

Evaluation of the Liver Cancer Prevention of Anthocyanin Extracts from Mulberry (*Morus alba* L.) Variety PR-01

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Abstract

This study aims to evaluate the preventive effects of anthocyanins extracts (MAEs) from mulberry variety PR-01 against *N*-nitrosodiethylamine (NDEA)-induced hepatocarcinogenesis in rats. It was found that 150 mg·kg⁻¹ MAEs treatment significantly reduced the NDEA-induced hepatic nodules incidence and hepatocellular carcinoma incidence by 58.30% and 41.70% compared to the model group. Meanwhile, MAEs significantly restored the elevated the liver function enzymes, inhibited the tumor necrosis factor alpha and interleukin-6 levels, elevated the serum interleukin-10 and interferon- γ and increased hepatic glutathione-S-transferase and UDP-glucuronosyltransferase 2B1 enzyme activity. Moreover, 150 mg·kg⁻¹ MAEs supplement enhanced glutathione content and the activities of superoxide dismutase, catalase, glutathione peroxidase activities but reduced the malondialdehyde and thiobarbituric acid-reactive substances content by 37.90% and 44.52%. Furthermore, MAEs pretreatment maintained nuclear factor erythroid 2-related factor 2 (Nrf2), Kelch-like ECH-associated protein 1, heme oxygenase-1, and NAD(P)H: quinine oxidoreductase1 stimulation and inhibited the expression of TNF- α , nuclear factor-kappaB (NF- κ B), and cyclooxygenase-2 (COX-2), indicating that MAEs exhibit effectively prevention effects against liver cancer via decreased lipid peroxidation, induced Nrf2-mediated antioxidant enzymes and attenuating the inflammatory mediators COX-2 through NF- κ B pathway. Thus, MAEs of mulberry variety PR-01 may be used as a good functional dietary supplement against liver cancer.

Keywords

Mulberry (*Morus alba* L.) Anthocyanins Extracts, *N*-Nitrosodiethylamine,

1. Introduction

According to the histological classification, primary liver cancer can be divided into hepatocellular carcinoma (HCC), cholangiocarcinoma and mixed liver cancer, among which HCC is the most common [1] [2]. Despite significant improvements in the clinical diagnosis and treatment of early HCC, the advanced HCC is a highly aggressive tumor with poor or no response to conventional treatment [3] [4]. Therefore, it is urgent to find effective ways or measures to prevent liver cancer.

In the past few decades, dietary studies have shown that regular consumption of fruits and vegetables reduces the risk of chronic diseases such as heart disease and cancer [5] [6] [7]. Colored fruits and vegetable foods contain abundant anthocyanins, including cyanidin-3-glucoside (C3G), pelargonidin-3-glucoside (P3G), malvidin-3-glucoside (M3G), peonidin-3-glucoside, petunidin-3-glucoside, and delphinidin-3-galactoside, etc. [8] [9]. Epidemiological and laboratory data have shown that anthocyanins have many biological activities, such as inhibition of the growth of cancerous cells, improving immune responses [7] [10], reducing cardiovascular diseases [11], and promoting anti-obesity effects [12].

Mulberry (*Morus alba* L.) fruit has been traditionally used in Chinese medicines for its pharmacological effects including antioxidant activity, anti-inflammatory, hepatoprotective effect, and anti-diabetic properties, etc. [13]. Mulberry is rich in anthocyanins (The anthocyanins content was 11.20 mg·100g⁻¹ - 193.00 mg·100g⁻¹ in mulberry varieties), among which, the C3G and cyaniding-3-rutinoside (C3R) have been demonstrated to dose-dependently inhibit the migration and invasion of A549 human lung carcinoma cells [14]. Yan *et al.* [15] showed that mulberry anthocyanin extract ameliorates oxidative damage in HepG2 Cells and prolongs the lifespan of *Caenorhabditis elegans* through mitogen-activated protein kinase and nuclear factor erythroid 2-related factor 2 (Nrf2) pathways. Li, *et al.* [10] reported that mulberry anthocyanin has a strong protective effect on the carbon tetrachloride-induced liver fibrosis. Furthermore, it was recently demonstrated that the mulberry extract can increase the protein expression of liver antioxidant, accelerate the metabolism and excretion of nitrophenol, and detoxify the toxicity of nonyl phenol-induced rats [6]. Results of the above studies suggested that mulberry anthocyanins may have potential effects in reducing the risk of cancers by anti-inflammatory, detoxication and chemoprotective properties. However, the anti-cancer activity and the potential anti-cancer mechanisms of mulberry anthocyanins *in vivo* have not been well elucidated.

In our preliminary work, we have found that black mulberry fruit is abundant in anthocyanins, especially the new cultivars, mulberry variety PR-01, which the

anthocyanins content is 193.00 mg·100g⁻¹, 10.16-fold that of the common variety (19.00 mg·100g⁻¹) [16]. Therefore, the present study is to determine the mulberry anthocyanin extracts (MAEs) composition of mulberry variety PR-01 fruits and investigate its prevention effects against the *N*-nitrosodiethylamine (NDEA)-induced hepatocarcinogenesis in rats and its possible mechanisms.

