Research Article

Total Flavonoids from Clinopodium chinense (Benth.) O. Ktze Protect against Doxorubicin-Induced Cardiotoxicity In Vitro and In Vivo

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1. Introduction

Doxorubicin (DOX) is an anthracycline derivative widely used to treat various cancers, including acute leukemias, malignant lymphomas, and a variety of solid tumors [1, 2]. However, the prolonged use of DOX may cause cardiotoxicity and lead to cardiomyopathy and heart failure, which severely limit its clinical applications [3]. Numerous mechanisms have been proposed and oxidative stress is believed to play a key role in the intracellular signaling pathways and apoptosis in this pathophysiology [4–6]. Antioxidants reportedly exert protective effects on DOX-treated cardiomyocytes [7]. However, no single active ingredient has been clinically proved effective and safe for preventing this severe side effect. Therefore, the discovery of plant extracts with cardioprotective effects is important to continue the clinical use of DOX.

Intrinsic pathway and extrinsic pathway are the two canonical apoptosis signaling pathways. The activation of p53-initiated intrinsic pathway has been implicated in DOX-induced cardiotoxicity and subsequent chronic cardiomyopathy [8]. It is also known that oxidative stress induced activation of mitogen-activated protein kinase (MAPK) signaling pathways which are the primary intermediates for the induction of apoptosis [9]. MAPK includes three major

1. Doxorubicin (DOX) is an anthracycline derivative widely used to treat various cancers, including acute leukemias, malignant lymphomas, and a variety of solid tumors [1, 2]. However, the prolonged use of DOX may cause cardiotoxicity and lead to cardiomyopathy and heart failure, which severely limit its clinical applications [3]. Numerous mechanisms have been proposed and oxidative stress is believed to play a key role in the intracellular signaling pathways and apoptosis in this pathophysiology [4–6]. Antioxidants reportedly exert protective effects on DOX-treated cardiomyocytes [7]. However, no single active ingredient has been clinically proved effective and safe for preventing this severe side effect. Therefore, the discovery of plant extracts with cardioprotective effects is important to continue the clinical use of DOX.

Intrinsic pathway and extrinsic pathway are the two canonical apoptosis signaling pathways. The activation of p53-initiated intrinsic pathway has been implicated in DOX-induced cardiotoxicity and subsequent chronic cardiomyopathy [8]. It is also known that oxidative stress induced activation of mitogen-activated protein kinase (MAPK) signaling pathways which are the primary intermediates for the induction of apoptosis [9]. MAPK includes three major
signaling cascades: the extracellular signal-related kinases (ERK1/2), the c-Jun N-terminal kinases (JNK), and the p38 kinase (p38). The literature suggests that DOX induces the activation of MAPKs in cardiac pathophysiology [10]. In the present study, we investigated the function of MAPKs during DOX-induced myocardium apoptosis. The phosphoinositide 3-kinase (PI3K)/AKT signaling pathway is an important signaling transduction pathway that plays an important role in the development heart disease and protects cells undergoing apoptosis [11, 12]. We also studied the effect of DOX on this signaling pathway in the present study.

Clinopodium chinense (Benth.) O. Ktze is a traditional Chinese herbal medicine and is originally documented in the “Jiuhuang Herbal,” which belongs to the family Labiatae. It was reported that the stem and leave of Clinopodium chinense (Benth.) O. Ktze possess various pharmacological effects, such as anti-inflammatory and immunity [13, 14], lowering blood glucose [15], and antitumor and antiradiation [16, 17]. The major components have been identified in the C. chinense (Benth.) O. Ktze, including flavonoids, triterpenoid saponins, and volatile oil [16, 18]. Total flavonoids from C. chinense (Benth.) O. Ktze (TFCC) play an important role for the treatment of cardiovascular disease [16]. The instant hypotensive effect of TFCC has been found in normal rabbit. TFCC perfusion reduced contractility and slow rhythm of the heart isolated from toad. Besides, TFCC could also suppress elevation of ECG ST-T induced by pituitrin and improve the ability of hypoxia tolerance in the heart of rabbit injected with epinephrine and norepinephrine [19]. The alcohol extract from C. chinense (Benth.) O. Ktze can effectively reduce oxidative stress injury in HUVECs induced by H₂O₂ [20]. However, there has been no study about the protective effect of C. chinense (Benth.) O. Ktze on DOX-induced cardiotoxicity. The present study aimed to examine the protective effects of TFCC against DOX-induced cardiotoxicity in vitro and in vivo and determine its underlying mechanisms.

2. Materials and Methods

2.1. Ethics Statement. All animal experiments were approved by the Medical Ethics Committee of Peking Union Medical College and were in accordance with the National Institutes of Health Regulations for the Care and Use of Animals. All efforts were made to minimize suffering.

2.2. Plant Materials. C. chinense (Benth.) O. Ktze was purchased from Chinese Medicinal Material Markets (Bozhou, China) in 2012. Effective plant overground part identification was provided by Professor Shichun Yu of the School of Pharmacy, Anhui University of Chinese Medicine. The voucher specimen (20130312) was deposited in the Key Laboratory of Bioactive Substances and Resources Utilization of Chinese Herbal Medicine, Ministry of Education, Beijing, China.

