Pristimerin protects against doxorubicin-induced cardiotoxicity and fibrosis through modulation of Nrf2 and MAPK/NF-κB signaling pathways

Dina S El-Agamy¹,², Khaled M El-Harbì³, Saad Khoshhal³, Nishat Ahmed³, Mohamed A Elkablawy⁴,⁵, Ahmed A Shaaban³,⁶, Hany M Abo-Haded³,⁷

¹Department of Pharmacology and Toxicology, College of Pharmacy, Taibah University, Al-Madinah Al-Munawwarah 30001, Saudi Arabia; ²Department of Pharmacology and Toxicology, Faculty of Pharmacy, Mansoura University, Mansoura 35516, Egypt; ³Cardiogenetic Team, Department of Pediatrics, College of Medicine, Taibah University, Al-Madinah Al-Munawwarah 30001, Saudi Arabia; ⁴Department of Pathology, Faculty of Medicine, Taibah University, Al-Madinah Al-Munawwarah 30001, Saudi Arabia; ⁵Department of Pharmacology, Faculty of Pharmacy, Aqaba University of Technology, Aqaba 77110, Jordan; ⁶Pediatric Cardiology Unit, Department of Pediatrics, Faculty of Medicine, Mansoura University, Mansoura 35516, Egypt

Background/purpose: Pristimerin (Pris) is triterpenoid compound with many biological effects. Until now, nothing is known about its effect on doxorubicin (DOX)-induced cardiotoxicity. Hence, this study investigated the impact of Pris on DOX-induced cardiotoxic effects.

Materials and methods: Rats were treated with Pris 1 week before and 2 weeks contaminant with repeated DOX injection. Afterwards, electrocardiography (ECG), biochemical, histopathological, PCR, and Western blot assessments were performed.

Results: Pris effectively alleviated DOX-induced deleterious cardiac damage. It inhibited DOX-induced ECG abnormalities as well as DOX-induced elevation of serum indices of cardiotoxicity. The histopathological cardiac lesions and fibrosis were remarkably improved in Pris-treated animals. Pris reduced hydroxyproline content and attenuated the mRNA and protein expression of the pro-fibrogenic genes. The antioxidant activity of Pris was prominent through the amelioration of oxidative stress parameters and enhancement of antioxidants. Furthermore, Pris enhanced the activation of nuclear factor-erythroid 2 related factor 2 (Nrf2) signaling pathway as it increased the mRNA and protein expression of Nrf2 and Nrf2-dependent antioxidant genes (GCL, NQO1, HO-1). Additionally, the anti-inflammatory effect of Pris was obvious through the inhibition of mitogen activated protein kinase (MAPK)/nuclear factor kappa-B (NF-κB) signaling and subsequent inhibition of inflammatory mediators.

Conclusion: This study provides evidence of the cardioprotective activity of Pris which is related to the modulation of Nrf2 and MAPK/NF-κB signaling pathways.

Keywords: doxorubicin, pristimerin, cardiotoxicity, Nrf2, MAPK/NF-κB

Introduction

Doxorubicin (DOX) is a broad spectrum anthracycline antineoplastic agent that is used in various malignancies. Its antitumor action is mediated through inhibition of topoisomerase-II enzyme, DNA damage, prevention of DNA repair, and cell replication which lead to cell death in highly proliferative tissues.¹ Treatment with DOX has been associated with fatal cardiotoxic adverse effects which may range from pericarditis, transient controllable arrhythmias, and reversible hypotension observed immediately after treatment to congestive heart failure with mortality rate exceeding 50% in case of chronic treatment.² Chronic use of DOX produces cardiac fibrosis and remodeling. This is probably due to the imbalance between the synthesis and degradation of extracellular matrix (ECM) collagens. Alteration in ECM usually results in left ventricle dysfunction and progression of heart failure.³–⁵

The exact molecular mechanisms underlying DOX-induced cardiotoxicity are complex and still not fully clarified. Studies have proposed that oxidative stress,
inflammation, and apoptosis are the main players that control DOX pathogenesis. Excessive generation of free radicals during DOX metabolism augments oxidative stress, lipid peroxidation, membrane damage, and cardiomyocyte death.

The nuclear factor-erythroid 2 related factor 2 (Nrf2) is a transcription factor that controls the expression of enzymes involved in the anti-oxidative stress response and hence modulates the cellular defense pathway. Recent studies have shown that Nrf2 activation could counteract DOX-induced cardiotoxicity via suppression of DOX-induced oxidative stress and impairment of autophagy.

Activation of cellular inflammatory pathways is greatly involved in DOX-induced cardiac injury. DOX activates nuclear factor kappa-B (NF-κB) which results in pronounced inflammatory reactions and cellular damage. Inhibition of NF-κB pathway by different agents was shown to confer protection against DOX-induced cardiotoxicity. Even though DOX is still used, only few chemicals are adopted to protect the heart against deleterious adverse effects of DOX. Hence, efforts must continue to find new therapeutics that can oppose DOX-induced cardiotoxicity.