2. Materials and Methods

2.1. Materials and Chemicals

Fresh mature mulberry variety PR-01 fruits was obtained in May 2016 from the Institute of Agricultural Product Quality, Fujian Agriculture and Forestry University (Fujian, China) and stored at -80°C on the same day.

All chemicals were ordered from Sigma-Aldrich (Shanghai, China) or Beijing Dingguo Changsheng Biotech Co., Ltd. (Beijing, China). TRIzol reagent, Hi-Script II Q RT SuperMix for qPCR (+cDNA wiper) and AceQTM qPCR SYBR Green Master Mix were purchased from Takara Co., Ltd. (Dalian, China). Rabbit polyclonal antibodies nuclear factor-kappaB (NF- κ B), cyclooxygenase-2 (COX-2), tumor necrosis factor alpha (TNF- α), Kelch-like ECH-associated protein 1 (Keap1), NAD(P)H:quinine oxidoreductase1 (NQO1), heme oxygenase-1 (HO-1) and Nrf2 were purchased from Santa Cruz Biotechnology or Cell Signaling (Danvers, MA, US).

2.2. Preparation of Anthocyanin Rich-Extract of Mulberry

The PR-01 mulberry anthocyanin rich-extract was extracted as previously described [17]. Freeze mulberry fruits were dried in an ALPHA 1-2LD PLUS lyophilizer (Beijing, China), then ground and sifted for homogenization and stored at -80°C. The finely ground mulberries (3 kg) were extracted with 300 L of methanol-hydrochloric acid (999:1) at 37°C under the condition of ultrasonic power 200 w and frequency 40 kHz for 30 min. This process was repeated three consecutive times. The extracts is then mixed and evaporated into syrup using RE-2000E rotatory evaporator (Xi'an Taikang Biotechnology Co., Ltd., Xi'an, China). The syrup was dissolved in 1000 mL of 0.01 mol/L HCl, and was defatted with 1200 mL of ethyl acetate. The water solution was subjected to column chromatography of Amberlite XAD-7 orderly eluted with water and methanol-formic acid (9:1) [18] [19]. The methanol-formic acid elution solution was evaporated into syrup and freeze-dried to yield 403 g of MAEs. The MAEs stored at -80°C for further use.

2.3. pH Difference Method Analysis of Total Anthocyanins Content

Reference to previous study [20], the MAEs was diluted 10-fold with hydrochloric acid-sodium chloride of pH 1.0 and pH 4.5, respectively. The absorbance of MAEs diluents was measured at wavelengths of 520 nm and 700 nm by 755b UV-Vis spectrophotometer (Xi'an Heb Biotechnology Co., Ltd, Xi'an, China), respectively. The absorbance (*A*) was calculated by the Equation (1):

$$A = (A_{520 \text{ nm}} - A_{700 \text{ nm}})_{\text{pH 1.0}} - (A_{520 \text{ nm}} - A_{700 \text{ nm}})_{\text{pH 4.5}} \quad (1)$$

The total anthocyanin content of MAEs was calculated according to the formula derived by Wrolstad *et al.* [21]:

$$\text{Total anthocyanin content (mg}\cdot\text{g}^{-1}) = \frac{A \times MW \times DF \times V \times 1000}{\epsilon \times l \times M} \quad (2)$$

where MW is the molecular weight of cyclamine-3-glucoside (449.2), DF is the dilution, ϵ is the molar absorptivity (29,600), l is the optical path (1.0 cm), V is the total volume of MAEs diluents (1.0 ml), and M (0.01 g) is the weight of the MAEs.

2.4. Electrospray Ionization Mass Spectrometry (ESI-MS) and Ultra-Performance Liquid Chromatography (UPLC) Analysis of Anthocyanins

Quantifications of anthocyanin monomers were performed on a LC-MS/MS (Thermo Fisher Scientific Inc., SAN JOSE, Calif.) with a MRM mode. The LC-MS analysis was carried out using a LC-MS system comprising of a LC (2010, Finnigan, USA) and LCQ Fleet Ion Trap LC/MS (LCQ Fleet, Thermo Fisher Scientific). Mass spectrum condition: positive ion scanning (ESI⁺, m/z 200 ~ 1200), the collision induced dissociation voltage was 150 V; capillary voltage: 3.8 kV; taper hole voltage: 30 v. Voltage of photoelectric multiplier: 650 V; Ion source temperature: 120°C; Take off the solvent temperature: 250°C.

Separation of anthocyanins in MAEs was on a 2.1 × 150 mm, 1.7 μm ACQUITY UPLC BEH C18 column (Waters, UK) analyzed by UPLC using a ACQUITY UPLC H-Class system with a 2996 PDA detector (Waters, Milford, MA, USA). Anthocyanin standards (Sigma-Aldrich, Shanghai, China) were used for quantitative determination.

