ORIGINAL RESEARCH

Ginsenoside Rg1 ameliorates cardiac oxidative stress and inflammation in streptozotocin-induced diabetic rats

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Background and purpose: Ginsenoside Rg1 (GS Rg1), as an important active substance of Panax ginseng, has been proven to have elaborate cardioprotective effects. The purpose of this study was to detect that GS Rg1 attenuates cardiac oxidative stress and inflammation in streptozotocin (STZ)-induced diabetic rats (DM).

Methods: Cardiac function was assessed by heart rate and blood pressure. Markers relevant to myocardial oxidative stress and antioxidant capacity, and inflammatory reaction factors were detected. The mRNA and protein expression were detected by RT-qPCR and Western blot, respectively.

Results: GS Rg1 treatment significantly reduced the symptoms of cardiac hypertrophy and hypertension, and also decreased oxidative stress, inflammation response, NF-κB expression and NLRP3 inflammasome expression. GS Rg1 enhanced mitochondrial biogenesis by increasing PGC-1α, complex III and complex Ⅳ expression. GS Rg1 treatment significantly increased the expression of AMPK, Nrf2 and HO-1 in cardiac tissues.

Conclusion: GS Rg1 exhibited protective effect against STZ-induced cardiac dysfunction, which is potentially associated with AMPK/Nrf2/HO-1 signal pathway.

Keywords: Ginsenoside Rg1, oxidative stress, inflammation, AMPK, Nrf, HO-1

Introduction

At present, diabetes has become a global problem that threatens human health.[1] As the incidence of diabetes increases, the complications of diabetes have caused considerable concern. Diabetic cardiomyopathy is considered to be a disease of myocardial structural and functional abnormality independent of hypertension, coronary heart disease and related heart disease.[2] Among them, the occurrence of cardiovascular complications can easily lead to high morbidity and mortality in diabetic patients. At the same time, the risk of developing coronary heart disease and congestive heart failure increases in diabetic patients.[3] The risk of developing congestive heart failure events in diabetic patients is much higher than in non-diabetics.[4] According to clinical features, the main feature of diabetic cardiomyopathy is myocardial dilatation or hypertrophy, while left ventricular systolic or diastolic function is impaired. However, the underlying mechanisms for the pathogenesis of diabetic cardiomyopathy are not completely understood.

Oxidative stress, ammoniation, myocardial fibrosis, apoptotic cell death and other molecular mechanisms can consider to be used as potential factors in the pathogenesis of diabetic cardiomyopathy.[5] For oxidative stress, activation of 5′ adenosine

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phosphate-activated protein kinase (AMPK) plays a key role in improving mitochondrial activity and regulating cell metabolism.[6] Recently, studies have indicated that AMPK can suppress oxidative stress by stimulating upregulation of nuclear factor erythrocyte 2 related factor 2 (Nrf2)-dependent blood oxygenase (HO-1), and significant crosstalk is observed in mammalian inflammatory systems.[7] Therefore, we hypothesized that dysfunction of AMPK/ Nrf2/HO-1 signaling pathway might play an important role in streptozotocin (STZ)-induced cardiac oxidative stress and inflammation.

GS Rg1 is an active ingredient from the Panax noto-ginseng, known for its cardioprotective effects.[8] Currently, many studies have reported the protective effect of GS Rg1, such as inhibiting cardiovascular cell apoptosis and reducing oxidative stress.[9] Moreover, most of the diseases studied are mainly hypertrophic cardio-myopathy, acute myocardial infarction, myocardial ische-mia/reperfusion.[10–12] However, few studies have demonstrated the myocardial protective effect of GS Rg1 by decreasing oxidative stress and inflammatory response in diabetic rats. Therefore, in the current study, we observed the effects of GS Rg1 on oxidative stress and inflammatory response in STZ-induced DM rats.

Materials and methods
Animal experimental
This study was approved by the Ethics Committee of Qingdao University. Male Wistar rats (200±20) g were purchased from laboratory animal center of Qindao University and randomly divided into diabetic group and control group. The rats of diet group were administered intraperitoneally with STZ (40 mg/kg body weight). The rats of control rats were administered with 0.1 M citrate buffer. One week after the administration of STZ, blood glucose concentration >300 mg/dL were considered diabetic.

The STZ-induced DM rats subsequently apportioned randomly and equally to a DM control group (administered saline, ip), GS Rg1 group (20 mg/kg/day, ip) and AMPK inhibitor compound C group (CC, 20 mg/kg/day, ip). The assigned treatments were administered for 8 weeks. Based on previous studies, we use GS Rg1(20 mg/kg) as pharmacological approach for the administration of diabetic-induced cardiomyopathy.[13,14] The rats in the non-diabetic control group were normal, and these rats served as non-diabetic controls in the following experiment. Body weight of rats was measured weekly.

Heart rate and blood pressure measurement
Heart rate and blood pressure were measured using a non-invasive tail-cuff method (Narco BioSystem, Houston, TX, USA) in all rats. The rats were placed at 37°C for 10 mins at room temperature and then placed in an acrylic restrainer. The mean arterial pressure (MAP) was read and recorded. Finally, the average value of six individuals is counted as final value.

Preparation of myocardial tissue and blood samples
After 8 weeks of treatment, all rats were sacrificed under an overdose of 10% chloral hydrate (700 mg/kg body weight). When blood sample was obtained from the heart of the rat, 10% chloral hydrate (350 mg/kg body weight) was used to anesthetize rats. Then, 3 mL of blood samples was collected for subsequently determining the levels of blood glucose, lactate dehydrogenase (LDH), creatine kinase (CK)-MB and aspartate aminotransferase (AST). Subsequently, rat heart tissues were rapidly dissected and frozen at −80°C until RT-qPCR and Western blot. The myocardial hypertrophy index (ratio of heart weight/body weight) was determined.