Various active compounds derived from medicinal plants have been considered as an attractive alternative approach for multiple organ dysfunction. Pristimerin (Pris) is a natural triterpenoid compound (Figure 1A). It is extracted from Celastraceae plants and has been used in Chinese traditional medicine because of its multiple biological properties. Studies have demonstrated that Pris can be an effective therapeutic for different tumors, and autoimmune and inflammatory disorders. The anti-inflammatory activity of Pris was initially linked to suppression of the release of nitric oxide (NOx) via inhibition of inducible nitric oxide synthase (iNOS). More recently, Pris was shown to inhibit NF-κB and its inflammatory cascade. Based on these data, we hypothesized that Pris might have potential protective activities against DOX-induced myocardial damage. The current study evaluated the effects of Pris treatment on DOX-induced cardiac injury and fibrosis in rats. Furthermore, the impact of Pris on different molecular pathways was assessed.

**Materials and methods**

**Drugs and reagents**

DOX was purchased as ampoules (Ebewe Pharma GmbH, Austria). Pris was purchased from Tocris Bioscience (Bristol, UK) and dissolved in dimethyl sulfoxide (DMSO). Ketamine was obtained as ampoules (Tekam; Hikma Pharmaceuticals, Amman, Jordan). Creatine kinase isoenzyme-MB (CK-MB) kit was obtained from ELITech, France. Lactate dehydrogenase (LDH) activity kit was purchased from Human Biochemica und Diagnostica GmbH (Wiesbaden, Germany). ELISA kits for troponin I (cTnI) and troponin T (cTnT) were purchased from Kamiya Biomedical Co (Tukwila, WA, USA). Hydroxyproline colorimetric assay kit was purchased from Abcam (Cambridge, UK). Malondialdehyde (MDA), superoxide dismutase (SOD), reduced glutathione (GSH) kits were purchased from Biodiagnostic Company (Giza, Egypt). ELISA kits for 4-hydroxynonenal (4-HNE), 8-hydroxy-2-deoxyguanosine (8-OHdG), and protein carbonyl (PC) were purchased from MyBioSource Inc. (San Diego, CA, USA). Nuclear extraction and NF-κB p65 activation kits were bought from Abcam. ELISA kits for NOx, tumor necrosis factor-α (TNF-α), and interleukin-6 (IL-6) were purchased from R&D Systems, Minneapolis, MN, USA. Total protein extraction kit was purchased from KeyGEN Biotech Co, Ltd (Nanjing, China). Bicinchoninic acid (BCA) Protein Assay Kit was purchased from Thermo Fisher Scientific Inc. (Waltham, MA, USA).

Primary antibodies were purchased from Abcam and Thermo Fisher Scientific Inc. for the following proteins: transforming growth factor-beta1 (TGF-β1), cat no ab92486, metalloproteinase-2 (MMP-2, cat no ab37150), metalloproteinase-9 (MMP-9, cat no ab38898), tissue inhibitor of metalloproteinase-1 (TIMP-1, cat no ab61224), Nrf2 (cat no ab137550), glutamate-cysteine ligase catalytic subunit (GCLc, cat no ab80841), NAD(P)H dehydrogenase (quinone) 1 (NQO1, cat no ab28947), heme-oxygenase-1 (HO-1, cat no ab13243), NF-κB p65 (cat no PA5-16545), p-NF-κB p65 (cat no PA5-37718), inhibitor of NF-κB kinase alpha (IKKα, cat no PA5-17803), phospho-IKKα (p-IKKα, cat no PA5-37524), phospho-inhibitor of NF-κB (p-IxBα, cat no PA5-38603), goat anti-rabbit IgG (H+L) secondary antibody, biotin (cat no 31820), mitogen activated protein kinase (MAPK) isoforms: p38 (p38 MAPK, cat no AHO1202), p-p38 MAPK (cat no MA5-15182), c-JUN N-terminal kinase (JNK, cat no LF-PA0047), phospho-JNK (p-JNK, cat no PA5-37698), extracellular signal-regulated kinase (ERK, cat no LF-MA0215), phospho-ERK (p-ERK, cat no 36-8800), and β-actin (cat no ab8227). Other chemicals were of high analytical purity.

**Animals**

Animal handling and use were in accordance with the guidelines of the Guide for the Care and Use of Laboratory Animal of the National Institutes of Health (NIH, Bethesda, MD, USA). The experimental protocol was approved by the Research Ethics Committee, Taibah University, Saudi Ara-
bia and Research Ethics Committee, Faculty of Pharmacy, Mansoura University, Egypt. Male Wistar rats (120–150 g) were obtained from the Animal Facility, College of Pharmacy, Taibah University. Animals were held under standard conditions of 12-hour dark/light cycle, temperature (25±2°C), and humidity (60%–70%). Animals were freely fed standard rat chow and water.

**Experimental protocol**

Rats were randomly divided into four groups of eight animals each and were treated as follows.

1. Control group where rats received oral DMSO once daily for 2 weeks.
2. DOX group: rats injected with DOX (2.5 mg/kg) six times over 2 weeks. This would result in a cumulative dose of 15 mg/kg body weight.\(^{18}\)
3. DOX + Pris groups: in the two remaining groups, rats were treated with Pris at two different doses (0.5, 1 mg/kg, intraperitoneal [IP]) once daily for 1 week before and 2 weeks contaminant with DOX injection. Pris doses were selected based on previous studies that reported its potent anti-inflammatory activity.\(^{15}\)

At the end of the treatment period, rats were weighted and anesthetized using ketamine (75 mg/kg, IP) and then subjected to electrocardiography (ECG) recording. Blood samples were obtained; serum was separated and frozen at −20°C for estimation of biochemical parameters. Rats were humanely killed; their hearts were dissected and washed with PBS (10 mM PO\(_4\)\(^{3−}\), 137 mM NaCl, and 2.7 mM KCl; pH 7.4) and then plotted dry on filter paper and weighted. Relative heart weight was determined as the ratio of the absolute weight of the heart divided by the body weight multiplied by 100.