2.5. Animals and Experimental Design

All animal studies were conducted in accordance with Chinese National Guidelines for the Care of Laboratory Animals and approved by the Animal Ethics Committee of the Laboratory Animal Center, Fujian University of Traditional Chinese Medicine (Fujian, China). Four-week-old SPF-grade male Wistar rats were purchased from Shanghai slac laboratory animal Co., Ltd. with animal production license No.: SCXK (HU) 2012-0002. All animals were housed in an SPF barrier system of the Laboratory Animal Center, Fujian University of Traditional Chinese Medicine. Breeding conditions were as follows: temperature 20°C ~ 25°C, relative humidity 40% ~ 60%, 12 h day/night alternation.

Rats were fed with a rodent diet and had tap water *ad libitum*. After a week of adaptive feeding, the rats were randomly divided into five groups. The control group (Control), model group (NDEA), NDEA + low-dose MAEs (MAEs-75), NDEA + high-dose (MAEs-150), and NDEA + cyanidin-3-glucoside (C3G-150). The animal experiment design is shown in **Table 1**. Body weights were measured weekly. Twenty-four hours after the last MAEs administration, the rats were anesthetized by ip of pentobarbital sodium (40 mg·kg⁻¹) for collection of

Table 1. Sequences of primers used for qRT-PCR analysis.

UniProt ID	Gene symbol	Forward primer (5'-3')	Reverse primer (5'-3')
AF304364.1	Nrf2	ACCCACCGCCTGGGTTTCAGT	TGTGCCCTTGAGCTGGCGAC
NM_173295.1	UGT2b1	TTAGACCTGGAGCCTGTGGAAA	GCCGAAGATACAAGAACCGTGA
NM_017014.1	GST	AGAGCAGCCGCTACCTCTCAAC	CCAATGTGGACAGGTCTCCCT
NM_001159613.1	NQO1	CCAGCAGCCCGCCAATCTG	AGGTCCGACACGGCGACCTC
NM_182864.2	Keap1	TGATGGACAAACCCAACTCA	CACTGGACAGGAAACCACCT
NM_001004027.1	HO-1	AGGCTGAGAATGCCGAGTTC	TGTGGTACAAGGACGCCATC
NM_214022.1	TNF- α	GGAAGTGGCAGAGGAGGCGC	CCCCGCCACGAGCAGGAATG
NM_199267.2	NF- κ B	TGATGACATACTCCACAAG	CAATATCCCCAGACCTAAG
S67722.1	COX-2	ACCAGCAGTTCAGTATCAGA	AAGTGAGCAAGTCCGTGTTC
NM_017008.4	GAPDH	CGGAGTCAACGGATTTGGTCTGAT	AGCCTTCTCCATGGTGGTGAAGAC

blood samples and blood samples were centrifuged at 3000 rpm at 4°C for 10 min to obtain serum. Livers were fully excised from the animal and accurately weighed [6]. The size and number of liver nodules were measured as described earlier [3]. Liver tissue was collected for biochemical and histopathological analyses.

2.6. Biochemical Assays

The levels of serum aspartate transaminase (AST), alanine aminotransferase (ALT), γ -glutamyl transpeptidase (GGT), total bilirubin (TBiL), alkaline phosphatase (ALP), Blood Urea Nitrogen (BUN), and creatinine were detected according to the instructions of commercial kits (Nanjing Jiancheng Bioengineering Engineering Institute, Nanjing, China) using Infinite M200 PRO Varioskan Flash (TECAN, Switzerland). The levels of serum alpha fetal protein (AFP), carcino embryonic antigen (CEA), TNF- α , interleukin-6 (IL-6), interleukin-10 (IL-10) and interferon- γ (IFN- γ) were measured by ELISA kit (CUSABIO Biotech Co., Ltd., Wuhan, China) according to manufacturer's instructions.

Precisely 1 gram of liver was taken, rinsed with cold PBS, pH 7.4, dried with paper wipe, and placed in a 10-mL centrifuge tube. Then the tissue homogenate was centrifuged at 3500 rpm for 15 min. The supernatant was collected in a 10-mL centrifuge tube (namely 10% liver homogenate), immediately frozen in liquid nitrogen and stored at -20°C until use [6]. The contents of malondialdehyde (MDA), glutathione (GSH), and the activities of superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GSH-Px) were assayed using standard commercially available kits (Nanjing Jiancheng Bioengineering Engineering Institute, Nanjing, China) according to the manufacturer's instructions. The serum thiobarbituric acid-reactive substances (TBARS) concentrations were assayed by rat TBARS ELISA kit (QIYI Biological technology Co., Ltd., Shanghai, China). The levels of Glutathione S-transferase (GST) and UDP-glucuronosyltransferase 2B1 (UGT2b1) in the liver homogenates were measured with ELISA kit (CUSABIO Biotech Co., Ltd., Wuhan, China).