The dried overground part (1 kg) of C. chinense was crushed with a grinder (Tianjin Taisite Instrument Co., Ltd., Tianjin, China) and extracted twice with 8 L of 70% ethanol (v/v) at 70°C for 2 h. The extract was filtered and evaporated to remove ethanol completely. The residue was suspended in water and successively extracted with petroleum ether and ethyl acetate. Approximately 56 g residue of ethyl acetate was loaded on a polyamide column (700 mm × 75 mm) and then eluted with 2 L of 20% ethanol, 8 L of 20% ethanol (v/v), 15 L of 50% ethanol (v/v), and 15 L of 70% ethanol (v/v). The effluents of 50% and 70% ethanol were collected and evaporated at 55°C under vacuum. At last, we obtained 26 g of the 40% and 70% ethanol fractions of the total flavonoids.

2.3. Ultraperformance Liquid Chromatography (UPLC) Analysis. UPLC was performed using an Acquity UPLC system (Waters, USA) with an Agilent phenyl column (100 mm × 2.1 mm i.d., 1.7 μm). The mobile phase was composed of water with 0.2% formic acid (A) and acetonitrile (B) with the following gradient elutions: 0 min to 3 min, 2% to 16% B; 3 min to 6 min, 16% to 20% B; 6 min to 12 min, 20% B; 12 min to 13 min, 20% to 26% B; 13 min to 16 min, 26% to 35% B; 16 min to 20 min, 35% to 50% B; and 20 min to 23 min, 50% to 100% B. The flow rate was set at 0.4 mL/min, the column temperature was 40°C, and the sample injection volume was 10 μL.

2.4. Mass Spectrometry. Mass spectrometry detection was performed on a Synapt G2 MS system (Waters, USA) equipped with an electrospray ionization (ESI) source. Masslynx 4.1 (Waters, USA) was used to control the instrument. Nitrogen gas was used for nebulization. The detection mode of the flight tube was selected to be a “V” pattern. The negative ion spectra of the column elute were recorded within the range of 100 m/z to 1200 m/z. The ESI source conditions were as follows: capillary voltage, 2.5 kV; sampling cone voltage, 4.0 V; extraction cone voltage, 4.0 V; ESI source temperature, 120°C; desolvation temperature, 450°C; cone gas flow, 50 L/h; and desolvation gas flow, 800 L/h. The lock mass compound used was leucine enkephalin (554.2615 m/z), and the interval scan time was 0.02 s.

2.5. Drugs and Reagents. DOX was purchased from Shenzhen Main Luck Pharmaceuticals, Inc. (Shenzhen, China). Cell culture materials were purchased from Gibco (Grand Island, NY). Caspase-3 fluorometric assay kits were acquired from BioVision (CA, USA). The fluorescent dye (JC-1) was purchased from Molecular Probes (Eugene, OR). The kits for determining malondialdehyde (MDA), glutathione peroxidase (GSH-Px), superoxide dismutase (SOD), and catalase (CAT) were purchased from Nanjing Jiancheng Institute of Biological Engineering (Nanjing, China). The kits for measuring lactate dehydrogenase (LDH), creatine kinase (CK), and aspartate aminotransferase (AST) were obtained from Biosino Biotechnology and Science Inc. (Hong Kong, China). All antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA), and other chemicals were purchased from Sigma (St. Louis, MO).

2.6. Animals and Pharmacological Treatment In Vivo. Male Sprague-Dawley (SD) rats weighing 220 g to 250 g were used in the experiments. All SD rats were randomly assigned to the following groups: control (distilled water), TFCC (80 mg/kg), DOX, and TFCC + DOX (20, 40, and 80 mg/kg). These rats were intragastrically treated with distilled water or TFCC for
15 d and then intraperitoneally injected with normal saline or a single dose of DOX (3 mg/kg) every two days for a total of three injections. The TFCC group was only treated with the highest TFCC concentration (80 mg/kg). After 14 d of the first administration of DOX, the rats were euthanized for morphological and cellular studies. The body and heart weights were measured. Serum samples were assayed for the determination of serum cardiac enzymes (e.g., CK, AST, and LDH) by kits using an automatic chemical analyzer (7060, Hitachi, Tokyo, Japan). The left ventricle was excised for histopathological examination, and myocardial homogenates were prepared for MDA, GSH-Px, CAT, and SOD detection by the corresponding kits.

2.7. Heart Histopathological Examination. The heart apex was trimmed and embedded in paraffin blocks after fixation with 4% paraformaldehyde. The heart apex was sectioned and stained with hematoxylin and eosin. The structure was then examined under a light microscope (CKX41, 170 Olympus, Tokyo, Japan) by a pathologist blinded to the groups under study.

2.8. Cell Culture and Treatment. Rat embryonic cardiomyoblast-derived H9c2 cells were cultured in DMEM supplemented with 10% (v/v) fetal bovine serum, 2 mM L-glutamine, 100 U/mL penicillin, and 100 mg/mL streptomycin. The cells were cultured in a humidified incubator with 95% air/5% CO2 at 37°C [21]. For all experiments, the cells were plated at an appropriate density according to the experimental design and grown for 36 h before treatment. After being treated with TFCC (0 μg/mL to 50 μg/mL) with or without 1 μM DOX (4 h after) for 24 h, the cells were harvested at 4°C for further molecular and biochemical analyses.