Detection of markers relevant to myocardial oxidative stress and antioxidant capacity
The supernatants of myocardial tissues from each group were evaluated for levels of thiobarbituric acid reactive substances (TBARS), hydrogen peroxide, peroxynitrite, superoxide dismutase (SOD), catalase, GPx and glu-tathione peroxidase (GSH) using commercial kits (Cayman Chemicals, USA). ROS in cardiac homogenates were estimated fluorometrically.[15]

Determination of inflammatory reaction factors
Inflammatory reaction factors (IL-1β, IL-6 and TNF-α) were detected using ELISA kits (Thermo Fisher Scientific, USA) in heart homogenate and plasma, respectively.

cDNA synthesis and quantitative real-time PCR
Total RNA (1 μg) was extracted from the cardiac tissue using TRIzol reagent (Takara, Tokyo, Japan). Using 1 μg total

RNA, first-strand cDNA was synthesized by the AMV enzyme in 20 μL reaction mixture (Takara, Tokoyo, Japan). Using 2 μL reverse transcriptase products, quantitative PCR was performed in a final volume of 20 μL using gene-specific primers. The primers sequence is listed in Table 1. Amplification was carried out as follows: 95°C, 30 s, 1 cycle; 95°C, 5 s and 60°C 1 min for 40 cycles; subsequently, the melting curve was determined. Gene transcripts were quantified with SYBR Premix Ex Taq kit (Takara, Tokyo, Japan). The levels of gene expression were evaluated by quantitative real-time PCR with CFX96 system (Thermo Fisher Scientific, USA). The target gene expression level relative to the endogenous control gene (GAPDH) was calculated by $2^{-\Delta\Delta Ct}$ method.[16]

Analysis of relevant protein levels by Western blot

The cardiac tissues were harvested, washed three times in sterile saline and then homogenized in RIPA buffer (Beyotime Institute of Biotechnology, Suzhou, People's Republic of China) containing a protease inhibitor cocktail (Thermo Scientific, Rockford, IL, USA). Following centrifugation at 16,000 × g at 4°C for 30 mins, the supernatant was collected. Protein was quantified using a BCA Protein Assay kit (Beyotime Institute of Biotechnology),

then separated by 10% SDS-PAGE. Following transfer to a PVDF membrane (Beyotime Institute of Biotechnology), the primary antibodies were then incubated at 4°C overnight. Subsequently, they were incubated with the appropriate horseradish peroxidase-conjugated anti-rabbit (ZB-2301) secondary antibodies (ZSGB-BIO) for 1 hr at room temperature and visualized under a chemiluminescence system (Bio-Rad, USA). The bands were quantified using MultiGauge version 3.2 software. Relative expression levels of proteins were calculated by integrated grey values of the bands normalized with reference protein. The primary antibodies: AMPK (1:800; ab3759, Abcam, Boston, MA, USA), p-AMPK (Thr 172) (1:500; ab23875, Abcam), Nrf2 (1:800; ab137550, Abcam), HO-1 (1:1000; ab13243, Abcam), PGC-1α (1:800; ab54481, Abcam), NF-κB (1:800; ab131546, Abcam).

Statistical analysis

All experimental data were analyzed using SPSS 20.0 software (Chicago, IL, USA). Student's *t*-test was utilized as the comparison between the two groups. *P*<0.05 indicated that the data were significantly different.

Results

Effects of GS Rg1 treatment on MAP, heart rate, body weight and hypertrophy index in STZ-induced diabetic rats

As shown in Figure 1A, B, compared with control rats, the MAP, heart rate and hypertrophy index of DM rats were significantly increased (*p*<0.05). Figure 1C shows no change in the body weight of diabetic rats (*p*>0.05).

After 8 weeks of GS Rg1 injections in DM rats, we found that the MAP, heart rate and body weight of GS Rg1 treatment were significantly decreased (*p*<0.05; Figure 1A–C). Furthermore, GS Rg1 administration also remarkably decreased hypertrophy index of heart in DM rats (*p*<0.05; Figure 1D).

Effects of GS Rg1 on glycaemic control and myocardiac injury markers in STZ-induced diabetic rats

As depicted in Figure 1, the STZ administration significantly increased blood glucose concentrations (*p*<0.05), which was efficiently decreased by GS Rg1 (*p*<0.05; Figure 1E). Previous research has shown that serum enzyme activities of LDH, CK-MB and AST usually can be used as a marker to detect