2.7. Histological Examination

Hepatic tissues from each group were washed with cold physiological saline and rapidly excised after the rats were sacrificed. The tissue samples were immediately fixed in 4% phosphate-buffered paraformaldehyde for 48 h, then dehydrated in a graded ethanol series, cleared in xylene and embedded in paraffin. Sections (4 μm thick) were stained with hematoxylin and eosin (H&E) according to published methods [22]. The samples were also sectioned and Masson's trichrome staining was performed as previous studies [23]. For histological analysis, the tissue sections were photographed with a microscope (Nikon E200, Nikon Corp., Japan).

2.8. Immunohistochemical Analysis

The proteins expression of inflammatory markers in liver tissue was detected by the SABC immunohistochemical method. The main steps are as follows: paraffin section (4 μm thick) was de-waxed and incubated with sodium citrate buffer (pH 6.0) at 37°C for 5 min (twice), then washed with 0.1 M PBS for 5 min (3 times), incubated with 3% hydrogen peroxide at 37°C for 10 min, washed with PBS for 5 min (3 times) and placed in a blocking solution at room temperature for 1.5 - 3 h. The sections were incubated with rabbit polyclonal antibodies, NF- κB , COX-2, and TNF- α (Cell Signaling, Danvers, MA, USA) with TBS and Tween 20 overnight at 4°C. Then sections were washed with PBS and incubated with HRP-labeled sheep-anti-rabbit secondary antibody at 37°C for 1 - 2 h followed by streptavidin-biotin-peroxidase at room temperature for 30 min. The slides were washed and the immunoprecipitation was visualized by treating with 3, 3'-diaminobenzidine for color development for 25 min. Then slides were counterstained with hematoxylin and the brown color signifying the presence of antigen bound to antibody was detected by light microscopy. For the negative control, TBS was used instead of a primary antibody. From ten randomly selected sections of each slide, 500 cells were counted. The percentage of positive cells for each group was calculated using the Image-Pro Plus 6.0 image analysis system.

2.9. Western Blot Analysis

Liver tissue was lysated in RIPA lysis buffer containing protease inhibitor cocktail protease inhibitors and phosphatase inhibitors according to the M-PER (R) Mammalian Protein Extraction Reagent (Thermo Scientific, Fair Lawn, NJ, USA). The protein extraction was collected and the concentrations were quantified using a BCA kit (Biotechnology Development Co., Ltd., Beijing, China). The same amount of proteins was separated using 10% SDS-polyacrylamide gel and transferred to polyvinylidene fluoride (PVDF) membranes. After blocked with 5% skim milk for 1 h at room temperature, membranes were incubated with various antibodies against Nrf2 (1:2000), NQO1 (1:1000), HO-1 (1:2000), Keap1 (1:500), NF- κB (1:4000), COX-2 (1:500), TNF- α (1:1000), and β -actin (1:2000) which purchased from Cell Signaling Technology (Danvers, MA, USA) or Santa

Cruz Biotechnology (Santa Cruz, CA, USA) overnight at 4°C. Then membranes were washed and exposed to HRP-conjugated secondary antibodies (1:10,000) at room temperature for 1 h. Immunoreactive bands were detected by enhanced cheiluminescence solution (CUSABIO Biotech Co., Ltd., Wuhan, China) and exposed to X-ray film using the Bio-Rad Chemi Doc XRS imaging System. The immunoreactive bands were visualized and quantified using Quantity One software and normalized to β -actin.

2.10. Real-Time PCR Analysis

Total RNA was extracted from liver tissue using a TRIzol reagent (Invitrogen, Thermo Fisher Scientific Inc., Beijing, China) according to the manufacturer's instructions. Real-time PCR was performed using the SYBR Green Kit (Takara Biomedical Technology Co., Ltd., Beijing, China) on the ABI Step One RT-PCR system. Primers designed with Primer Premier 5 and Beacon Designer 8.1 was listed in **Table 2**.

2.11. Statistical Analysis

Data was presented by mean standard deviation (SD). Data from study was dealt with SPSS 21.0 statistical package. Statistical analysis was performed by one-way analysis of variance (ANOVA) with Tukey's multiple comparisons or Student's t-test. A P-value of <0.05 was considered statistically significant. The GraphPad Prism 6.01 was used for the graphical evaluations.

3. Results and Discussion

3.1. PR-01 Mulberry Anthocyanins Extract Component Analysis

The total anthocyanins content of the PR-01 mulberry anthocyanins rich-extract was $(718 \pm 8.9) \text{ mg}\cdot 100\text{g}^{-1}$. UPLC-ESI-MS analysis showed the pelargonidin 3-glucoside (P3G), C3R and C3G is the main anthocyanins (**Figure 1**), which was in agreement with that reported by Song [17] and Huang [24]. UPLC analysis

Table 2. Experimental design of the study.