2.9. Cell Viability Analysis. Cell viability was determined by MTT assay [22]. Briefly, the cells were seeded in 96-well plates at 1 × 10^3 cells/well. After being pretreated with TFCC at different concentrations for 4 h and incubated with 1 μM DOX for 24 h, the cells were incubated with 20 μL 5 mg/mL MTT solution at 37°C for 4 h. The supernatants were discarded, and the metabolized MTT in each well was dissolved in 150 μL of DMSO. Optical density was measured at 570 nm with a microplate reader (BioTek, Vermont).

2.10. Measurement of LDH and MDA Levels and SOD, CAT, and GSH-Px Activities. H9c2 cells were cultured in six-well plates at 3 × 10^5 cells/well. After the experimental procedures, the supernatant was used to measure the level of LDH release using an LDH assay kit. The cells were collected, ultrasonicated, and centrifuged at 1000 rpm for 5 min at 4°C. The supernatant was used to assess activities of MDA, SOD, CAT, and GSH-Px according to the corresponding detection kits (Nanjing Jiancheng Bioengineering Institute, Nanjing, China).

2.11. Hoechst 33342 Staining. Hoechst 33342 staining, which distinguishes apoptotic cells from normal cells on the basis of nuclear chromatin condensation and fragmentation, was used to qualitatively analyze the apoptotic cells. H9c2 cells were cultured on cover slips in 24-well plates for 36 h. After treatment, the cells were incubated with 5 mg/mL Hoechst 33342 at 37°C for 15 min, washed twice with phosphate-buffered saline (PBS), and then observed via fluorescence microscopy (Leica, Heidelberg, Germany).

2.12. TUNEL Staining. Cell apoptosis in the heart tissue and H9c2 cells was detected using TUNEL assay according to the manufacturer’s protocol. After dewaxing and rehydration, the heart sections were incubated with proteinase K (20 mg/mL) at room temperature for 15 min. After rinsing with PBS, the slices were incubated with working-strength terminal deoxynucleotidyl transferase enzyme at 37°C for 1 h in a humidified chamber. After rinsing in a stop/wash buffer, the sections were incubated with working-strength antidigoxigenin conjugate at room temperature for 30 min. The slices were stained with 4′6-diamidino-2-phenylindole and then observed under a fluorescence microscope (Leica, Heidelberg, Germany). H9c2 cells were cultured on cover slips for 36 h. After the treatment, the cells were fixed in 1% paraformaldehyde at room temperature for 30 min. Subsequently, the cells were treated with a permeabilizing solution (0.1% 162 Triton X-100) at 4°C for 2 min, incubated in the TUNEL reaction mixture at 37°C for 60 min, and then visualized by fluorescence microscopy (Leica, Heidelberg, Germany).

2.13. Measurement of Mitochondrial Membrane Potential (ΔΨm). The change of mitochondrial transmembrane potential was determined using JC-1. JC-1 accumulates to form J-aggregates and emits red fluorescence when there are mitochondria with high membrane potential. However, JC-1 emits green fluorescence when this compound dissociates into monomers in mitochondria with no cross-membrane electrochemical gradient. Therefore, the ratio of green to red fluorescence can be a reliable estimate of mitochondrial ΔΨm impairment. H9c2 cells from different treatment groups were washed with PBS and incubated with JC-1 (1 μM) in DMEM at 37°C for 20 min. After washing with PBS, the cells were immediately analyzed under a fluorescence microscope. The ratio of aggregated and monomeric JC-1 was used to quantify the change of mitochondrial membrane potential. Decreased JC-1 ratio represents depolarization of the mitochondria, indicating a decrease in mitochondrial membrane potential.

2.14. Caspase-3 Activity Assessments. Caspase-3 activation was measured using a fluorescein active caspase-3 staining kit according to the manufacturer’s instructions (BioVision, CA, USA). After treatment, approximately 300 μL (1 × 10^6 cells/mL) of the cultures was incubated with 1 μL of the substrate FITC-DEVD-FMK at 37°C for 1 h. These cultures were then centrifuged at 3000 rpm for 5 min, and the supernatant was removed. After washing twice with cold PBS, the cells were resuspended in 100 μL of wash buffer. The cell suspension was transferred to each well in a black microtiter plate. Fluorescence intensity was detected using a microplate reader at excitation and emission wavelengths of 485 and 535 nm, respectively.
2.15. Detection of Intracellular ROS Production. The total ROS detection kit was used to monitor the level of intracellular ROS according to the manufacturer's protocol (Enzo Life Sciences, Farmingdale, NY). After treatment, the cells were harvested, placed in 10 mL round bottom polystyrene tubes, and then washed with 1x wash buffer. The cells were centrifuged at 1000 g for 5 min at room temperature, and the supernatant was discarded. The cells were resuspended in 500 mL of ROS detection solution, stained at 37°C for 30 min in the dark, and then analyzed by flow cytometry.