Table 1 List of primer sequences

| Target gene		Primer sequence
HO-1	Forward Reverse	5′-CGTGCAGAGAATTCTGAGTTC-3′ 5′-AGACGCTTTACGTAGTGCTG-3′
Nrf 2	Forward Reverse	5′-GATTCGTGCACAGCAGCA-3′ 5′-GCCAGCTGAACTCCTTAGAC-3′
NF-κB	Forward Reverse	5′-GGGTCAGAGGCCAATAGAGA-3′ 5′-CCTAGCTTTCTCTGAACTGCAAA-3′
PGC-1α	Forward Reverse	5′-AGGTCCCCAGGCAGTAGAT-3′ 5′-CGTGCTCATTGGCTTCATA-3′
TLR4	Forward Reverse	5′-GTGGGTCAAGGACCAGAAAA-3′ 5′-GAAACTGCCATGTCTGAGCA-3′
NLRP3	Forward Reverse	5′-GCTGCTCAGCTCTGACCTCT-3′ 5′-AGGTGAGGCTGCAGTTGTCT-3′
ASC	Forward Reverse	5′-TTATGGAAGAGTCTGGAGCTGTGG-3′ 5′-AATGAGTGCTTGCCTGTGTTGG-3′
GAPDH	Forward Reverse	5′-TGATGACATCAAGAAGGTGGTGAAG-3′ 5′-TCCTTGGAGGCCATGTGGGCCAT-3′
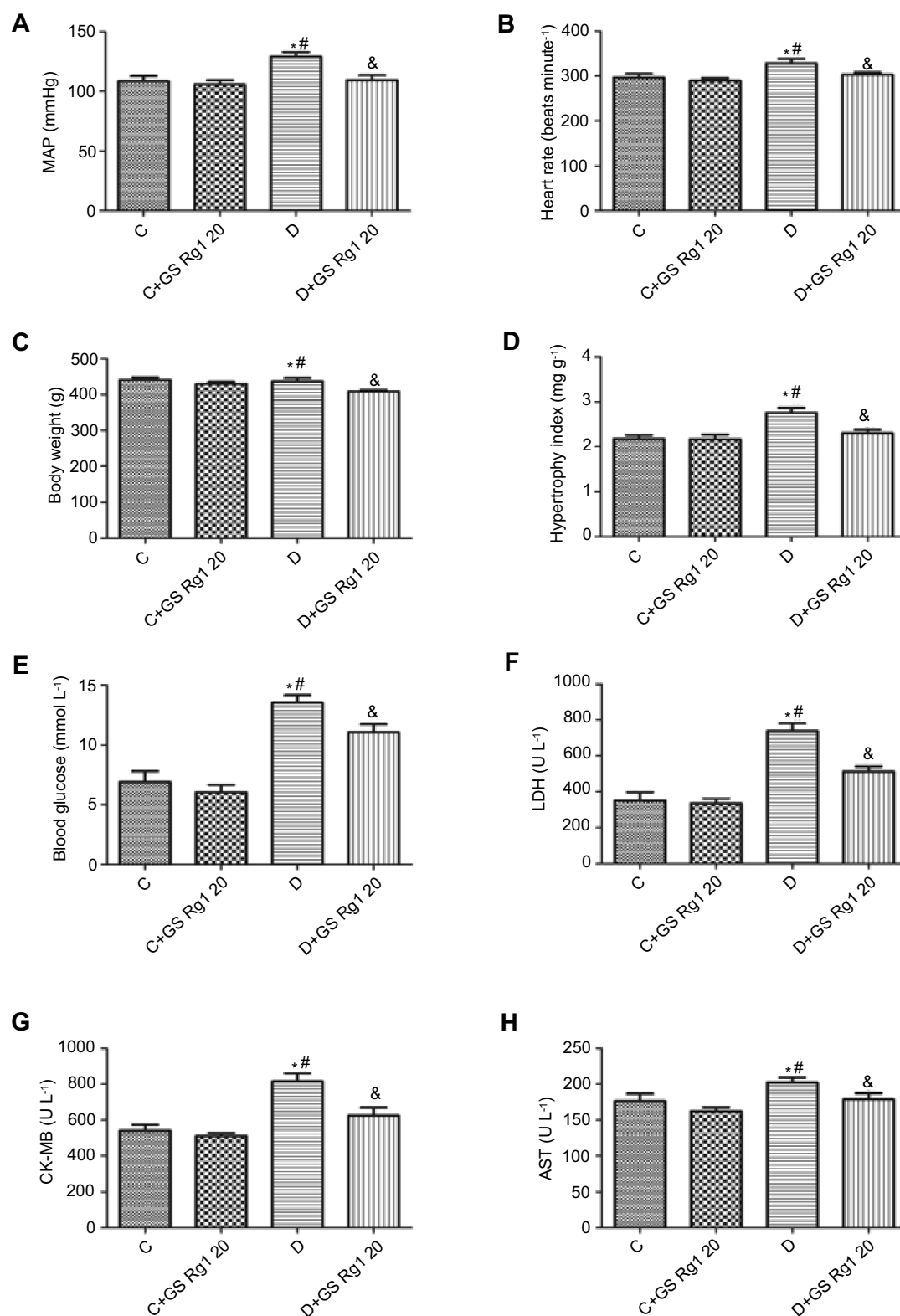
Figure 1 Effects of GS Rg1 on the indicators in the control and diabetic rats. (**A**) The MAP levels. (**B**) The heart rate. (**C**) The body weight. (**D**) The hypertrophy index. (**E**) The blood glucose levels. (**F**) The LDH levels. (**G**) The CK-MB levels. (**H**) The AST levels. *P*<0.05. (*) vs C; (#) vs C+GS Rg1 20; (&) vs D.
Abbreviations: MAP, mean arterial pressure; LDH, lactate dehydrogenase; CK-MB, creatine kinase-MB; AST, aspartate aminotransferase; C, control; D, STZ-induced diabetic; GS Rg1 20, ginseng Rg1 20mg/kg/day.

myocardial infarction.[16] Therefore, we tested the expression of the above indicators. The results are shown in Figure 1. Compared with control rats, LDH ($p<0.05$; Figure 1F), CK-MB ($p<0.05$; Figure 1G) and AST ($p<0.05$; Figure 1H) expression levels were significantly increased in the plasma in DM rats, which all were efficiently lessened by GS Rg1 treatment.

Effects of GS Rg1 on myocardial oxidative stress in STZ-induced diabetic rats

Studies have shown that oxidative stress is associated with diabetic cardiomyopathy.[17] Therefore, we tested the relevant indicators expression of cardiac oxidative stress. As shown in Figure 2, the expression levels of TBARS ($p<0.05$; Figure 2A), ROS ($p<0.05$; Figure 2B), hydrogen peroxide ($p<0.05$; Figure 2C) and peroxynitrite ($p<0.05$; Figure 2D) significantly increased in DM group rats compared with control group, suggesting that STZ administration significantly increased oxidative stress in the hearts of diabetic rats. On the contrary, GS Rg1 treatment remarkably reduced these oxidative stress markers in the myocardium of STZ-induced diabetic rats ($p<0.05$; Figure 2A-D).

Effects of GS Rg1 on antioxidant defense system in STZ-induced diabetic rats myocardium

Based on the results of the above oxidative stress indicators, we speculated whether GS Rg1 would affect the antioxidant defense system of STZ-induced diabetic rats. The results are shown in Figure 2. Compared with the rats of control group, total SOD ($p<0.05$; Figure 2E), catalase ($p<0.05$; Figure 2F), GSH ($p<0.05$; Figure 2G) and GPx ($p<0.05$; Figure 2H) expression levels were significantly decreased in the antioxidant defense system in STZ-induced diabetic rats, indicating STZ treatment significantly attenuated the role of the antioxidant defense system in the myocardium. However, GS Rg1 administrated significantly reversed these indicators of antioxidant defense system in STZ-induced diabetic rats ($p<0.05$; Figure 2E–H).

Effects of GS Rg1 on inflammation in STZ-induced diabetic rats

DM patients often have chronic inflammation. Therefore, to validate the anti-inflammatory effect of GS Rg1 in STZ-induced diabetic rats, the levels of inflammatory factors IL-1β, IL-6 and TNF-α were examined. As shown in Figure 3, compared with control group rats, the expression levels of IL-1 ($p<0.05$; Figure 3A), IL-6 ($p<0.05$; Figure 3B) and TNF-α ($p<0.05$; Figure 3C) were significantly elevated in the myocardial of STZ-induced diabetic rats. However, GS Rg1 treatment significantly attenuated the expression levels of IL-1β, IL-6 and TNF-α in the myocardium of STZ-induced diabetic rats (Figure 3). In addition, we also tested these inflammatory cytokines in the plasma. We found that a similar kind of results was observed in plasma inflammatory cytokines from GS Rg1 administered STZ-induced diabetic rats and control rats ($p<0.05$; Figure 3D–F).