Small piece of left ventricle was immediately flesh frozen and kept at −80°C for measurements of gene expression. Another piece of the ventricle was homogenized in PBS and then centrifuged at 10,000×g for 20 minutes at 4°C. The supernatant was kept at −20°C till further analysis of oxidative stress parameters, antioxidants, and cytokines. The remaining portion of the left ventricle was fixed in 10% neutral buffered formaldehyde and submitted for histopathological and immunohistochemical (IHC) analysis.

**ECG**

Needle electrodes were inserted under the skin of the animals in lead II position. ECG was recorded and analyzed using an iWorx data recorder and the Labscribe2 software (model 214; iWorx Systems Inc., Dover, NH, USA).

**Serum biochemical analysis**

Serum CK-MB, LDH, cTnI, and cTnT levels were determined based on the instructions of the kits.

**Histopathological and IHC analysis**

Heart specimen was stained with H&E and examined using a light microscope. Heart lesions were determined based on the degree of myofibrillar loss, apoptosis, necrosis, and fibrosis. The severity of myocardial injury was graded according to a semiquantitative score ranking from 0 to 3 depending on the degree and extent of histopathological alteration.\(^{13}\) Another heart specimen was stained with Masson trichrome (MT) stain for assessment of fibrosis development and deposition of ECM in the cardiac tissue. The fibrosis percentage was calculated as the ratio of the interstitial fibrosis area relative to the total specimen area.\(^{18}\)

IHC for NF-κB p65 was automatically performed using Ventana BenchMark XT system (Ventana Medical Systems, Oro Valley, AZ, USA) as described previously.\(^{13}\) In brief, paraffin-embedded heart slides were dewaxed and rehydrated. Heat-induced antigen retrieval was achieved by boiling tissue sections in 10 mM citrate buffer (pH 6.0) for 10 minutes in microwave followed by cooling at room temperature for 20 minutes. Endogenous peroxidase activity was blocked with 3% hydrogen peroxide. Heart sections were then incubated with primary antibodies against rabbit polyclonal anti-NF-κB p65 (1:100) antibody at room temperature for 30 minutes. Biotin-conjugated anti-rabbit secondary antibody (1: 300) was incubated at 37°C for 2 hours. Diaminobenzidine was utilized for visualization of IHC reactions. Hematoxylin was used for counterstaining of sections. Semiquantitative analysis was performed using the open resource digital image analysis software (ImageJ, 1.46a; NIH), implemented with the IHC Profiler plug-in developed by Varghese et al.,\(^{19}\) which creates a pixel-by-pixel analysis profile of a digital IHC image, and further assigns an OD score in a four tier system: 1 – negative, 2 – low positive, 3 – positive, and 4 – high positive. All images were captured at the same magnification and with the same levels of contrast and brightness.

**Estimation of hydroxyproline content**

The hydroxyproline content of the cardiac tissue reflects the total amount of collagen and so it was quantified to assess the degree of cardiac fibrosis. Detection was performed according to the kit procedures. Briefly, 100 mg of the right ventricle was homogenized in dH\(_2\)O. Tissue homogenates were mixed with concentrated NaOH (10 N) in a tightened screw-capped polypropylene vial and then heated at 120°C.
for 1 hour. The alkaline lysate was cooled on ice, neutralized, and centrifuged to obtain supernatants. The hydrolysates were dried, oxidized, and finally reacted with 4-(dimethylamino) benzaldehyde. The resultant colored product was detected at 560 nm and found to be proportional to the hydroxyproline present in the sample.

Oxidative stress and antioxidant markers
Levels of 4-HNE, MDA, PC, 8-OHdG, GSH, and SOD were determined in the cardiac homogenates following the protocols of the kits.

Real-time PCR
Total RNA was extracted from heart tissues using RNeasy Mini Kit according to the manufacturer’s protocol (Qiagen NV, Venlo, the Netherlands). Each sample was subjected to purity quantification using OD 260/280 nm absorbance ratio. Pure samples with a 260/280 ratio between 1.8 and 2.1 were selected for subsequent experiments. cDNA was obtained from RNA (1 µg) using a cDNA reverse transcription kit as described in the manufacturer’s protocol (Qiagen NV). Real-time quantitative PCR (SYBR Green PCR Master Mix Kit) was used to determine the mRNA expression of targeted genes as described previously. Sequences of the primers are listed in Table 1. Specificity of PCR products was verified by melting curve analysis.