2.16. Western Blot Analysis. Heart tissues or H9c2 cells were lysed on ice with tissue or cell protein extraction reagent that contains a 0.1 mM dithiothreitol and protease inhibitor cocktail. The protein concentration was determined using a BCA kit (Pierce Corporation, Rockford, USA). Equal amounts of protein fractions were separated by 12% SDS-PAGE and then transferred onto nitrocellulose membranes (Millipore Corporation, USA) in Tris-glycine buffer at 100 V for 55 min. The membranes were blocked with 5% (w/v) nonfat milk powder in Tris-buffer that contains a 0.1 mM dithiothreitol and proteinase inhibitor cocktail. The membranes were washed thrice with TBST, incubated with secondary antibodies for 2 h at room temperature, and then washed again thrice with TBST. Protein blots were developed using an enhanced chemiluminescence solution. The protein expression levels were visualized with Image Lab Software (Bio-Rad, USA).

2.17. Statistical Analysis. Results from at least three independent experiments were expressed as mean ± SE. Statistical comparisons between different groups were measured by using Student's t-test or ANOVA with Prism 5.00 software. Statistical significance was considered as P < 0.05.

3. Results

3.1. Characterization of TFCC Using UPLC/Q-TOF-MS/MS. TFCC were characterized using UPLC/Q-TOF-MS/MS, in which 28 peaks were observed (Figure 1). The 28 peaks were assigned by MS fragments and the previous literatures [23, 24] (Table 1).

3.2. TFCC Reduced DOX-Induced Myocardial Injury in Rats. We first measured the general toxicities of DOX and TFCC in vitro. As shown in Figure 2(a), the rats treated with DOX showed significantly less body and heart weights after 14 d. By contrast, TFCC pretreatment (20, 40, and 80 mg/kg i.p.) resulted in a dose-dependent recovery of body and heart weights. However, no significant difference in the relative heart weight index (heart weight-to-body weight ratio) was detected among all six groups after 14 d (Figure 2(a)).

The levels of cardiac enzymes (CK, AST, and LDH) in serum were evaluated to assay the cardioprotective effects of TFCC. TFCC significantly ameliorated the DOX-induced increase in serum cardiac enzyme levels in a dose-dependent manner. TFCC treatment alone did not cause any apparent abnormalities (Figure 2(b)). Based on the histological assessments of the different cardiac segments of the experimental animals, we observed that DOX injection disorganized the normal radiating pattern of cell plates in the heart. TFCC reduced such changes, and the hearts appeared similar to those of the normal group.

3.3. TFCC Protected against DOX-Induced Cytotoxicity in H9c2 Cells. We evaluated the effect of DOX on cell viability using MTT assay. H9c2 cells were treated with different concentrations of DOX (0.25, 0.5, 1, 2, and 3 μM) for 24 h. As shown in Figure 3(a), cell viability significantly decreased at 0.5 μM DOX and decreased by approximately 50% at 1 μM DOX. DOX caused severe cell damage, and cell viability was only 11.1% and 9.1% at 2 and 3 μM, respectively (Figure 3(a)). Therefore, 1 μM DOX for 24 h was used for subsequent experiments. Before testing the protective effects of TFCC against DOX-induced cytotoxicity in H9c2 cells, we analyzed the direct effect of TFCC on H9c2 cells. Cells were treated with different concentrations of TFCC (6.25, 12.5, 25, and 50 μg/mL) for 4 h. As shown in Figure 3(b), no significant difference in cell viability was observed between the TFCC-treated and control groups. Therefore, the tested TFCC concentrations could not induce cell injury in H9c2 cells and will be used in the next experiments. As shown in Figures 3(c) and 3(d), DOX significantly reduced cell viability and increased LDH release, whereas pretreatment with 25 μg/mL TFCC significantly maintained cell viability at approximately 78% and reduced LDH release. A high concentration of TFCC (50 μg/mL) showed no additional benefit. Therefore, a 25 μg/mL dose was selected for subsequent experiments in vitro.

3.4. TFCC Enhanced Antioxidant Capacity in H9c2 Cells and Cardiac Tissues. The effects of DOX and TFCC with respect to oxidative stress damage were further evaluated in vivo and in vitro. DOX induced oxidative stress damage in both rat heart and H9c2 cells, as indicated by decrease in SOD, CAT, and GSH-Px activities and increase in MDA production (Figures 4(a) and 4(b)). These actions were dose-dependently ameliorated by TFCC. These findings proved the efficacy of TFCC in alleviating DOX-induced oxidative stress.

3.5. TFCC Reduced DOX-Induced ROS Generation in H9c2 Cells. Flavonoids can change the cellular ROS level [25, 26].
Table 1: The MS<sup>+</sup> data and proposed compounds of 28 peaks in total flavonoids from <i>Clinopodium chinense</i> (Benth.) O. Ktze (TFCC).