Groups	No. of animals	Ip administration	Gavage administration
		Every Tuesday and Friday 8:30 am from 2 weeks to 12 weeks	Every day 10:00 am from 1 weeks to 18 weeks
Control	10	Physiological saline	Distilled water
NDEA	12	25 mg/kg NDEA	Distilled water
C3G-150	12	25 mg/kg NDEA	150 mg/kg C3G
low-dose MAE (MAEs-75)	12	25 mg/kg NDEA	75 mg/kg MAEs
high-dose MAEs (MAEs-150)	12	25 mg/kg NDEA	150 mg/kg MAEs

Ip, intraperitoneal injection; C3G, cyanidin-3-glucoside; MAEs, mulberry anthocyanins extract; NDEA, *N*-nitrosodiethylamine. Gavage doses of MAEs intervention groups were calculated by C3G contents. The MAEs and C3G were dissolved in distilled water.

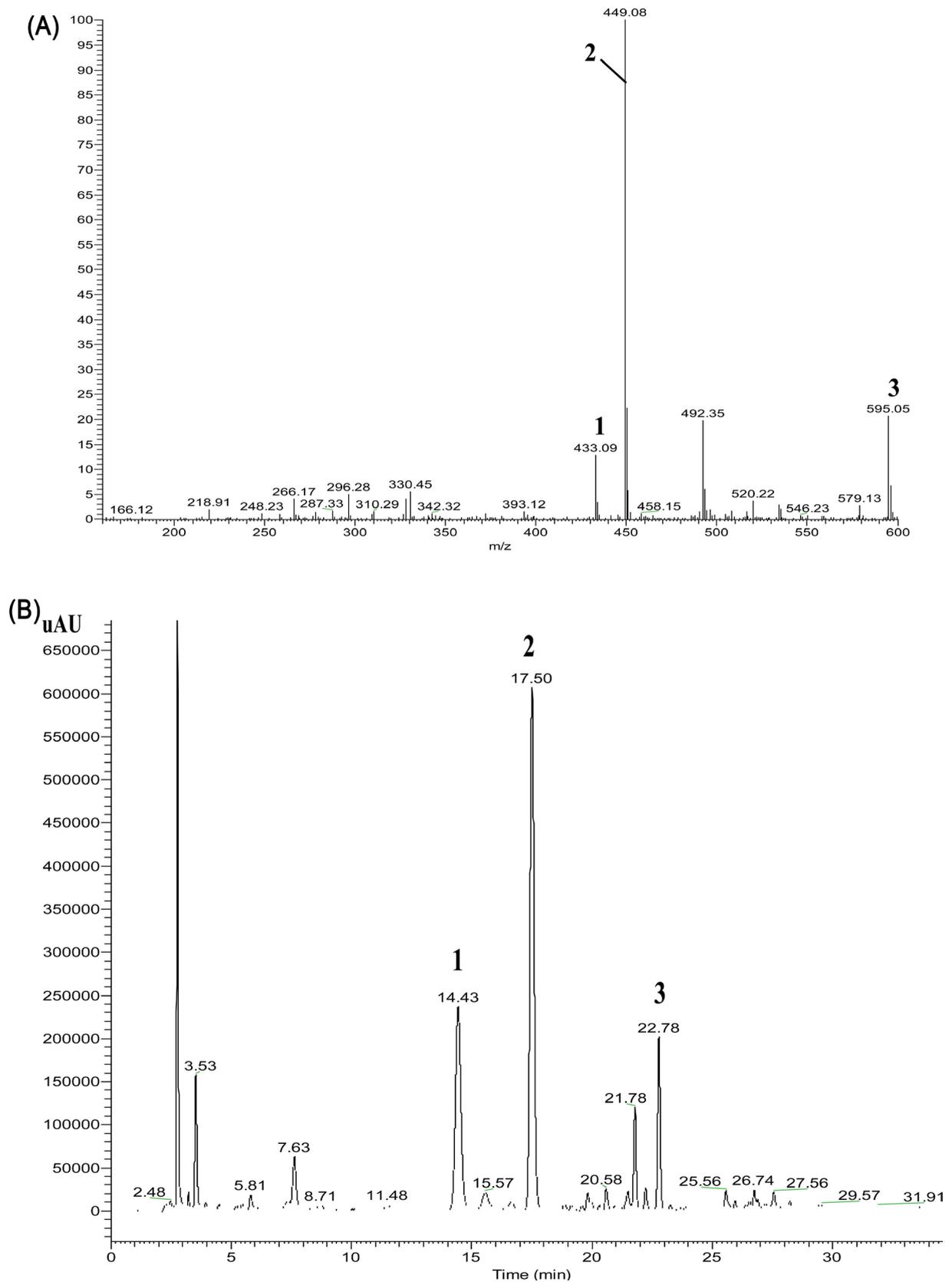


Figure 1. The mass spectrogram and chromatogram of the PR-01 mulberry anthocyanins rich-extract. (A) ESI-MS of MAEs; (B) chromatogram of MAEs. The three peaks correspond to (1) pelargonidin 3-glucoside; (2) C3G; (3) C3R.

showed that the content of P3G, C3R and C3G in the MAEs is (75.4 ± 6.5) $\text{mg}\cdot\text{g}^{-1}$, (105.8 ± 4.1) $\text{mg}\cdot\text{g}^{-1}$, and (593.0 ± 12.6) $\text{mg}\cdot\text{g}^{-1}$, respectively. The mass spectrum of MAEs is shown in **Figure 1**, while the characterization of chemical constituents of MAEs is presented in **Table A1**.