Western blot analysis
This was conducted as previously described. In brief, total tissue proteins were extracted according to the instructions of the total protein extraction kit and the protein concentration was measured using the BCA Protein Assay Kit. Equal amounts of protein (20 µg) were loaded into the wells of 8% SDS-PAGE gel, transferred to a cellulose acetate membrane, and then blocked with 5% nonfat milk. The membranes were incubated (4°C) overnight with the primary antibodies including TGF-β1, MMP-2, MMP-9, TIMP-1, Nrf2, GCLc, NQO1, HO-1, NF-κB p65, p-NF-κB p65, IKKα, p-IKKα, p-IκBα, p38 MAPK, p-p38 MAPK, JNK, p-JNK (diluted to 1:1,000), ERK (diluted to 1:2,000), and p-ERK (diluted to 1:250). The membranes were incubated for 1 hour with the secondary antibodies (1:200) at room temperature before the detection of immunoreactive proteins. β-Actin (1:1,000) was used as an internal control.

ELISA assay
NF-κB p65 activation
Nuclear extract of the cardiac tissue was obtained using the extraction kit. NF-κB p65 activation was measured based on the instruction manual of the kit. Results were represented as OD 450 nm.

Cytokines
Levels of NOx, TNF-α and IL-6 were quantified in the cardiac homogenates based on the protocols provided by the manufacturer of the kits.

Statistical analysis
One-way ANOVA followed by Tukey–Kramer multiple comparison test was used for statistical comparisons between different groups. Values are presented as means ± standard error for eight rats in each group. P-value <0.05 was considered significant.

Table 1 Primers used for real time-PCR measurements

<table>
<thead>
<tr>
<th>Gene</th>
<th>Forward primer</th>
<th>Reverse primer</th>
<th>Product size</th>
<th>NCBI reference sequence</th>
</tr>
</thead>
<tbody>
<tr>
<td>TGF-β1</td>
<td>AGGGCTACCTAGCCTCAACCTTC</td>
<td>CCACGTAGTAGACGATGCGG</td>
<td>168</td>
<td>NM_021578</td>
</tr>
<tr>
<td>MMP-2</td>
<td>AGAAGGCTGTGTTCTGCGA</td>
<td>AAAGGAGGAGTCTACTTCTG</td>
<td>128</td>
<td>NM_031054</td>
</tr>
<tr>
<td>MMP-9</td>
<td>GATCCCTCCAGAGGCTACCTCG</td>
<td>GGTGGAGAACACACAGC</td>
<td>132</td>
<td>NM_031055</td>
</tr>
<tr>
<td>TIMP-1</td>
<td>TGCTCAAGGTCAGATGACCT</td>
<td>CCAGGAGGTTGCTAGTGTC</td>
<td>200</td>
<td>NM_053819</td>
</tr>
<tr>
<td>Fibronectin</td>
<td>GATGCCCTCCTCCAGAGAAGT</td>
<td>GGGTGGAGGAGCACTTC</td>
<td>188</td>
<td>NM_019143</td>
</tr>
<tr>
<td>Coll1</td>
<td>GAGGCTCAGAGGTGCTCCTCC</td>
<td>CGGCCACCATCTTGGAGCTT</td>
<td>185</td>
<td>NM_053304</td>
</tr>
<tr>
<td>Nrf2</td>
<td>TTTGATAGATGCATGATGATCCG</td>
<td>TGTCCTGCTATGCTGCT</td>
<td>142</td>
<td>NM_031789</td>
</tr>
<tr>
<td>GCLc</td>
<td>CCACTGTCAGGTTGAGGC</td>
<td>CTTGCTACACCACTACACCA</td>
<td>126</td>
<td>NM_012815</td>
</tr>
<tr>
<td>GCLm</td>
<td>TCAAGCTCACAATCAGGGG</td>
<td>CAAACCACACACATACCAGG</td>
<td>134</td>
<td>NM_017305</td>
</tr>
<tr>
<td>NQO1</td>
<td>GCCCATCATGTTGGCGAGTCTC</td>
<td>TCATTGGAGAAGCGA</td>
<td>145</td>
<td>NM_017000</td>
</tr>
<tr>
<td>HO-1</td>
<td>TCTGCGAGGGGAGAATGCTG</td>
<td>TTGGATGAGGAAGTGC</td>
<td>135</td>
<td>NM_012580</td>
</tr>
</tbody>
</table>

Abbreviations: Coll1, collagen type I alpha 1; GCLc, glutamate-cysteine ligase catalytic subunit; GCLm, glutamate-cysteine ligase modifier; HO-1, heme-oxygenase-1; MMP-2, metalloproteinase-2; MMP-9, metalloproteinase-9; Nrf2, nuclear erythroid 2-related factor 2; TGF-β1, transforming growth factor-beta1; TIMP-1, tissue inhibitor of metalloproteinase-1.
Results

**Pristiminer protects against DOX-induced cardiotoxicity and fibrosis**

Repeated DOX injections resulted in marked cardiac dysfunction in rats. ECG analysis revealed that DOX induced ECG abnormalities as there was significant increase in heart rate, QTc interval, ST segment elevation, and R wave amplitude compared to control animals (Figure 1C). Moreover, DOX administration caused a significant decrease in the relative heart weight (P<0.001) compared to control animals (Figure 1B). Serum biochemical analysis showed that there was significant elevation in the indices of cardiac damage (CK-MB, LDH, cTnI, cTnT) in the DOX group compared to control group (Figure 1D).