| Number | $t_R$ (min) | Measured mass (Da) | Calculated mass (Da) | Fragments in MS | Proposed compounds |
|--------|-------------|--------------------|---------------------|----------------|-------------------|
| 1      | 5.01        | 593.15             | 593.15              | 447.09, 285.04 | Kaempferol-3-O-rhamnopyranoside-7-O-glucopyranoside (or kaempferol-3-O-rhamnopyranoside-7-O-glucopyranoside isomer) |
| 2      | 5.57        | 593.15             | 593.15              | 447.10, 285.05 | Kaempferol-3-O-rhamnopyranoside-7-O-glucopyranoside (or kaempferol-3-O-rhamnopyranoside-7-O-glucopyranoside isomer) |
| 3      | 6.00        | 595.17             | 595.17              | 433.16, 287.06 | Eriocitrin |
| 4      | 6.45        | 595.17             | 595.17              | 449.11, 287.06 | Neoeriocitrin |
| 5      | 6.64        | 463.09             | 463.09              | 417.21, 371.21, 287.06 | Tetrahydroxyflavanone-7-O-glucuronide |
| 6      | 6.92        | 593.15             | 593.15              | 283.04 | Luteolin-7-O-rutinoside (or luteolin-7-O-neohesperidoside) |
| 7      | 6.97        | 593.15             | 593.15              | 283.04 | Luteolin-7-O-rutinoside (or luteolin-7-O-neohesperidoside) |
| 8      | 7.41        | 579.17             | 579.17              | 271.06 | Narirutin (or naringin) |
| 9      | 8.13        | 579.17             | 579.17              | 271.06 | Narirutin (or naringin) |
| 10     | 8.42        | 609.18             | 609.18              | 579.16, 433.10, 301.07, 271.06 | Hesperidin |
| 11     | 8.52        | 609.18             | 609.18              | 433.11, 301.07, 271.06 | Neohesperidin |
| 12     | 8.63        | 577.16             | 577.16              | 269.05 | Apigenin-7-O-rutinoside (or apigenin-7-O-neohesperidoside) |
| 13     | 8.70        | 577.16             | 577.16              | 269.05 | Pigenin-7-O-rutinoside (or apigenin-7-O-neohesperidoside) |
| 14     | 8.81        | 447.09             | 447.09              | 271.06 | Naringenin-7-O-glucuronide (or its isomer) |
| 15     | 8.93        | 447.09             | 447.09              | 271.06 | Naringenin-7-O-glucuronide (or its isomer) |
| 16     | 9.33        | 577.16             | 577.16              | 269.05 | Apigenin-7-O-rutinoside isomer |
| 17     | 10.01       | 607.17             | 607.17              | 299.06 | Diosmin/chrysoeriol-7-O-rutinoside |
| 18     | 10.24       | 445.08             | 445.08              | 269.04 | Apigenin-7-O-glucuronide |
| 19     | 11.16       | 461.07             | 461.07              | 285.04 | Luteolin-7-O-glucuronide |
| 20     | 13.87       | 593.19             | 593.19              | 285.08 | Didymin (or poncirin) |
| 21     | 14.17       | 593.19             | 593.19              | 285.08 | Didymin (or poncirin) |
| 22     | 14.44       | 591.17             | 591.17              | 283.06 | Linarin (or its isomer) |
| 23     | 14.51       | 591.17             | 591.17              | 283.06 | Linarin (or its isomer) |
| 24     | 15.10       | 447.13             | 447.13              | 285.08 | Isosakuranetin-7-O-glucopyranoside |
| 25     | 15.87       | 491.10             | 491.10              | 327.22 | Isosalvianolic acid C |
| 26     | 16.34       | 271.06             | 271.06              |  | Naringenin |
| 27     | 16.78       | 269.04             | 269.04              |  | Apigenin |
| 28     | 19.55       | 285.08             | 285.08              |  | Isosakuranetin |

Therefore, we determined whether or not TFCC show a potent ROS-scavenging effect. The results of fluorescence assay (Figure 5(a)) and quantitative flow cytometry (Figures 5(b) and 5(c)) showed that H9c2 cells exposed to DOX displayed a significant increase in ROS production (approximately 76% versus control). By contrast, TFCC pretreatment effectively reduced DOX-induced ROS production. However, TFCC alone had no effect on ROS levels compared with the control treatment.

3.6. TFCC Protected against DOX-Induced Apoptosis In Vivo and In Vitro. We examined the nuclear morphology after
Figure 2: Effects of TFCC on DOX-induced myocardial injury \textit{in vivo}. Rats were treated with vehicle or DOX (i.p., 3 mg/kg every other day) with or without TFCC pretreatment (20, 40, and 80 mg/kg i.p.). At day 14, body weight, absolute heart weight, and relative heart weight index (heart weight-to-body weight ratio) were determined (a); effects of TFCC and DOX on AST, LDH, and CK activities were measured (b); effects of TFCC and DOX on histological changes in rat hearts by HE staining (200x) (c). Results are represented as the mean ± SE. *$P < 0.05$ relative to the control group, and $^*$ $P < 0.05$ relative to the DOX group.
Figure 3: Effects of DOX and TFCC on cardiomyocyte viability and LDH release in vitro. H9c2 cells were treated for 24 h with different concentrations of DOX, and cell viability was measured by MTT assay (a). H9c2 cells were treated with different concentrations of TFCC for 4 h, and cell viability was measured by MTT assay (b). Control and TFCC-treated cells were further exposed to 1 𝜇M DOX for 24 h. Cell viability (c) and LDH release (d) were measured. Results are represented as the mean ± SE. *P < 0.05 relative to the control group, and #P < 0.05 relative to the DOX group.

