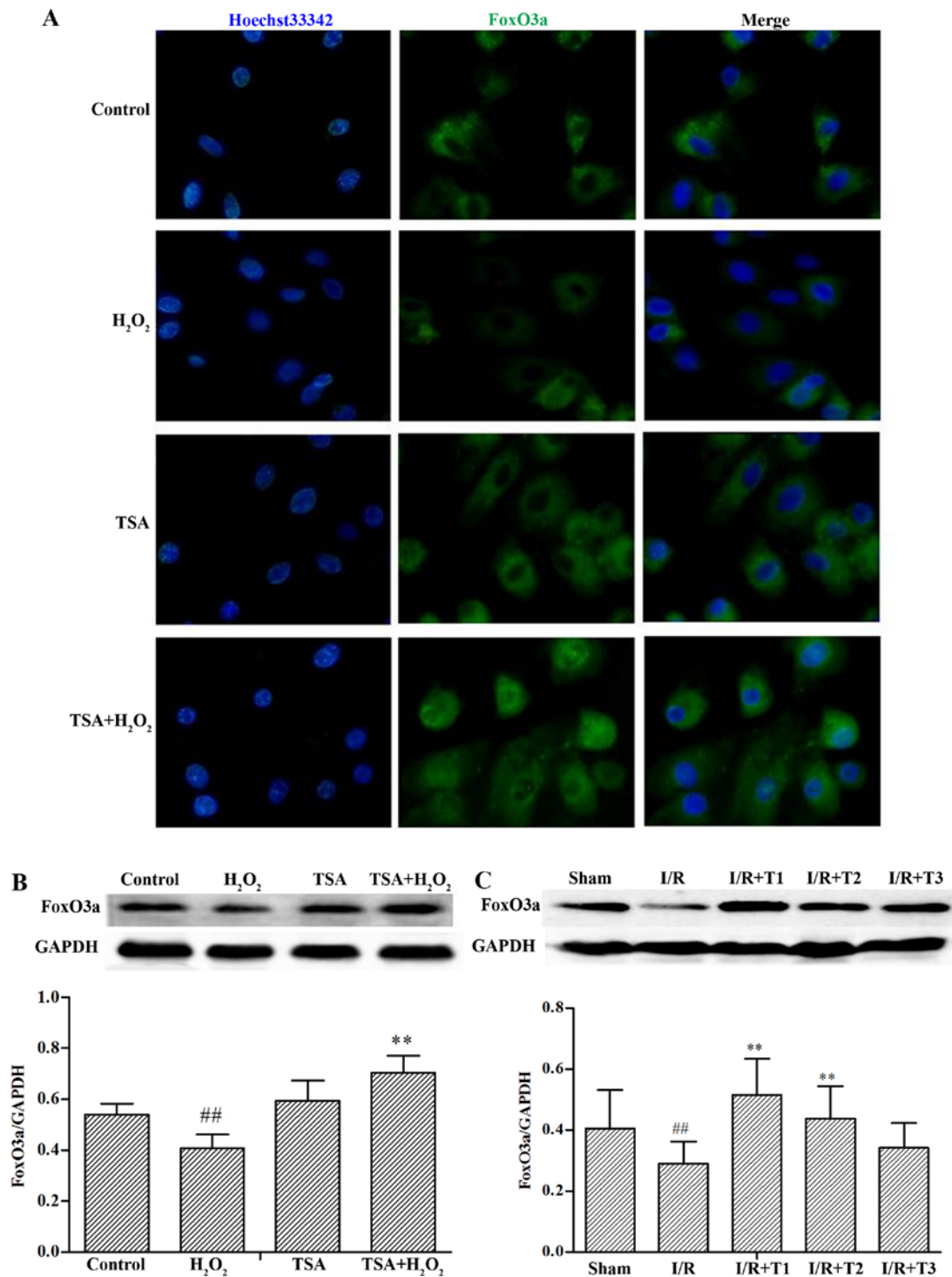


Figure 4. Effect of trichostatin A (TSA) on the expression of forkhead box O3a (FoxO3a). (A and B) Using immunofluorescence and western blot analysis to detect the expression of FoxO3a in H9c2 cells. FoxO3a was significantly upregulated following treatment with 50 nmol/l TSA in the absence or presence of H₂O₂. Data are the means \pm SD; ^{##}p<0.01 vs. control; ^{**}p<0.01 vs. H₂O₂. (C) TSA upregulated the expression of FoxO3a after ischemia for 30 min and 24 h of reperfusion in the rats. Data are the means \pm SD; ^{##}p<0.01 vs. sham-operated (sham) group; ^{**}p<0.01 vs. ischemia/reperfusion (I/R) group.

catalase in the H9c2 cells markedly decreased following exposure to H₂O₂, whereas treatment with TSA elicited a significant increase in SOD2 and catalase levels in the H9c2 cells (p<0.01) compared with cells exposed to H₂O₂ alone (Fig. 5).