Histopathological examination of the heart confirmed the ECG and biochemical results. Control animals exhibited normal architecture of the heart while the DOX group showed extensive deteriorated cardiac damage. Lesions occurred in the form of degeneration of cardiac muscles, coagulative necrosis, and fibrotic areas. Replacement of the cardiac tissue with fibrous tissue was well defined in the heart specimens stained with MT (Figure 2). Hydroxyproline estimation cleared the significant increase of hydroxyproline content of the heart of the DOX-treated animal compared to control indicating the increase in collagen deposition (Figure 2).

Interestingly, Pris treatment significantly reversed all of the DOX-associated cardiotoxic events in a dose-dependent manner. Pris significantly reversed DOX-induced abnormalities in the ECG and increased the relative heart weight compared to the DOX group in a dose-dependent manner. It attenuated and even normalized the serum parameters of cardiotoxicity compared to the DOX group. Pathological lesions were greatly improved and fibrosis was remarkably repressed in Pris-treated animals. Furthermore, Pris attenuated DOX-induced elevation in the hydroxyproline content of the heart tissue compared to the DOX group.

**Pristiminer reduces the expression of fibrogenic genes**

Repeated DOX administration caused elevation of the mRNA expression of the fibrogenic mediators TGF-β1, MMP-2, MMP-9, fibronectin, and coll1-α1 as compared to the control group (Figure 3). Conversely, the mRNA of TIMP-1 was significantly decreased in the DOX group. Results of Western blot were in the same direction as DOX increased the protein expression of TGF-β, MMP-2, MMP-9 contaminant with decrease in TIMP-1. Pris treatment significantly ameliorated DOX-induced alterations of the expression of these fibrogenic genes which may be correlated to the reduction of the collagen deposition in Pris-treated animals.

**Pristiminer inhibits DOX-induced oxidative stress**

As shown in Figure 4, repeated DOX injection induced elevation of lipid peroxidation markers, 4-HNE, MDA, PC, and 8-OHdG, compared to the control group. In addition, DOX caused a significant reduction in GSH content and SOD activity compared to the control group. Pris treatment significantly inhibited the elevation of these lipid per-oxidative parameters simultaneously with the remarkable elevation of GSH and SOD compared to the DOX group.

**Pristiminer activated Nrf2 signaling pathway**

Nrf2 is an important upstream regulator for antioxidant response element-driven expression of antioxidant enzymes, such as GCL, NQO1, and HO-1. As presented in Figure 5, DOX chronic administration resulted in downregulation of mRNA expression of Nrf2 as well as Nrf2 target gene (GClc, glutamate-cysteine ligase modifier, NQO1, and HO-1) compared to control animals. Consistent with mRNA results, Western blot analysis revealed significant reduction of the protein expression of Nrf2, GCL, NQO1, and HO-1 in the DOX group. Additionally, the level of HO-1 was significantly decreased in the hearts of DOX-treated animals. On the contrary, Pris treatment significantly upregulated the mRNA and protein expression of Nrf2 and its target gene. Pris caused significant elevation of HO-1 level compared to the DOX group.

**Pristiminer inhibited activation of MAPK/NF-κB signaling pathways**

To assess the mechanisms of the anti-inflammatory activity of Pris, we examined MAPK and NF-κB signaling pathways. Repeated DOX administration resulted in the activation of MAPK as noticed in the Western blot analysis. The phosphorylation levels of the MAPKs isofoms (p38, JNK, ERK) were elevated in the DOX-treated group although there was no change in the total protein of these isofoms. Meanwhile, Pris treatment dramatically decreased the phosphorylation of MAPKs compared to the DOX group (Figure 6A–C).

Concerning NF-κB signaling pathway, it was greatly activated in the case of repeated DOX administration. IHC and ELISA detection of p65 (the activated subunit of NF-κB) revealed the marked elevation of NF-κB p65 compared to control animals (Figure 6D and E). Western blot analysis showed the marked elevation of the levels of phosphorylated subunits of NF-κB (p-p65, p-1IKκα, and p-1xBκα) compared
to the control group indicating the activation of NF-κB. Conversely, Pris treatment effectively counteracted the activation of NF-κB and suppressed the phosphorylation of p65, IKKα, and IκBα (Figure 6F).

Activation of NF-κB is closely associated with the enhancement of inflammatory cytokines release. This was clearly evident through the elevation of NOx, TNF-α, and IL-6 levels. However, NF-κB signaling upswing was effectively depressed simultaneously with reduction of the levels of NOx, TNF-α, and IL-6 in the case of Pris treatment (Figure 6G).

**Discussion**

DOX-induced cardiotoxicity remains a major clinical challenge to its ubiquitous therapeutic use in oncology. Hence, new safe chemo-protective agents that could prevent the detrimental effects of DOX need to be found. Results of the present study provide new evidences of the cardioprotective activity of Pris against DOX-induced cardiomyopathy and fibrosis which may be mediated via Nrf2 activation and MAPK/NF-κB inhibition.