Hoechst 33342 staining to determine the effect of TFCC on DOX-induced apoptosis in H9c2 cells. DOX induced nuclear changes in H9c2 cells with heterogeneous intensity and chromatin condensation. Pretreatment with 25 𝜇g/mL TFCC for 4 h exhibited a strong antiapoptotic effect (Figure 6(a)). Caspase-3 activity was significantly upregulated in the cells treated with DOX and was inhibited by TFCC pretreatment (Figure 6(b)). Afterward, we examined the DNA fragmentation pattern to show the apoptotic changes in H9c2 cells and rat myocardial tissues. DOX caused DNA ladder fragmentation (Figure 6(c)), whereas TFCC treatment effectively reduced DNA laddering in the DOX-treated cardiomyocytes. However, TFCC alone showed no significant effect on apoptotic processes in both in vivo and in vitro experiments.

3.7 TFCC Prevented DOX-Induced Apoptosis through Mitochondrion-Dependent Apoptotic Pathway. The ultrastructural changes in cardiomyocytes were examined to evaluate the myocardial damage induced by chronic DOX treatment. The control and TFCC groups showed no significant morphological abnormalities. However, DOX treatment resulted in myofibril loss, cytoplasm vacuolization, and mitochondrial swelling with membrane disruption and these results were significantly attenuated by TFCC pretreatment (Figure 7(a)). Mitochondrion is the key organelle related to cell energy supply and cellular apoptosis. A decrease in the mitochondrial membrane potential causes membrane depolarization and triggers a cascade of apoptotic signaling [27]. In the present study, JC-1 staining showed that TFCC pretreatment abated DOX-induced decrease in ΔΨ in H9c2 cells (Figure 7(b)). Disruption of mitochondrial membrane permeability results in mitochondrial cytochrome c release to the cytosol, which in turn activates the apoptotic effector caspase-3 to induce cell apoptosis [28]. Immunofluorescence results showed that DOX caused mitochondrial cytochrome c release into
the cytosol and subsequent activation of caspase-9/3 and poly ADP-ribose polymerase (PARP) cleavage. In contrast, TFCC inhibited DOX-induced mitochondrial cytochrome c release and prevented the cleavage of caspase-9/3 and PARP (Figure 7(c)).

3.8. TFCC Preserved DOX-Induced Disturbance in the Expression Patterns of Bcl-2 and Bax In Vivo and In Vitro. TFCC showed protective effect against DOX-induced mitochondrial damage. Therefore, we subsequently evaluated the expression
levels of Bcl-2 proteins, mainly including the proapoptotic Bax and the antiapoptotic Bcl-2, which are upstream regulators of mitochondrial potential [29]. The results of Western blot assay showed that DOX caused elevated Bax protein levels and concomitance with decreased Bcl-2 protein levels in both rat heart tissues and H9c2 cells (Figure 8(a)). However, TFCC mitigated DOX-induced Bax upregulation and Bcl-2 downregulation. Statistical calculation of the Bcl-2 to Bax ratio with respect to protein levels revealed that TFCC alleviated the DOX-induced disturbance in Bcl-2 and Bax levels of Bcl-2 proteins, mainly including the proapoptotic Bax and the antiapoptotic Bcl-2, which are upstream regulators of mitochondrial potential [29]. The results of Western blot assay showed that DOX caused elevated Bax protein levels and concomitance with decreased Bcl-2 protein levels in both rat heart tissues and H9c2 cells (Figure 8(a)). However, TFCC mitigated DOX-induced Bax upregulation and Bcl-2 downregulation. Statistical calculation of the Bcl-2 to Bax ratio with respect to protein levels revealed that TFCC alleviated the DOX-induced disturbance in Bcl-2 and Bax levels of Bcl-2 proteins, mainly including the proapoptotic Bax and the antiapoptotic Bcl-2, which are upstream regulators of mitochondrial potential [29]. The results of Western blot assay showed that DOX caused elevated Bax protein levels and concomitance with decreased Bcl-2 protein levels in both rat heart tissues and H9c2 cells (Figure 8(a)). However, TFCC mitigated DOX-induced Bax upregulation and Bcl-2 downregulation. Statistical calculation of the Bcl-2 to Bax ratio with respect to protein levels revealed that TFCC alleviated the DOX-induced disturbance in Bcl-2 and Bax
Figure 6: Effects of DOX and TFCC on apoptosis in vivo and in vitro. H9c2 cells were treated with vehicle or DOX (1 μM) with or without TFCC (25 μg/mL for 4 h prior to DOX exposure). After 24 h, Hoechst 33342 staining (a) and TUNEL assay (d) were used to measure the apoptosis in H9c2 cells. Caspase-3 activity (units per microgram of protein) was assayed as described in Materials and Methods (b). Rats were treated with vehicle or DOX (i.p., 3 mg/kg every other day) with or without TFCC pretreatment (40 mg/kg i.p. before DOX administration). At day 14, heart tissue section was obtained, and TUNEL assay of heart tissue was measured (c). Cont, vehicle treatment; TFCC, TFCC treatment; DOX, doxorubicin treatment; TFCC + DOX, TFCC and DOX cotreatment. The results are represented as the mean ± SE. *P < 0.05 relative to the control group, and #P < 0.05 relative to the DOX group.
Figure 7: Effects of DOX and TFCC on mitochondrial apoptotic pathway in vivo and in vitro. Rats were treated with vehicle or DOX (i.p., 3 mg/kg every other day) with or without TFCC pretreatment (20, 40, and 80 mg/kg i.p.). At day 14, the ultrastructure of cardiomyocytes was observed using electron microscope (a). H9c2 cells were treated with vehicle or DOX (1 μM) with or without TFCC (25 μg/mL for 4 h prior to DOX exposure). After 24 h, cells were stained with JC-1 dye and were visualized by fluorescence microscopy. Quantitative analysis of JC-1 staining was evaluated (green to red fluorescence ratio) (b). Cytochrome c (cyt c) in mitochondrial and cytosolic protein extracts and caspase-3/9 and PARP levels in rat heart tissues and H9c2 cells were determined using Western blot analysis (c). Cont, vehicle treatment; TFCC, TFCC treatment; DOX, DOX treatment; TFCC + DOX, TFCC and DOX cotreatment. The results are represented as the mean ± SE. *P < 0.05 relative to the control group, and **P < 0.05 relative to the DOX group.
3.9. Effects of TFCC and DOX on Phosphorylation of Different MAPKs In Vitro. MAPK plays a key role in DOX-induced apoptosis [30]. To determine the involvement of the MAPK signaling pathway, the phosphorylation of ERK1/2, p38, and JNK kinases was examined in H9c2 cells of the control, TFCC, DOX, and TFCC + DOX groups. As shown in Figures 9(a) to 9(d), DOX induced a significant increase in the phosphorylation of ERK1/2, JNK, and p38 (after 24 h incubation) and was neutralized by TFCC pretreatment. However, TFCC alone had no influence on MAPK activation (Figures 9(a) to 9(d)).