Effects of GS Rg1 on myocardial Nf-κB expression and inflammasome in STZ-induced diabetic rats

As shown in Figure 4, vcompared with control group rats, the mRNA and protein expression of NF-κB in the myocardial were significantly increased in STZ-induced diabetic rats ($p<0.05$). On the contrary, GS Rg1 treatment significantly reduced mRNA ($p<0.05$; Figure 4A) and protein ($p<0.05$; Figure 4E) expression of cardiac NF-κB in DM rats whereas CC treatment enhanced the same. Meanwhile, co-administration with CC markedly reversed the inhibitory effect of GS Rg1 on NF-κB ($p<0.05$; Figure 4A, E). Next, we evaluated the expression of inflammasome components (NLRP3 and ASC) and its upstream activator (TLR4). The results were shown that the mRNA expression levels of TLR4 ($p<0.05$; Figure 4B), NLRP3 ($p<0.05$; Figure 4C) and ASC ($p<0.05$; Figure 4D) significantly increased in DM rats. GS Rg1 administrate effectively decreased mRNA expression of above gene, whereas CC markedly enhanced them ($p<0.05$; Figure 4B–D). However, co-administration with CC prevented the inhibitor effect of GS Rg1 on mRNA expression of inflammasome components.

Effects of GS Rg1 on mitochondrial biogenesis in STZ-induced diabetic rats

Studies have shown that oxidative stress is known to cause mitochondrial dysfunction.[18] As shown in Figure 5, compared with control group rats, the mRNA levels of PGC-1α ($p<0.05$; Figure 5A) and protein levels of myocardial PGC-1α ($p<0.05$; Figure 5B), complex III ($p<0.05$; Figure