DOX-induced cardiotoxicity may occur in the form of acute or chronic cardiomyopathy. Acute cardiotoxic effects of DOX occur after administration of a high dose of DOX while chronic cardiotoxicity results from repeated DOX administration and it is dose dependent. Cardiac fibrosis is noted in chronic DOX administration and it contributes to cardiac stiffness and dysfunction. Fibroblasts overproduce collagen which replaces the necrotic or apoptotic myocytes.
The present study revealed that repeated DOX administration produced profound cardiac damage and dysfunction. DOX-associated depression of the cardiac function was evident through ECG abnormalities. Similar results were reported in previous studies. In addition, there was an increase in the serum biomarkers of myocardial toxicity, namely CK-MB, LDH, cTnI, and cTnT, indicating loss of cell membrane integrity of myocytes. These ECG and biochemical data were supported by the results of the histopathological examination of the heart specimen which revealed severe cardiac lesions and remarkable fibrosis, that was clearly noticed in the heart specimen stained by MT stain. Collagen deposition and development of cardiac fibrosis were further confirmed by the significant increase in the hydroxyproline content which is the main constituent of collagen. These findings agree with the outcomes of previous studies. Interestingly, Pris ameliorated all DOX-induced toxic events as presented by the amelioration of ECG abnormalities, lowering of cardiotoxic biomarkers, and improvement of the heart lesions and fibrosis. These data indicate that Pris enabled marked preservation of the myocardium integrity and restoration of the cardiac function.

MMPs are a family of zinc-dependent proteases that play a significant role in the degradation of fibrillar ECM proteins and maintain normal tissue growth and remodeling. There is a balance between MMPs and their TIMPs which is altered during myocardial injury and inflammation resulting in the deregulation of ECM turnover and myocardial fibrosis. Previous studies have linked the rise of MMPs expression with the progression of DOX-induced cardiac dysfunctions and fibrosis. In line with the previous context, our results showed the elevation of the expression of MMP-2 and MMP-9 in the heart of DOX-treated animals simultaneously with depressed expression of TIMP-1. Conversely, Pris markedly downregulated the expression of MMP-2 and MMP-9 contaminant with upregulation of TIMP-1. These results suggest that Pris could effectively modulate the principal regulators of ECM homeostasis which may explain in part Pris-induced amelioration of cardiac fibrosis in DOX-treated animals.
The pathogenesis of fibrosis includes several mechanistic pathways. The inflammatory and growth factor signaling pathways mediated by TGF-β are the basic pathways for fibrosis. TGF-β1 is secreted by cardiac myofibroblasts and controls the progression of cardiac fibrosis. DOX-induced cardiac fibrosis is associated with the overexpression of TGF-β1.18,26 Our results confirmed these previous findings as DOX administration caused elevation of the pro-fibrogenic markers TGF-β, fibronectin, and Col1α1. On the other hand, we observed that Pris-treated groups exhibited attenuation of these pro-fibrogenic genes leading to depression of fibrosis development. Taken together, these data provide evidence that Pris could effectively attenuate fibrosis development in DOX-treated animals via modulation of fibrogenic genes.

ROS produce direct tissue injury and induce inflammatory responses via the activation of different pro-inflammatory cytokines.29 Increased oxidative stress and the resultant lipid peroxidation cause tissue necrosis and inflammation.
and hence promote the progression of tissue fibrogenesis.\textsuperscript{30} As the myocardium has a restricted regenerative capacity, and also is characterized by its relative lack of endogenous antioxidants, this makes the cardiac tissue more vulnerable to oxidative damage.\textsuperscript{2} Overproduction of ROS extensively harms myocardial tissue and destructs cell membrane integrity causing leakage of cardiac diagnostic markers.\textsuperscript{31} Previous investigations have shown the basic role of ROS and oxidative stress in the mediation of DOX-induced cardiotoxicity.\textsuperscript{4,32–34} DOX is known to generate extensively ROS such as hydroxyl radicals and hydrogen peroxide that induce lipid peroxidation and accumulation of the highly reactive electrophile, 4-HNE, which is a good biomarker of DOX-induced oxidative stress.\textsuperscript{35} Furthermore, DOX administration has been shown to depress the level of GSH in the heart.\textsuperscript{36} Antioxidant treatment has shown its ability to attenuate the progression of DOX cardiotoxicity and to improve cardiac function.\textsuperscript{37} In the present study, DOX-induced deleterious oxidative stress was evident through the elevation of PC (marker of the protein oxidative injury), 4-HNE (marker of lipid peroxidation), and 8-OHdG (marker of oxidative DNA injury). However, these indices were significantly depressed in the Pris-treated groups. Additionally, DOX attenuated SOD activity and caused a depletion of myocardial GSH which were reversed by Pris treatment. Therefore, these data indicate the potent antioxidant activity of Pris which impeded DOX-induced oxidative injury. This hypothesis is reinforced by the data of an earlier report of Hui et al,\textsuperscript{38} which demonstrated the free radical scavenging capacity of Pris in LPS-induced tissue injury.

Nrf2 is a redox-sensitive transcription factor that controls detoxification gene activity and cellular responses toward oxidative and electrophilic stress. Normally, Nrf2 is repressed in the cytosol by Kelch-like ECH-associated protein 1 (Keap1). In response to oxidative or electrophilic stress, Nrf2 is released from Keap1 and translocated into the nucleus where it induces the expression of several antioxidant genes such as GCL, NQO1, and HO-1.\textsuperscript{1} Previously, it was reported that DOX administration reduces active nuclear Nrf2 in the heart. The inhibition of Nrf2 pathway has been

Figure 4 Impact of Pris on DOX-induced oxidative stress and depression of antioxidant parameters in cardiac tissue.