3.10. Involvement of PI3K/Akt in DOX-Induced Apoptosis. We next investigated whether PI3K/AKT signaling pathway was involved in the protective effect of TFCC on DOX-induced cardiotoxicity. Results showed that DOX significantly decreased AKT phosphorylation and PI3K protein levels (Figures 10(a) and 10(b)), whereas TFCC pretreatment significantly increased the phosphorylation of these proteins. Next, we evaluated the effects of TFCC and DOX on cell viability and caspase-3 activation in cardiomyocytes with or without LY294002 (specific inhibitor of PI3K/AKT). The results showed that LY294002 pretreatment significantly inhibited the protective effect of TFCC against DOX-induced death and apoptosis in H9c2 cells. However, LY294002 alone had no influence on cell viability and caspase-3 activation (Figures 10(c) and 10(d)). These results indicated that the PI3K/AKT activation may participate in the cardioprotective effects of TFCC.

3.11. TFCC Had No Effect on DOX Antitumor Activity. We subsequently evaluated the effect of TFCC on antitumor ability of DOX. As shown in Figure 11, DOX exhibited significant inhibitory effects on the viability of different cancer cell lines, including MCF-7, KB, and HepG2 cells. Cotreatment with TFCC had no effect on DOX-induced growth delay of these cells. TFCC alone also did not affect the growth of these cells. These results suggested that TFCC had no influence on DOX antitumor effect.

4. Discussion
First, our study explored the protective effects of TFCC on DOX-induced myocardial injury. Further study illustrated that TFCC counteract against oxidative stress induced by DOX partially via the mitochondrion-dependent and the p53-mediated apoptotic signaling. Additionally, TFCC also activated the PI3K/AKT signaling pathways which benefit cell survival. These results suggested that TFCC could be a possible antidote to DOX-induced cardiotoxicity in clinical practice.

DOX, used in cancer patients, might induce cardiomyopathy, which can lead to heart failure [2, 31, 32]. The results demonstrated that DOX administration increased the serum levels of LDH, AST, and CK. The increase of these enzymes can cause heart failure. Besides, it could also increase cardiac index and induce histological damage in the heart. However, TFCC pretreatment can reverse this pathological condition. Experiments in vitro also proved that pretreatment with TFCC can effectively increase cell viability and suppress DOX-induced LDH release in H9c2 cells. ROS overproduction and subsequent oxidative stress serve important functions in the initiation and progression of DOX-induced myocardial dysfunction [10]. The toxic effect of DOX on cardiomyocytes is largely attributed to its inherent chemical structure, which induces the generation of free radicals to provoke cell apoptosis [6]. In accordance with
Figure 9: The effects of DOX and TFCC on phosphorylation of different MAPKs in vitro. H9c2 cells were treated with vehicle or DOX (1 μM) with or without TFCC (25 μg/mL for 4 h prior to DOX exposure) for 24 h. The protein levels of phosphorylated and total ERK1/2, p38, and JNK were measured using Western blot analysis (a) and were expressed as the fold changes over the control ((b), (c), and (d)). Cont, vehicle treatment; TFCC, TFCC treatment; DOX, DOX treatment; TFCC + DOX, TFCC and DOX co-treatment. The results are represented as the mean ± SE. *P < 0.05 relative to the control group, and #P < 0.05 relative to the DOX group.