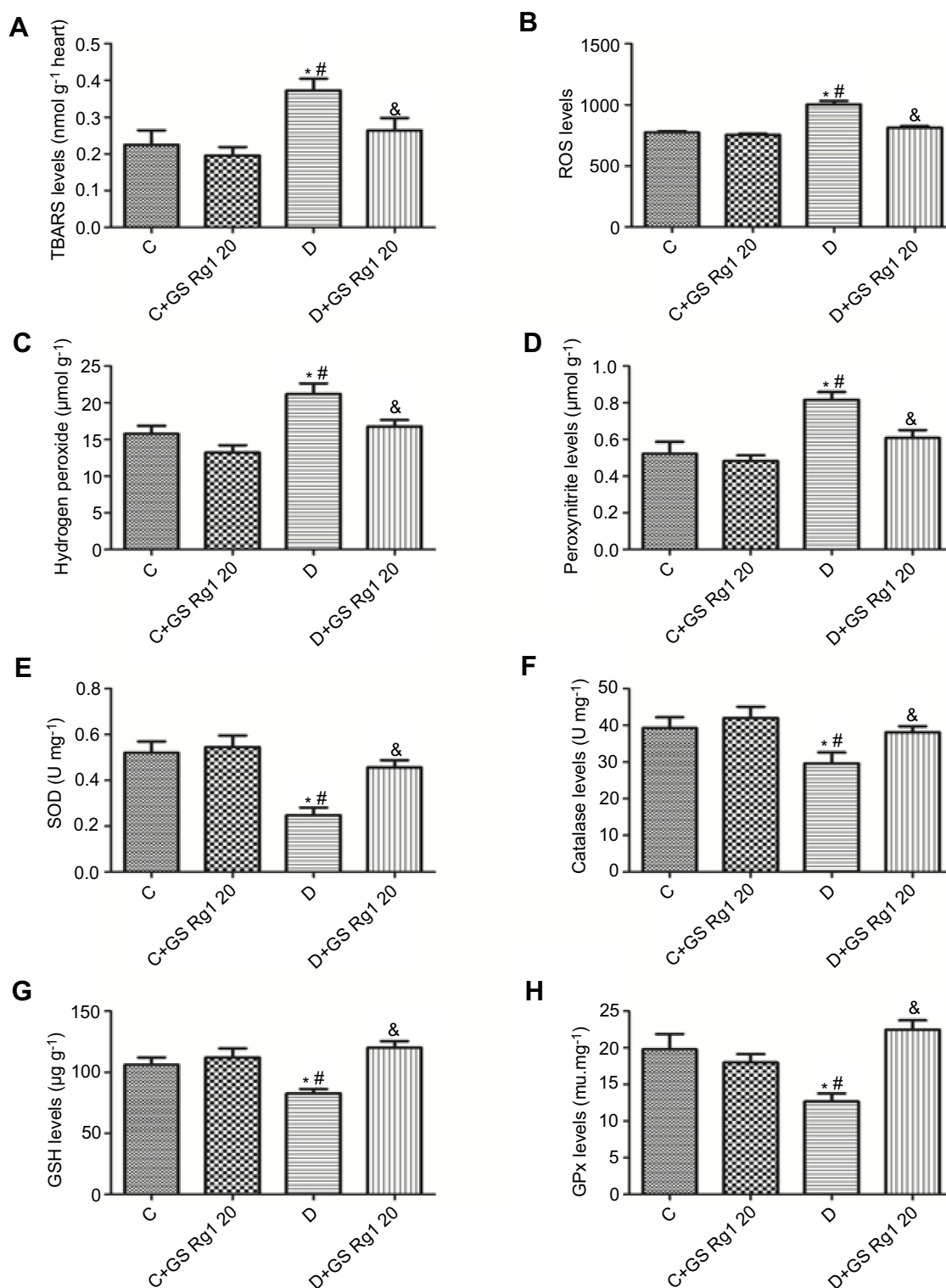
Figure 2 Effects of GS Rg1 on the cardiac oxidative stress and antioxidant defense system in control and diabetic rats. (**A**) Cardiac TBARS levels. (**B**) Cardiac ROS levels. (**C**) Cardiac hydrogen peroxide levels. (**D**) Cardiac peroxynitrite levels. (**E**) Myocardial SOD levels. (**F**) Myocardial catalase levels. (**G**) Myocardial GSH levels. (**H**) Myocardial GPx levels. $P<0.05$. (*) vs C; (#) vs C+GS Rg1 20; (&) vs D.
Abbreviations: C, control; D, STZ-induced diabetic; GS Rg1 20, ginseng 20mg/kg/day; TBARS, thiobarbituric acid reactive substances; ROS, reactive oxygen species; SOD, total superoxide dismutase; GSH, reduced glutathione; GPx, glutathione peroxidase.
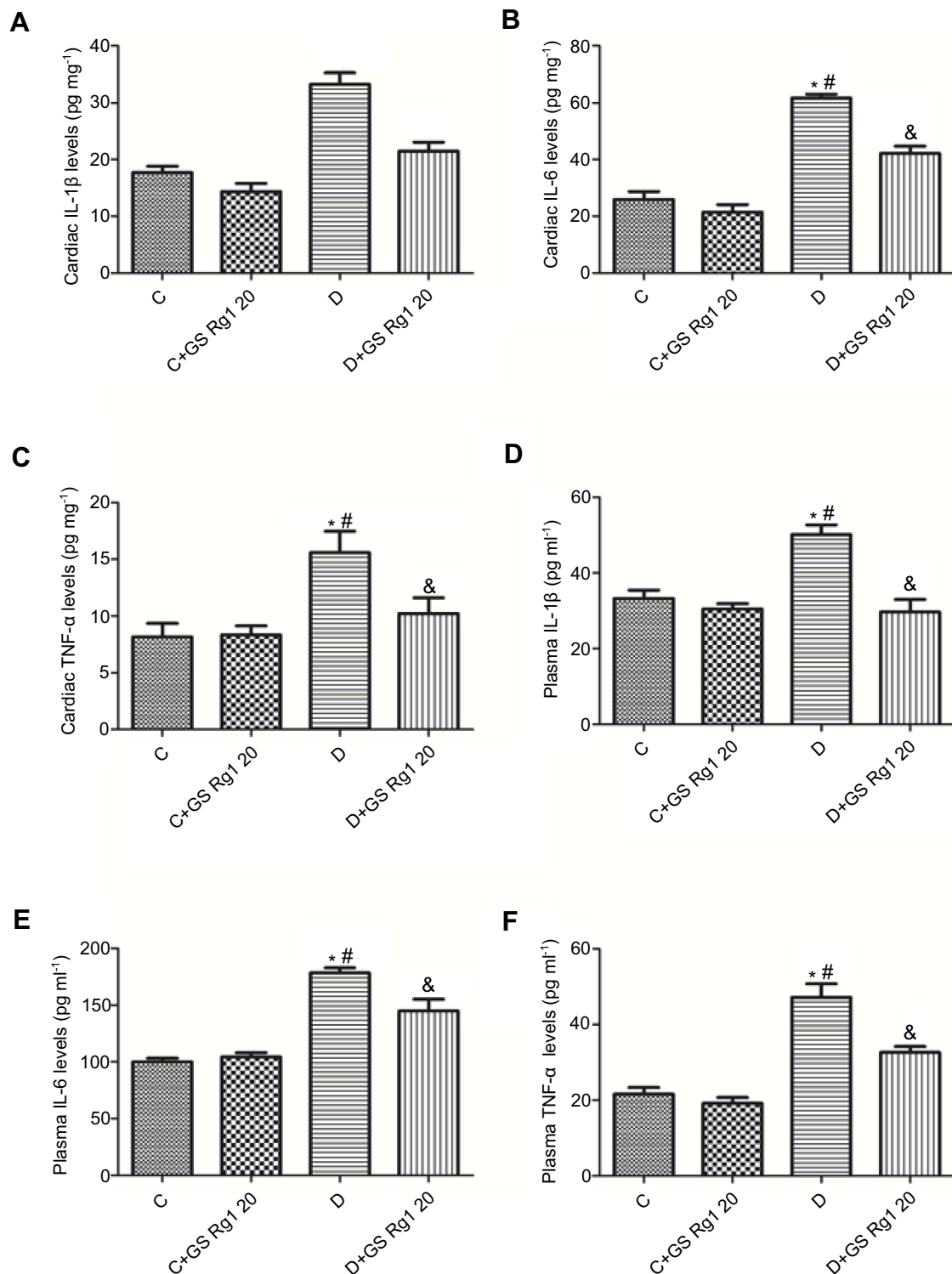
Figure 3 Effects of GS Rg1 supplementation on cardiac and plasma inflammatory cytokines of control and diabetic rats. (**A**) Cardiac IL-1β levels. (**B**) Cardiac IL-6 levels. (**C**) Cardiac TNF-α levels. (**D**) Plasma IL-1β levels. (**E**) Plasma IL-6 levels. (**F**) Plasma TNF-α levels. The significance was set at $P<0.05$. (*) vs C; (#) vs C+GS Rg1 20; (&) vs D.
Abbreviations: C, control; D, STZ-induced diabetic; GS Rg1 20, ginseng Rg1 20mg/kg/day; IL-1β, interleukin-1 beta; IL-6, interleukin-6; TNF-α, tumour necrosis factor-α.