3.2. Alterations in Body Weight, Food Intake, and Liver Index

As shown in **Table 3**, the body weight of rats in NDEA group was significantly reduced subsequent to NDEA compared with that in control group ($P < 0.01$), while the weight gains of animals treated with MAEs and C3G-150 were similar ($P > 0.05$). With regard to average daily feed consumption, the NDEA group consumed a lower amount of the feed than the control group ($P < 0.01$). However, after treatment with MAEs and C3G, the daily feed intakes were significantly increased than the NDEA group ($P < 0.05$). Compared to NDEA group, the relative liver weight was remarkably decreased in the MAEs-75 and MAEs-150 groups by 23.64% and 28.37%, respectively after MAEs treatment ($P < 0.01$).

3.3. Changes in Liver Function Biomarkers

NDEA causes liver damage and consequently releases liver enzymes in its selected dose [25]. The past research suggested that mulberry water extracts may serve as liver protective agents [4] [25]. In this study, the elevated activity of ALT, AST, Tbil, ALP, and GGT are indicative of poor hepatic function in the NDEA alone treated animals compared to the control animals (**Table 4**), as reported by Latief *et al.* [26]. After treated with MAEs, the levels of the elevated liver function enzymes above mentioned were significantly restored in MAEs-75 and MAEs-150 group animals, suggesting that MAEs has an efficacy function on liver protection. Pretreatment with the higher dose of MAEs ($150 \text{ mg}\cdot\text{kg}^{-1}$) also reversed the NDEA-induced increase in serum BUN and creatinine to normal levels (**Table 4**). In addition, the serum CEA and AFP levels were obviously higher in NDEA group than that of the control group ($P < 0.05$), while $75 \text{ mg}\cdot\text{kg}^{-1}$ MAEs and $150 \text{ mg}\cdot\text{kg}^{-1}$ MAEs treatment remarkably reduced the serum CEA and AFP levels by 25.99%, 39.32%, 32.57%, and 49.24%, respectively compared with the NDEA group (**Table 4**, $P < 0.05$). Pretreatment with higher dose of MAEs ($150 \text{ mg}\cdot\text{kg}^{-1}$) appeared to be more efficient in inhibiting NDEA-induced

Table 3. Effects of MAEs on body weight, feed intake, and liver index in NDEA-induced hepatocarcinogenesis rats.

Groups	Initial Body weight (g)	Final Body weight (g)	Body weight gain (g)	Feed intakes (g/day/rat)	Relative liver weights [#]
Control	141.25 ± 4.87 ^a	592.42 ± 28.54 ^a	451.17 ± 28.98 ^a	27.06 ± 3.48 ^a	2.92 ± 0.43 ^b
NDEA	141.42 ± 4.34 ^a	514.83 ± 38.32 ^c	373.42 ± 38.11 ^c	21.99 ± 1.45 ^c	4.23 ± 0.64 ^a
C3G-150	141.92 ± 7.97 ^a	531.33 ± 64.91 ^{bc}	389.42 ± 64.40 ^{bc}	24.92 ± 2.03 ^b	3.79 ± 0.62 ^a
MAEs-75	141.92 ± 6.52 ^a	554.42 ± 42.68 ^b	413.08 ± 43.53 ^b	25.28 ± 2.34 ^{ab}	3.23 ± 0.49 ^b
MAEs-150	141.08 ± 7.47 ^a	556.00 ± 21.86 ^b	414.75 ± 25.21 ^b	25.08 ± 3.50 ^{ab}	3.03 ± 0.38 ^b

Note: [#]Relative liver weights equal liver weight/body weight × 100. Values expressed as mean ± S.D. (n = 8 - 10). Values with different superscript letters (a, b, c, d, e) within cultivar are significantly different.

Table 4. Effects of MAEs on serum hepatic enzymes in NDEA-induced hepatocarcinogenesis rats.