In the previous study, we also proved that treatment with DOX could increase the production of oxidative free radicals and depressed expression of antioxidant enzymes in heart tissue and cardiomyocytes, indicating that DOX induced oxidative stress on myocardial cells through both evoking free radical generation and impeding intrinsic protective antioxidant capacity [33–35]. Administration of the antioxidant agent may prevent heart dysfunction in doxorubicin-treated rats [36]. Numerous studies investigated the cardioprotective properties of plant-derived polyphenolic nonsteroidal compounds, including flavonoids. It has been demonstrated that pretreatment with these agents elicits strong antioxidant activities [31, 37, 38]. In the study, TFCC could modulate the cardiomyocytes redox state by modulating the activity of enzymes involved in ROS metabolism and then effectively reduced DOX-induced alterations.

The mitochondrion is crucial in maintaining normal cell function, which is an important site of intracellular injury in cardiomyocytes after DOX exposure. Mitochondrion-dependent apoptosis in cardiomyocytes is the major factor in DOX-induced cardiotoxicity [27, 39]. We observed that DOX induced apoptosis (confirmed by DNA fragmentation analyses), upregulated proapoptotic (Bax) and downregulated antiapoptotic (Bcl-2) proteins, reduced mitochondrial membrane potential, and increased cytochrome c release into the cytosol and the cleavage of caspase-3 in cardiomyocytes. Additionally, DOX significantly increased intracellular Ca^{2+}. Previous study demonstrated that cytoplasmic Ca^{2+} level is elevated after ROS accumulation, and increased Ca^{2+} influx in the mitochondria leads to mitochondrial transmembrane potential [40]. The damage of the mitochondrial transmembrane potential resulted in the release of cytochrome c from the intermembrane space to the cytoplasm, where cytochrome c induces caspase activation and provokes cell apoptosis [41]. Mitochondrion-mediated apoptosis can be modulated by the proteins of Bcl-2 family [30]. Bax translocates to the outer mitochondrial membrane and influences the permeability of the outer mitochondrial membrane in response to enhanced oxidative stress, which ultimately induces cytochrome c release from the mitochondria into the cytosol. However, Bcl-2 can stabilize the mitochondrial transmembrane potential, preserve mitochondrial integrity,
and inhibit cytochrome c release [42, 43]. Thus, DOX could activate the mitochondrial-mediated apoptotic signal transduction pathways in cardiomyocytes. However, TFCC could antagonize all of these DOX-mediated proapoptotic events via inhibiting the production of ROS. Studies have shown that DOX induced apoptosis of cardiomyocytes via increasing the expression of p53 tumor suppressor protein, which activates the intrinsic apoptosis pathway [8, 44]. In the present study, we also observed that p53 expression increased in both hearts and cardiomyocytes exposed to DOX. TFCC reduced this activation of p53.

MAPK signaling pathways serve an important function in cell proliferation, differentiation, and death, which can be modulated by ROS-dependent redox cycling [45–47]. A recent study has revealed that DOX treatment increased phosphorylation of ERK1/2 persistently during the early stages of apoptosis in cardiomyocytes, whereas it had no effects on p-JNK and p-p38. However, another report has shown that p38 plays a key role in the regulation of cell apoptosis [48]. The difference in MAPK regulation may partially depend on the doses of DOX and the experimental conditions used. In the present study, TFCC pretreatment

Figure 10: Effects of DOX and TFCC on PI3K/AKT signaling pathway activation in vitro. H9c2 cells were treated with vehicle or DOX (1 μM) with or without TFCC (25 μg/mL for 4 h prior to DOX exposure) for 24 h. The PI3K and AKT protein levels were measured using Western blot analysis and were expressed as the fold changes over the control ((a) and (b)). H9c2 cells were pretreated with 10 mM LY294002 for 30 min and then treated with DOX (1 μM) with or without TFCC (25 μg/mL for 4 h prior to DOX exposure) for 24 h. Cell viability and caspase-3 activity were measured ((c) and (d)). Cont, vehicle treatment; TFCC, TFCC treatment; DOX, DOX treatment; TFCC + DOX, TFCC and DOX cotreatment. The results are represented as the mean ± SE. *P < 0.05 relative to the control group, #P < 0.05 relative to the DOX group, and $P < 0.05 relative to the TFCC + DOX group.
neutralized DOX-induced phosphorylation of ERK1/2, p38, and JNK, suggesting that TFCC, at least partially, prevented DOX-induced apoptosis by inhibiting the MAPK (ERK, JNK, and JNK) signaling pathway. It may also increase PI3K/AKT phosphorylation, which promotes cell survival. Therefore, the results of the present study elucidate the fact that TFCC may have implications in the long-term clinical use of DOX.

Conflict of Interests

The authors declare no conflict of interests regarding the publication of this paper.

Authors’ Contribution

Experimental concept and design were performed by Rong Chang Chen, Xu Dong Xu, Gui Bo Sun, and Xiao Bo Sun. Rong Chang Chen, Xu Dong Xu, Yin Di Zhu, Xi Dong, Jian Wang, and Hai Jing Zhang executed experimental performance. Rong Chang Chen, Xu Dong Xu, and Gui Bo Sun analyzed data. Rong Chang Chen and Yin Di Zhu wrote the paper. Rong Chang Chen and Xu Dong Xu equally contributed to this work.

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