5C) and complex IV ($p<0.05$; Figure 5D) in DM rats significantly decreased. However, GS Rg1 treatment significantly enhanced the same gene expression in DM rats, whereas CC markedly decreased the same gene levels (Figure 5). Meanwhile, co-administration with CC significantly reversed the effect of GS Rg1.
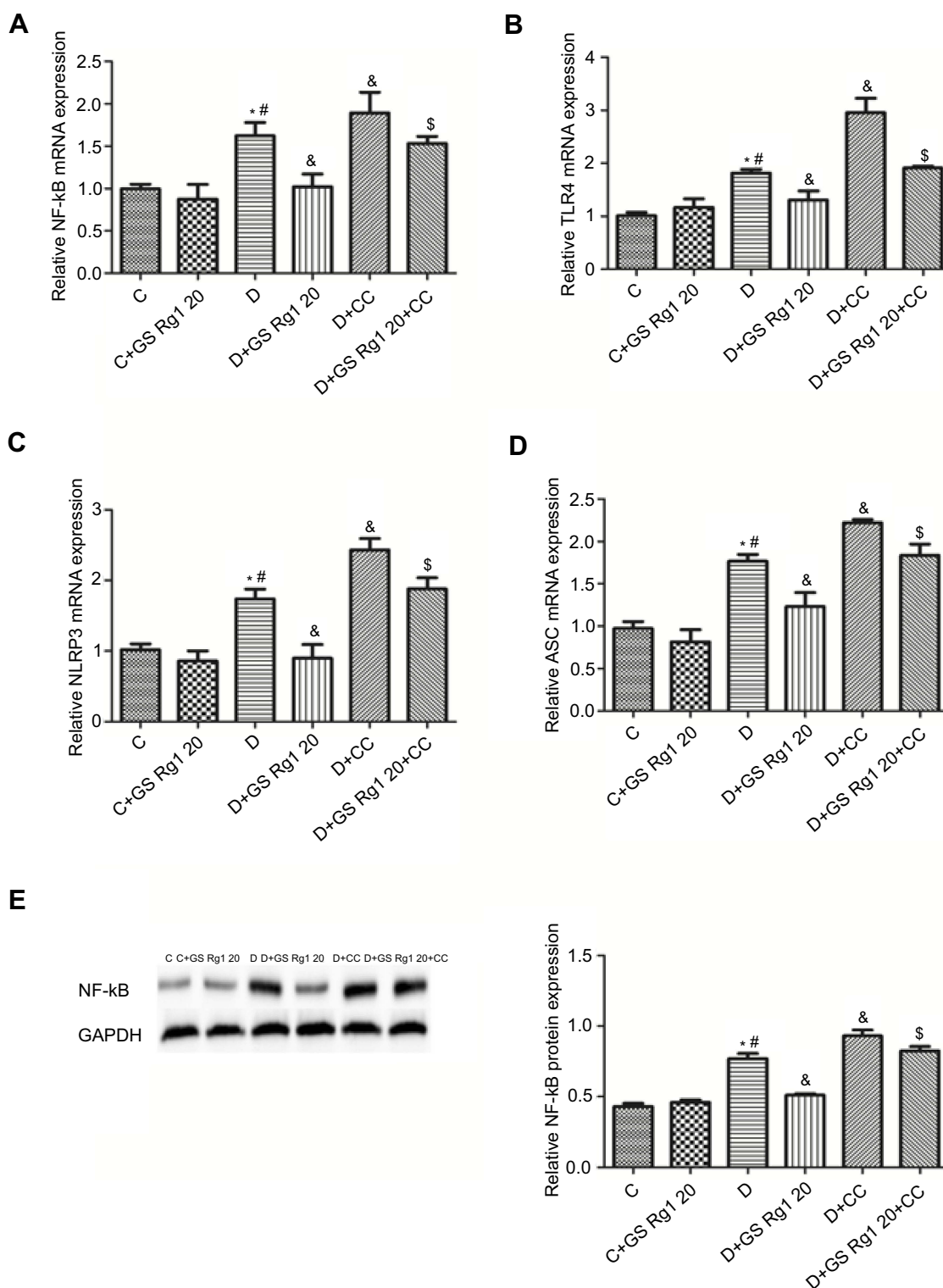
Figure 4 Effects of GS Rg1 treatment on inflammatory response in cardiac tissues of control and diabetic rats. (**A**) NF-κB mRNA expression. (**B**) TLR4 mRNA expression. (**C**) NLRP3 mRNA expression. (**D**) ASC mRNA expression. (**E**) NF-B protein expression. Values are represented as mean±SD. $P<0.05$. (*) vs C; (#) vs C+GS Rg1 20; (&) vs D; ($) vs D+GS Rg1 20.
Abbreviations: C, control; D, STZ-induced diabetic; GS Rg1 20, ginseng 20mg/kg/day; CC, compound C; NF-κB, nuclear factor κB; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.
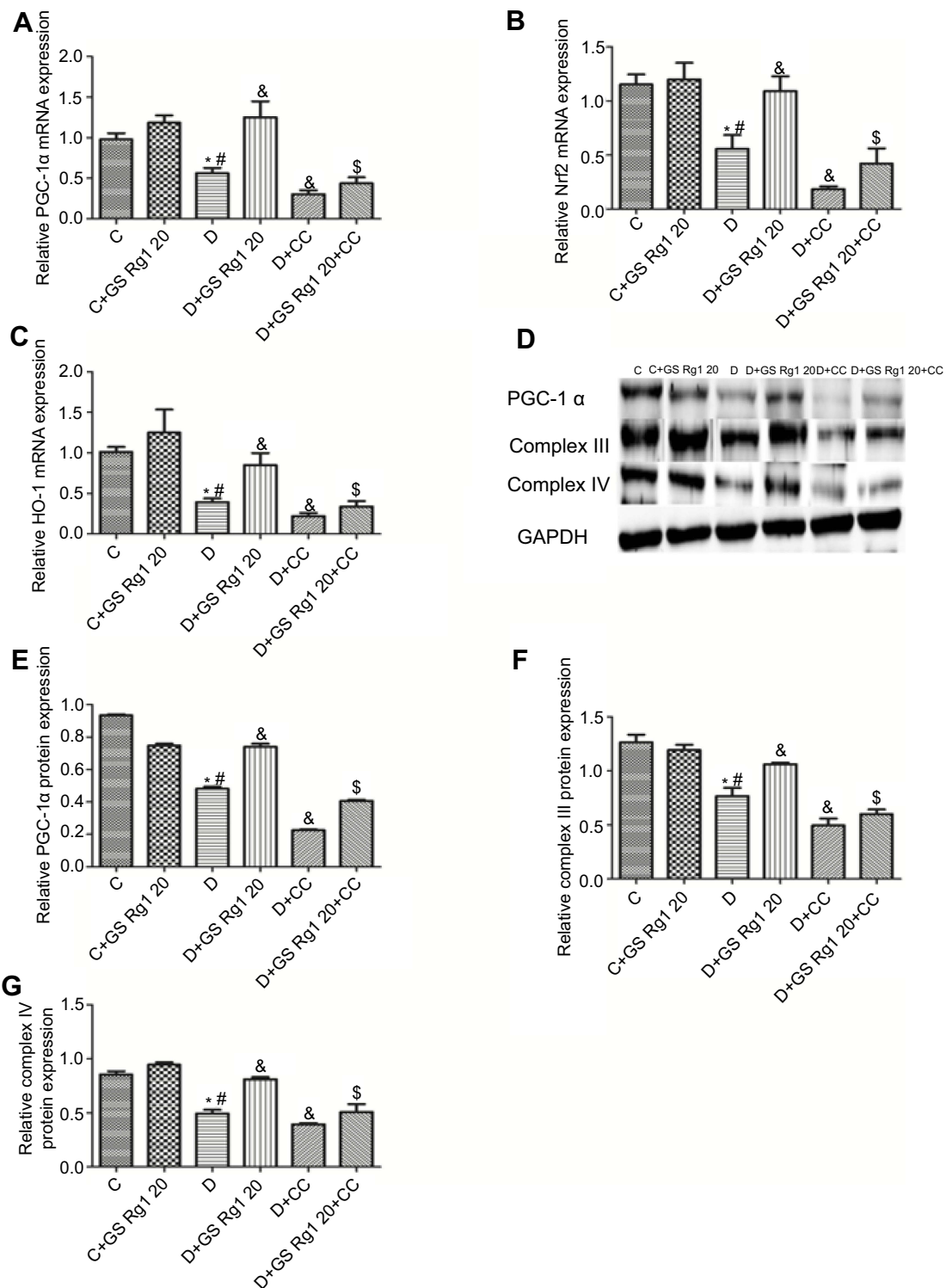
Figure 5 Effects of GS Rg1 administration on the expression of the mitochondrial biogenesis proteins, Nrf2, HO-1 in the hearts of control and diabetic rats. (**A**) mRNA expression of PGC-1α. (**B**) mRNA expression of Nrf2. (**C**) mRNA expression of HO-1. (**E**) protein expression of PGC-1α. (**F**) protein expression of complex III. (**G**) protein expression of complex V. $P<0.05$. (*) vs C; (#) vs C+GS Rg1 20; (&) vs D; ($) vs D+GS Rg1 20.