Groups	Control	NDEA	C3G-150	MAEs-75	MAEs-150
ALT (U/L)	40.70 ± 8.85 ^d	85.63 ± 17.78 ^a	61.50 ± 19.63 ^b	54.80 ± 11.67 ^{bc}	46.10 ± 11.14 ^{cd}
AST (U/L)	83.10 ± 12.71 ^c	173.13 ± 47.09 ^a	113.80 ± 23.70 ^b	98.10 ± 21.56 ^{bc}	93.20 ± 13.50 ^c
TBiL (mg/dl)	2.32 ± 0.26 ^c	4.45 ± 0.76 ^a	3.02 ± 0.41 ^b	2.89 ± 0.31 ^b	2.30 ± 0.17 ^c
ALP (U/L)	58.60 ± 9.16 ^b	90.63 ± 14.93 ^a	78.10 ± 22.81 ^a	60.20 ± 17.03 ^b	53.80 ± 9.37 ^b
GGT (U/L)	30.26 ± 3.62 ^d	115.43 ± 14.37 ^a	68.24 ± 6.08 ^c	50.55 ± 3.25 ^c	39.32 ± 6.54 ^{cd}
BUN (mmol/L)	5.40 ± 1.07 ^b	8.87 ± 1.13 ^a	7.12 ± 2.67 ^b	6.02 ± 2.26 ^b	5.81 ± 1.51 ^b
Creatinine(μmol/L)	23.96 ± 9.58 ^e	68.36 ± 9.86 ^a	47.15 ± 10.54 ^{bc}	33.50 ± 8.85 ^{de}	31.43 ± 14.44 ^{de}
CEA (ng/ml)	2.16 ± 0.44 ^d	7.35 ± 0.57 ^a	6.76 ± 1.18 ^{ab}	5.44 ± 1.85 ^{bc}	4.46 ± 1.87 ^c
AFP (ng/ml)	22.27 ± 6.18 ^c	67.54 ± 6.91 ^a	53.47 ± 16.69 ^a	45.54 ± 17.76 ^b	34.28 ± 13.2 ^{bc}

Values with different superscript letters (a, b, c, d, e) within cultivar are significantly different.

liver damage in rats.

3.4. Histopathological Study of Liver Samples

As one of the most important environmental hepatotoxin and carcinogen, NDEA can cause severe liver damage and lead to severe alterations in the lobular architecture and hampers liver functioning [26]. During the experiment, 2 rats died of severe hepatic tumor pathogenesis at 16th and 18th week in the NDEA group. No death was visible in other group animals. The results of this study shown that low-dose multiple injection of NDEA (25 mg·kg⁻¹ of NDEA twice a week) significantly increased the nodule incidence and the apparent HCC incidence (100.0%, 50.0%, respectively) and only a low mortality rate (16.7%) for 18 weeks in the NDEA group, indicating successful induction of liver cancer model. However, there was significant decrease in hepatic nodules incidence, maximum diameter, average number in the MAEs and C3G-150 group rats as compared with NDEA group animals (Table 5, $P < 0.05$). In particularly, treatment of 150 mg·kg⁻¹ MAEs prior to C3G showed a significant reduction of the nodule incidence (41.7%) compared to the NDEA group animals (100%).

As shown in Table 5, treatment of 150 mg·kg⁻¹ MAEs prior to C3G showed a significant reduction of the apparent HCC incidence (8.3%) compared to the NDEA group animals (50.0%). The liver phenotype of rats in the control group was not significantly changed, while the liver size and structure of NDEA group rats were significantly changed (Figure 2). Apparently the liver structure of injured rats was characterized by cells necrosis, hemorrhage, scars and hepatic nodule like structure, abundant eosinophilic cells, basophilic cells and egg cells hyperplasia obviously accompanied by inflammatory cells infiltration [26], which were observed in the liver of NDEA group rats. However, MAEs (75 mg·kg⁻¹ or 150 mg·kg⁻¹) and 150 mg·kg⁻¹ C3G administration regress these changes significantly (Figure 2(B)). The results of masson-stained sections showed that rats developed liver fibrosis after NDEA administration. However, the degree of

Table 5. Effects of MAEs on hepatic neoplasm-related lesions in NDEA-induced hepatocarcinogenesis rats.

Hepatic neoplasm-related lesions		Control (n = 10)	NDEA (n = 10)	C3G-150 (n = 11)	MAEs-75 (n = 12)	MAEs-150 (n = 12)
Macroscopic lesions	Nodule incidence [#]	0 (0)	10 (100)	8 (72.7)	7 (58.3)*	5(41.7)**
	Max nodule diameter (mm) [▲]	—	18.73 ± 7.13 ^a	9.42 ± 5.54 ^b	4.63 ± 2.46 ^c	1.87 ± 1.79 ^c
	Average nodule number [#]	ND	71.33 ± 62.11 ^a	48.10 ± 10.80 ^b	17.70 ± 3.42 ^d	6.13 ± 5.22 ^d
Microscopic lesions	AHF incidence	0 (0.0)	1 (10.0)	2 (16.7)	3 (25.0)	2 (16.7)
	HA incidence	0 (0.0)	3 (30.0)	4 (33.3)	3 (25.0)	2 (16.7)
	HCC incidence	0 (0.0)	5 (50.0)	3 (25.0)	2 (16.7)*	1 (8.3)**
	Total tumor incidence	0 (0.0)	8 (80.0)	7 (58.3)	5 (41.7)	3 (25.0)*

Data are means ± SDs or n (%). Values with different superscript letters (a, b, c, d) within cultivar are significantly different. * $P < 0.05$ and ** $P < 0.01$ compared with NDEA group by Fisher's exact test. [#]Visible nodules (diameter ≥ 1 mm). [▲]Max nodule diameter, mean maximum nodule diameter. ND, not detectable; AHF, altered hepatic foci; HA, hepatic adenoma; HCC, hepatocellular carcinoma.