Notes: (A) 4-HNE; (B) MDA; (C) PC; (D) 8-OHdG; (E) reduced GSH; (F) SOD. Rats were injected with DOX (2.5 mg/kg) six times over 2 weeks. Pris was administered (0.5, 1 mg/kg, intraperitoneal) once daily for 1 week before and 2 weeks contaminant with DOX injection. Data are means ± standard error (n=8). *P<0.05, **P<0.01, ***P<0.001 vs the control; #P<0.05, ##P<0.01, ###P<0.001 vs the DOX group (one-way ANOVA).

Abbreviations: 4-HNE, 4-hydroxynonenal; 8-OHdG, 8-hydroxy-2-deoxyguanosine; DOX, doxorubicin; GSH, glutathione; MDA, malondialdehyde; PC, protein carbonyl; Pris, pristimerin; SOD, superoxide dismutase.
Figure 5 Effect of Pris on DOX-induced alteration of Nrf2 signaling pathway.

Notes: (A) (i–v) mRNA expression of Nrf2, GCLc, GCLm, NQO1, and HO-1. (vi) Nrf2 binding capacity. (vii) HO-1 level in the heart. (B) Western blot for Nrf2, GCLc, NQO1, and HO-1, and their relative protein quantification. β-Actin served as an internal control. Rats were injected with DOX (2.5 mg/kg) six times over 2 weeks. Pris was administered (0.5, 1 mg/kg, intraperitoneal) once daily for 1 week before and 2 weeks contaminant with DOX injection. Data are the mean ± standard error (n=8). *P<0.05, **P<0.01, ***P<0.001 vs the control; #P<0.05, ##P<0.01, ###P<0.001 vs the DOX group (one-way ANOVA).

Abbreviations: DOX, doxorubicin; GCLc, glutamate-cysteine ligase catalytic subunit; GCLm, glutamate-cysteine ligase modifier; HO-1, heme-oxygenase-1; NQO1, NAD(P)H dehydrogenase quinone; Nrf2, nuclear erythroid 2-related factor 2; Pris, pristimerin.
Figure 6 Effect of Pris on DOX-induced alteration of MAPK/NF-κB signaling pathways.

Notes: Western blot showing the protein expression of (A) p-p38 and p38, (B) p-JNK and JNK, (C) p-ERK and ERK, and quantification of the protein intensities. β-Actin was used as an internal control. (D) Specimen of the cardiac tissue in different groups showing different protein expressions of NF-κB p65 via IHC staining and semiquantitative analysis of IHC expressed as OD across 10 different fields for each section; ×400, scale bar 25 µm. (E) NF-κB p65 activation in the cardiac tissue. (F) Western blot for NF-κB p65, p-NF-κB p65, IKKα, phospho-IKKα, p-IκBα, and quantification of the protein intensities. β-Actin was used as an internal control. (G) Cytokine levels in the cardiac tissue. Rats were injected with DOX (2.5 mg/kg) six times over 2 weeks. Pris was administered (0.5, 1 mg/kg, intraperitoneal) once daily for 1 week before and 2 weeks contaminant with DOX injection. Data are the mean ± standard error (n=8). *P<0.05, **P<0.01 vs the control; #P<0.05, ##P<0.01, ###P<0.001 vs the DOX group (one-way ANOVA).

Abbreviations: DOX, doxorubicin; ERK, extracellular signal-regulated kinase; IHC, immunohistochemical; IKKα, inhibitor of nuclear factor kappa-B kinase alpha; JNK, c-JUN N-terminal kinase; MAPK, mitogen activated protein kinase; NF-κB, nuclear factor kappa-B; p-ERK, phospho-ERK; p-IκBα, phospho-inhibitor of NF-κB; p-JNK, phospho-JNK; Pris, pristimerin.
focused on the fundamental role of NF-κB activation with depression of its target genes. The depression of Nrf2-targeted genes could be responsible in part in the depletion of GSH.\(^7\) In contrast, co-administration of Pris enhanced the expression and activity of Nrf2 and its target genes. These data lead us to the presumption that the cardioprotective activity of Pris may be partially mediated via activation of Nrf2 pathway which enhances the antioxidant capacity of the heart.

MAPKs families are serine/threonine protein kinases that control signal transduction involved in cell proliferation, differentiation, survival, and death in addition to their fundamental role in the regulation of inflammatory mediators. MAPK signaling pathways involve three kinase isoforms: p38, ERK, and JNK. Several lines of evidence indicate that ROS activate both JNK and p38 MAPK leading to the enhancement of apoptosis.\(^39\) Previous investigations have demonstrated the participation of MAPKs in the pathophysiology of DOX-induced cardiac injury.\(^40-42\) Our results were similar and confirmed the role of MAPK in the mediation of DOX-induced cardiotoxic effects. On the contrary, Pris supplementation suppressed DOX-induced MAPKs activation and so our results indicated that the cardioprotective effect of Pris might involve the modulation of MAPK signaling pathway. These findings are in agreement with those of Jin et al.\(^43\) who reported the inhibitory effect of Pris on MAPK signaling pathway in case of ovalbumin-induced allergic airway inflammation.