Abbreviations: C, control; D, STZ-induced diabetic; GS Rg1 20, ginseng 20mg/kg/day; CC, compound C; PGC-1α, peroxisome proliferator-activated receptor gamma coactivator; Nrf2, nuclear factor erythroid 2-related factor 2; HO-1, hemeoxygenase; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.
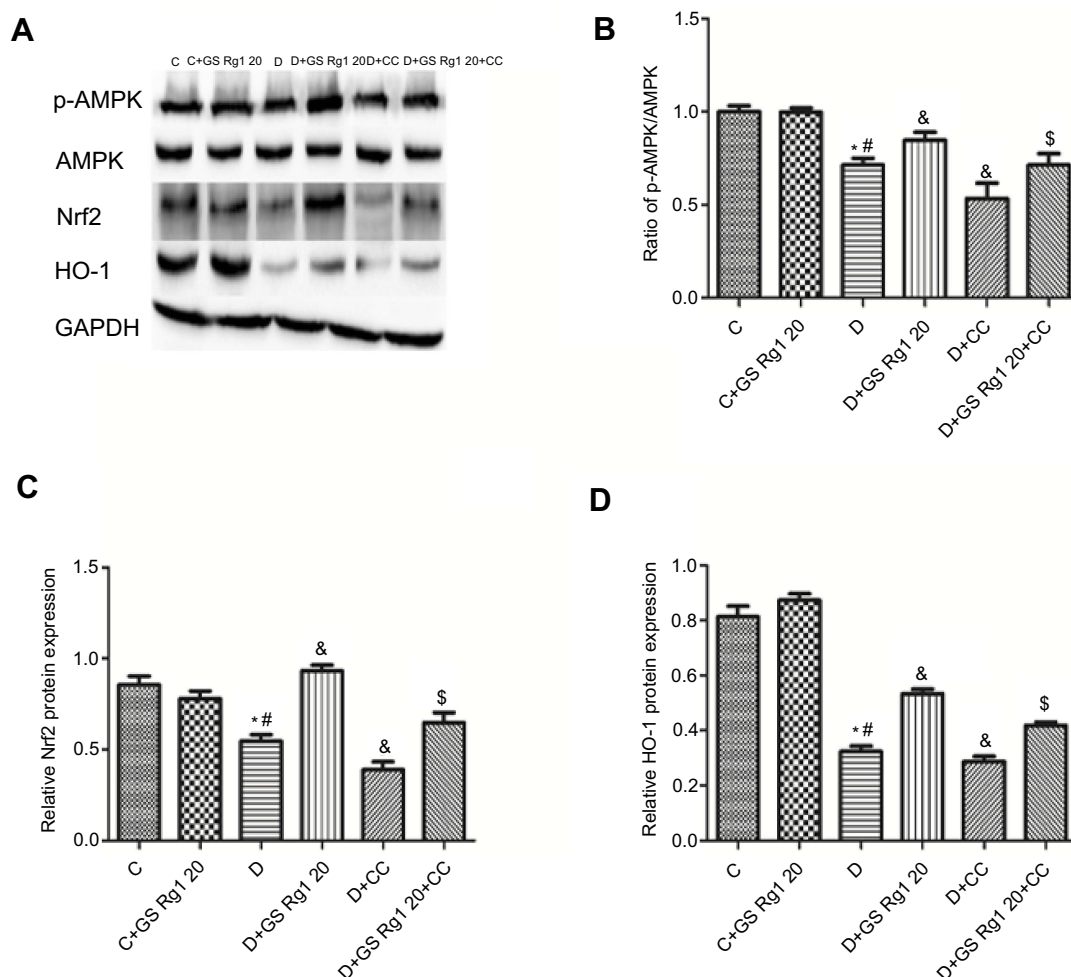
Figure 6 Effects of GS Rg1 administration on the protein expression levels of AMPK/Nrf2/HO-1 signaling components in the hearts of control and diabetic rats. Myocardial expression of (**A**) AMPK, *p*-AMPK; (**B**) Nrf2 and (**C**) HO-1. Values are represented as mean±SD. *P*<0.05. (*) vs C; (#) vs C+GS Rg1 20; (&) vs D; ($) vs D+GS Rg1 20.
Abbreviations: C, control; D, STZ-induced diabetic; GS Rg1 20, ginseng 20mg/kg/day; CC, compound C; Nrf2, nuclear factor erythroid 2-related factor 2; HO-1, hemeoxygenase; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