Effects of TSA on the acetylation levels of histone H3 and H4 in the promoter region of FoxO3a. To investigate whether TSA

affects the levels of histone acetylation of the FoxO3a gene promoter and then FoxO3a gene expression, we performed *in vitro* experiments to observe the changes in the H3 and H4 acetylation levels of FoxO3a. The H4 acetylation of the FoxO3a promoter region was markedly decreased in the H₂O₂-exposed cells compared with that of the control group (p<0.01), whereas TSA elicited a significant increase in H4 acetylation of the

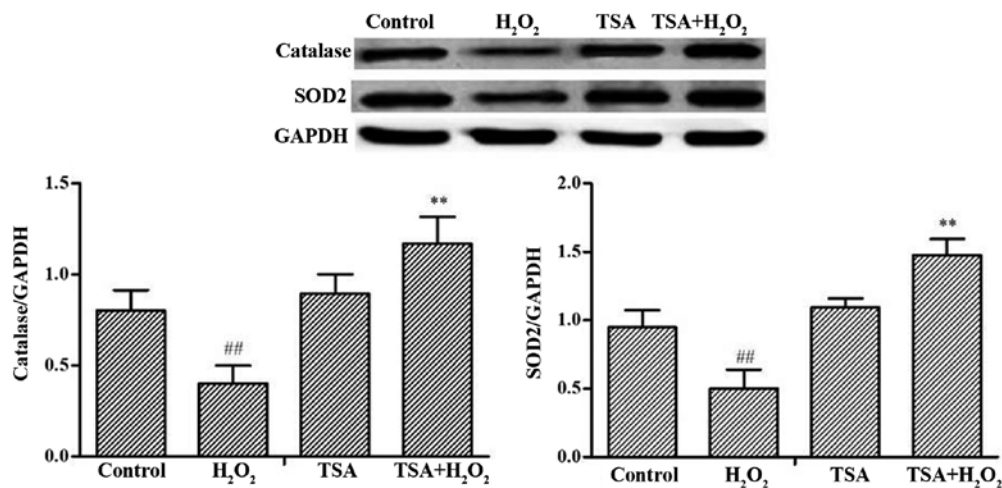


Figure 5. Effect of Trichostatin A (TSA) on superoxide dismutase 2 (SOD2) and catalase. Protein expression of SOD2 and catalase treated with/without 400 μ M H₂O₂ for 2 h in the absence or presence of 50 nmol/l TSA in H9c2 cells are shown. Mean \pm SD; [#]p<0.05 and ^{##}p<0.01 vs. control; ^{**}p<0.01 vs. H₂O₂.

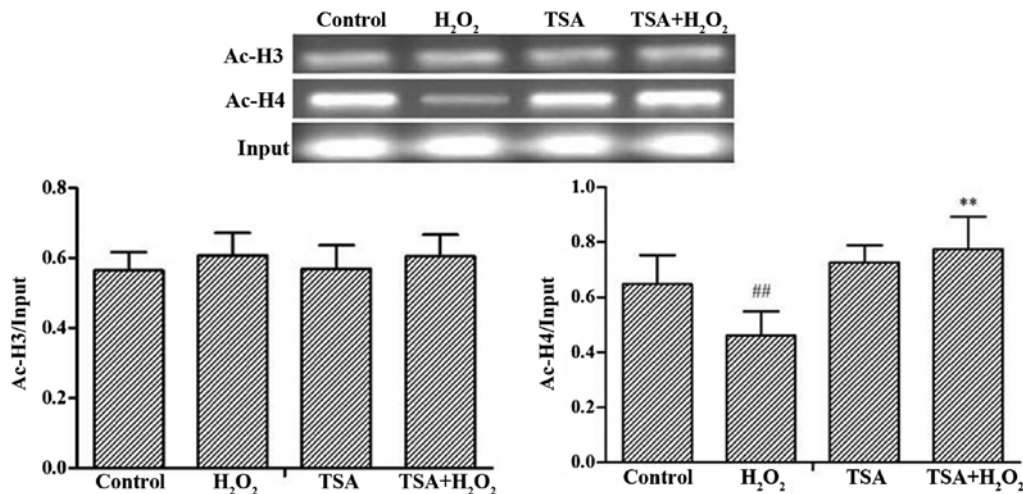


Figure 6. Effect of trichostatin A (TSA) on the acetylation levels of histone H3 and H4 in the promoter region of the forkhead box O3a (*FoxO3a*) gene. The acetylation levels of histone H3 and H4 in the promoter region of the *FoxO3a* gene following exposure or not to 400 μ M H₂O₂ for 2 h in the absence or presence of 50 nmol/l TSA in H9c2 cells are shown. Data are the means \pm SD; [#]p<0.01 vs. controls; ^{**}p<0.01 vs. H₂O₂-exposed group.