NF-κB signaling is another major signal transduction pathway that is activated in response to oxidative stress and has crucial roles in cell survival, inflammation, and immune responses. This ubiquitous transcription factor controls the induction of various inflammatory cytokines. Normally, NF-κB p65 subunit binds to its inhibitory counterpart IκBα and other IκB proteins to form an inactive complex that is present in the cytoplasm. Upon activation, IκBα is phosphorylated leading to activation and translocation of NF-κB p65 into the nucleus where it mediates the expression of inflammatory genes. Several studies have focused on the fundamental role of NF-κB activation with its subsequent inflammatory cascade in the mediation of DOX cardiotoxic effects.\(^10-12,25\) Correspondingly, elevations of TNF-α and IL-6 levels are considered as indicators of the inflammatory response in case of DOX intoxication. High expression of these inflammatory cytokines has been reported during DOX administration.\(^44\) Additionally, NF-κB activation has been shown to enhance the generation of NOx through the induction of expression of iNOS. NOx can generate peroxynitrite in cardiomyocyte which causes cell death and exaggerated cardiac damage. Peroxynitrite is also involved with other ROS in the inhibition of multiple mitochondrial enzymes leading to the enhancement of oxidative/nitrosative stress, persistent even in the absence of DOX in the circulation/myocardium.\(^45-47\) The strong correlation between the generated ROS and NF-κB activation results in the perseverance and exaggeration of their injurious impacts.\(^48\) Results of the present study confirmed the phosphorylation and activation of NF-κB signaling during DOX administration. Furthermore, the levels of the cytokines, NOx, TNF-α, and IL-6, were significantly increased. Notably, Pris remarkably inhibited the phosphorylation of NF-κB p65, IKKα, and IκBα in the cardiac tissue indicating that NF-κB activation was greatly depressed in Pris-treated animals. Hence, levels of the related downstream inflammatory cytokines, NOx, TNF-α, and IL-6 were dramatically decreased. These data are consistent with previous studies showing the inhibitory activity of Pris on NF-κB activation and subsequent release of inflammatory cytokines.\(^17,43\) The ability of Pris to inhibit DOX-induced NF-κB activation partially explains the underlying cardioprotective mechanism of Pris in DOX-induced cardiotoxicity. Collectively, it is acceptable to presume that Pris could repress DOX-induced cardiotoxicity via suppression of MAPK/NF-κB phosphorylation.

**Conclusion**

The present findings indicate the apparent protective effects of Pris against DOX-induced cardiotoxic damage. Pris showed potent antioxidant and anti-inflammatory activities which effectively suppressed DOX-induced oxidative damage and inflammatory responses, most likely due to activation of Nrf2 and inhibition of the transcriptional activity of MAPK/NF-κB signaling (Figure 7). Thus, it may be acceptable to suggest that the combined administration of Pris with DOX may provide a safer therapeutic strategy with a more improved therapeutic ratio of DOX. However, the cardioprotective effects of Pris against the cardiotoxic effect of DOX are worthy of further exploration.
Pristimerin protects against DOX-induced cardiotoxicity and fibrosis

Author contributions

Study design and conception: DS El-Agamy; experimental work: N Ahmed, DS El-Agamy, and AA Shaaban; drafting the manuscript: DS El-Agamy, S Khoshhal, HM Abo-Haded, and MA Elkablawy; genetic assessments: KM El-Harbi and S Khoshhal; histopathology and IHC: MA Elkablawy. The authors equally contributed to the acquisition of data and manuscript revision. All authors contributed to data analysis, drafting and revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

References


47. Morgan MJ, Liu ZG. Crosstalk of reactive oxygen species and NF-κB pathway.
Supplementary material

Table S1 PCR efficiency and melting temperature (Tm) of PCR primes used in this study

<table>
<thead>
<tr>
<th>Gene</th>
<th>PCR efficiency</th>
<th>Tm (°C)</th>
</tr>
</thead>
<tbody>
<tr>
<td>TGF-β1</td>
<td>104.9</td>
<td>83</td>
</tr>
<tr>
<td>MMP-2</td>
<td>109.9</td>
<td>85.5</td>
</tr>
<tr>
<td>MMP-9</td>
<td>105.2</td>
<td>89.5</td>
</tr>
<tr>
<td>TIMP-1</td>
<td>95.1</td>
<td>87</td>
</tr>
<tr>
<td>Fibronectin</td>
<td>109.2</td>
<td>85.5</td>
</tr>
<tr>
<td>Collα1</td>
<td>107.6</td>
<td>88.5</td>
</tr>
<tr>
<td>Nrf2</td>
<td>107.9</td>
<td>83.5</td>
</tr>
<tr>
<td>GCLc</td>
<td>105.1</td>
<td>80</td>
</tr>
<tr>
<td>GCLm</td>
<td>96.5</td>
<td>82.5</td>
</tr>
<tr>
<td>NQO1</td>
<td>103.5</td>
<td>84.2</td>
</tr>
<tr>
<td>HO-1</td>
<td>98.6</td>
<td>84</td>
</tr>
</tbody>
</table>

Abbreviations: Collα1, collagen type I alpha 1; HO-1, heme-oxygenase-1; MMP-2, metalloproteinase-2; MMP-9, metalloproteinase-9; GCLc, glutamate-cysteine ligase catalytic subunit; GCLm, glutamate-cysteine ligase modifier; NQO1, NAD(P)H dehydrogenase quinone; Nrf2, nuclear erythroid 2-related factor 2; TGF-β1, transforming growth factor-beta 1; TIMP-1, tissue inhibitor of metalloproteinase-1.