Effects of GS Rg1 on Nrf2 and HO-1mRNA expression levels in cardiac tissues of STZ-induced diabetic rats

As shown in Figure 6, compared with control group rats, the mRNA expression of Nrf2 (*p*<0.05; Figure 6A) and HO-1 (*p*<0.05; Figure 6B) significantly decreased in the cardiac tissue of DM rats. GS Rg1 administration significantly enhanced the Nrf2 and HO-1 mRNA expression in heart tissues of DM rats, whereas CC treatment significantly reduced Nrf2 and HO-1 mRNA expression (*p*<0.05; Figure 6A–B). However, co-administration with CC reversed the effect of GS Rg1.

Effects of GS Rg1 on AMPK/Nrf2/HO-1 signaling pathway in STZ-induced diabetic rats

As shown in Figure 6, we found that *p*-AMPK/AMPK ratio significantly decreased in DM rats myocardium, which was reversed after GS Rg1 administration (*p*<0.05; Figure 6C), whereas CC treatment significantly diminished *p*-AMPK/ AMPK ratio. However, co-administration with CC significantly blocked GS Rg1-mediated phosphorylation of AMPK (*p*<0.05; Figure 6C). Similarly, GS Rg1 administration significantly increased the Nrf2 and HO-1 expression levels in DM rats (*p*<0.05; Figure 6D–E). However,

co-administration with CC markedly reversed the effect of GS Rg1 ($p<0.05$; Figure 6D–E).

Discussion

In this study, we established a diabetes model with STZ 40 mg/kg injection. Then, we examined the relevant indicators of myocardium and determined that STZ administration can cause myocardial impairment.

Excessive production of ROS may exceed the activity of antioxidant enzymes, leading to the occurrence of oxidative stress. Growing evidence shows that excessive levels of superoxide radicals induced by hyperglycemia in diabetes can contribute to oxidative stress and changes in cardiac structure, which ultimately leads to diabetic myocardial damage.[17] Under normal physiological conditions, although ROS are continuously produced, the antioxidant defense system is sufficient to prevent ROS-related damage. Diabetic cardiomyopathy is an imbalance between ROS production and clearance, leading to oxidative stress being amplified by cascades and possibly leading to damage and apoptosis in normal tissues.[19] Hyperglycemia can increase the sensitivity of STZ-induced diabetic rats myocardial tissue to oxidative stress, mainly because the functional activities of SOD, catalase, GSH and GPX are decreased, and the antioxidant defense response is weakened. In this study, we used GS Rg1 administration to STZ-induced diabetic rats, found that GS Rg1 significantly reduced cardiac TBARS, ROS, hydrogen peroxide, peroxynitrite expression, and recovered cardiac antioxidant enzymes SOD, catalase, GPx and GSH levels. Therefore, the free radical scavenging properties and antioxidant effects of GS Rg1 contribute to the alleviation of STZ-induced myocardial oxidative stress.

It has been reported that hyperglycemia-induced myocardial tissue oxidative stress is associated with inflammation by overproduction of pro-inflammatory cytokines.[20] NF-κB is a pleiotropic transcription factor that controls the expression of multiple target genes and is primarily involved in inflammation. When NF-κB is blocked, oxidative stress and inflammatory responses in myocardial tissue are simultaneously reduced, and cardiac dysfunction in type 2 diabetic mice is alleviated.[21] In the heart, activation of NF-κB signaling is associated with different pathophysiological environments, including myocardial infarction, heart failure, cardiac hypertrophy and diabetic cardiomyopathy.[22] Our study found that the expression of NF-κB mRNA and protein was increased in STZ-induced diabetic rats, and GS Rg1 treatment decreased its expression, suggesting that GS Rg1 may inhibit NF-κB- mediated

cardiac inflammation. At the same time, studies have shown that NF-κB transactivation and Toll-like receptor 4 (TLR4)-mediated ROS production are mechanisms of inflammatory body initiation and regulation.[23] The NLRP3 pathway affects insulin sensitivity and increases myocardial cytokine levels as well as macrophage infiltration.[24] Moreover, NLRP3 inflammatory bodies regulate downstream inflammatory events of lipotoxicity and glucose toxicity during the development of T2 DM.[25] In the present study, we found that the expression of TLR4, NLRP3 and ASC in the heart of STZ-induced diabetic rats was significantly upregulated, while GS Rg1 reduced the inflammation of the NLRP3 inflammatory body. At the same time, surprisingly, after the combination of CC, we found that CC inhibited the inhibitory effect of GS Rg1 on myocardial inflammation in DM rats mediated by NF-κB and NLRP3 inflammatory neutrophils in an AMPK-dependent manner.

Previous studies have demonstrated that mitochondrial dysfunction promotes the production of oxidative free radicals, leading to cardiac oxidative stress in diabetic rats.[26] Overexpression of PGC-1α in the heart strongly enhances mitochondrial DNA content,[27] whereas knockdown of PGC-1α leads to decreased expression of the citrate cycle and phosphatidylation genes.[28] In our study, chronic GS Rg1 treatment enhanced the expression of PGC-1α, complex III and V subunits. At the same time, co-administration with CC, we found that CC significantly inhibited the promotion of GS Rg1 on myocardial mitochondrial biosynthesis in DM rats, indicating that GS Rg1 can be activated by AMPK pathway to improve mitochondrial biogenesis.

Next, we investigated the AMPK/Nrf2/HO-1 signal, a well-recognized metabolic stress sensor that reduces cardiac oxidative stress in the context of STZ-induced type 2 diabetes by GS Rg1 administration. Oxidative stress can cause AMPK phosphorylation, thereby stimulating the expression of Nrf2 and its downstream antioxidant enzymes, including HO-1.[29] In addition, activation of Nrf2 protects cardiomyocytes against doxorubicin-induced cardiomyopathy by inducing mitochondrial biosynthesis.[30] Surprisingly, Nrf2 also blocks the inflammatory response by inhibiting the transcription of pro-inflammatory cytokines.[31] Therefore, we believe that the Nrf2/HO-1 axis acts as a marker for antioxidant enzymes, and that GS Rg1 is found to be effective in increasing the expression of Nrf2, which activates AMPK in DM rats myocardium. To further reveal that the AMPK signaling pathway is involved in GS Rg1-induced cardioprotection effects,

AMPK inhibitors are given to diabetic rats with GS Rg1. CC combination therapy not only prevented GS Rg1 from inducing AMPK phosphorylation but also reduced the expression of Nrf2 and HO-1 in diabetic myocardium. AMPK stimulation regulates cellular energy content by increasing ATP generation through increasing mitochondrial pool via upregulation of mitochondrial biogenesis, which is requisite to restrain diabetes-triggered oxidative stress.[32] Meanwhile, p-AMPK may be involved in other pathway regulation. These results indicate that GS Rg1 promotes AMPK/Nrf2/HO-1 pathway to improve STZ-mediated myocardial oxidative stress and inflammation.

In conclusion, we have demonstrated that GS Rg1 administration has a beneficial effect on cardiac oxidative stress and inflammation in STZ-induced diabetic rats. The underlying mechanisms may be involved in AMPK/Nrf2/HO-1 pathway.

Ethical approval

All animal experiments were performed in accordance with the regulations of the Ethics Committee of the Affiliated Hospital of Qingdao University and were consistent with the Guidelines for the Care and Use of Laboratory Animals published by the US National Institutes of Health.

Disclosure

The authors declare that they have no conflicts of interest.

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