FoxO3a promoter region compared with that of the H₂O₂ group. The level of H3 acetylation of FoxO3a was not significantly altered (Fig. 6). The above-mentioned results thus indicate that TSA alters the H4 acetylation of the FoxO3a promoter region, thus affecting the expression of FoxO3a.

Discussion

The present study demonstrated that TSA attenuates myocardial injury mediated by oxidative stress *in vivo* and *in vitro*. This attenuation may be associated with the increased histone acetylation in the FoxO3a promoter region and the upregulation of the expression of FoxO3a, as well as the target proteins (SOD2 and catalase), which regulate the expression of ROS scavengers in response to ROS.

Oxidative stress is an important pathophysiological mechanism in myocardial injury (31). Oxidative stress means that

the body suffers various harmful stimulations, disrupting the balance between the generation and clearance of ROS; the overproduction of ROS as byproducts by the mitochondria results in myocardial injury (32,33). In this study, exposure to H₂O₂ at various concentrations for different periods of time in the H9c2 cells decreased cell viability in a dose- and time-dependent manner (Fig. 1A). Intracellular ROS levels in the H9c2 cells exposed to H₂O₂ for different periods of time increased gradually (Fig. 1B). This may be associated with the downregulated expression of FoxO3a (Fig. 2). Depending on the different environment and cell types, FoxO3a plays an important role in lifespan extension, energy metabolism and resistance to oxidative stress (34,35). FoxO3a also acts as the potential target for the treatment of several types of diseases (10). It has been demonstrated that FoxO3a plays an important role in maintaining cardiac function and antagonizing oxidative stress responses (36). Increasing the expression of FoxO3a reduces

ROS and promotes cardiomyocyte survival (20). FoxO3a also upregulates the expression of ROS scavengers to clear ROS (30,37). Mitochondria are considered to be important cellular organelles as the source of ROS (38), and $\Delta\psi_m$ is an important index maintaining mitochondrial function (39). A decrease in the $\Delta\psi_m$ results in mitochondrial dysfunction, causing an increase in ROS generation. MDA, which is the final product of lipid peroxidation, can reflect the production of oxygen free radicals; the content can not only reflect the formation of oxygen free radicals *in vivo*, it can also reflect the severity of tissue damage caused by oxygen free radicals. SOD, which is the specific scavenger, can reduce free radical damage to myocardial cells. SOD plays an important role in the free radical scavenging system. The present study revealed that the downregulation of FoxO3a in H9c2 cells was associated with the H_2O_2 -induced increase in intracellular ROS levels, and reduced $\Delta\psi_m$, and SOD2 and catalase levels. The opposite was observed following treatment with TSA (Figs. 3D and E, 4A and B and 5). Consistent with the experimental results obtained *in vitro*, TSA significantly reduced the MIS (Fig. 3A) and decreased the activities of serum CK, LDH and AST (Fig. 3B) in the rats. Moreover, TSA decreased the MDA level and increased SOD activity in the rats (Fig. 3C). These data suggest that TSA attenuates myocardial I/R-induced oxidative stress-related damage and upregulates the expression of FoxO3a, which is a possible mechanism ensuring the protective effects of TSA against I/R-induced myocardial injury (Fig. 4C).

Histone acetylation plays a pivotal role in the epigenetic modulation of gene expression and maintains a dynamic balance by affecting the activities of histone acetyltransferases and histone deacetylases (40). In most cases, the acetylation of transcription factors results in increasing their transactivation functions, principally by enhancing their DNA binding ability (41) and it has been shown that p300, an acetyltransferase, enhances the transcriptional activity of FoxO3a (42), reminding us that histone acetylation participates in FoxO3a regulation. Similar to a previous study showing that the acetylation level of the FoxO3a promoter region can be upgraded by HDAC inhibitors (21), we demonstrated that TSA reduced oxidative stress-induced myocardial injury by increasing the level of H4 acetylation of the FoxO3a promoter region, thereby enhancing the expression of FoxO3a (Fig. 6).

In conclusion, our data indicate that treatment with TSA attenuates oxidative stress-mediated myocardial damage by regulating the level of histone H4 acetylation in the promoter region of FoxO3a, and upregulating the expression of FoxO3a, SOD2 and catalase. Our findings provide a novel therapeutic strategy for TSA acting as a mediator of resistance to oxidative stress for myocardial protection.

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