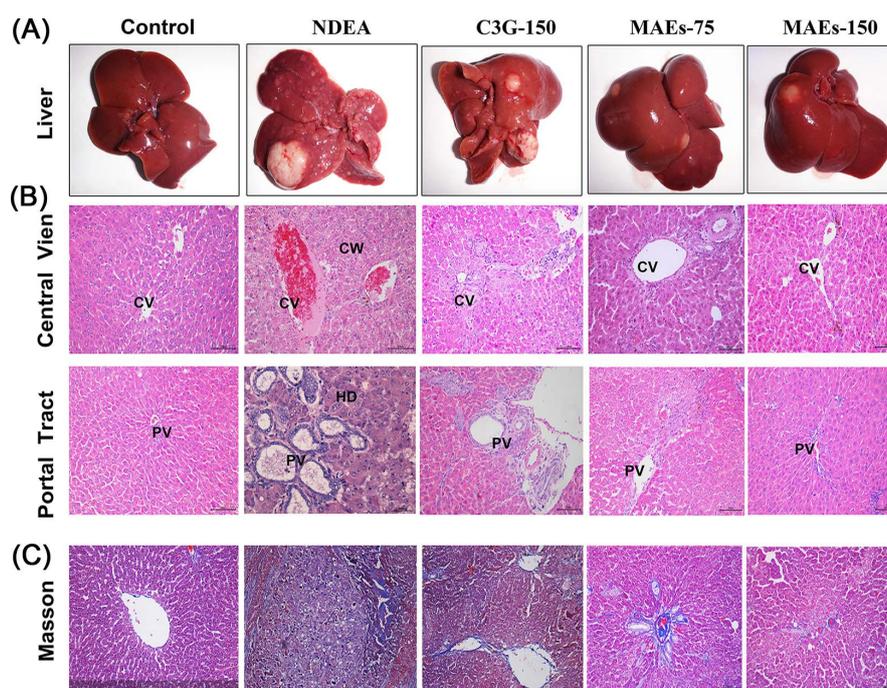


Figure 2. Effects of MAEs on hepatic pathological changes in NDEA-induced hepatocarcinogenesis rats. (A) Liver surface; (B) Liver sections with haematoxylin and esion (H & E, 200×). Basophilic cell focus (white arrow) is an altered hepatic focus. At the center of the lobule is the central vein (CV). PV, portal vein. (C) Liver sections with Masson staining (200×).

fibrosis in NDEA group animals was more severe compared to rats treated with MAEs or C3G (**Figure 2(C)**). Cellular vacuolization was also reduced in the MWEs-treated groups (**Figure 2(C)**).

These results suggested that MAEs is an effective chemopreventive agent for preventing or delaying NDEA-induced hepatocarcinogenesis in rats, among which the high dose of MAEs has the best effect and is superior to C3G.

3.5. Antioxidative Effect of MAEs on TBARS and Related Antioxidant Enzymes Activity

Excess reactive oxygen species produced in the processes of NDEA metabolic activation can interfere with the body's oxidation system, leading to oxidative damage and tissue carcinogenesis, which plays an important role in the pathogenesis of liver cancer [26]. TBARS concentrations were used as markers of oxidative stress [24]. In this study, compared with the control group, the TBARS level of liver tissue in the NDEA group was remarkably increased ($P < 0.001$) 1.88-fold, which is consistent with earlier reports [13] [27]. Compared with the NDEA group, the hepatic tissue TBARS were significantly reduced in the C3G-150, MAEs-75 and MAEs-150 groups by 39.01%, 40.85%, and 44.52%, respectively (Table 6, $P < 0.001$). Interestingly, these results showed that 150 mg·kg⁻¹ MAEs treatment almost restored the TBARS to the normal level. On this basis, we further detected the effect of MAEs on antioxidant capacity. The activities of antioxidant enzymes, GSH, GSH-Px, CAT and SOD in NDEA group were remarkably lower than that in the control group ($P < 0.01$), while these activities were dramatically promoted by MAEs and C3G ($P < 0.05$). Among which, the 150 mg·kg⁻¹ MAEs treatment rats had the highest levels of antioxidant enzymes in livers. After NDEA treatment, MDA content was obviously elevated in liver of NDEA group rats, which was approximately 2.44-fold higher than in control group rats ($P < 0.01$), and significantly decreased after MAEs treatment (Table 6, $P < 0.05$). Our results showed that NDEA administration changes the antioxidant status, leading to oxidative stress, Whereas MAEs can inhibit lipid peroxidation and enhance antioxidant capacity, so the prevention effect of MAEs against NDEA-induced hepatocarcinogenesis in rats may be closely related to its antioxidant activity. Chen *et al.* [13] [18] [27] also found that anthocyanins and water extracts from mulberry fruit can scavenge free radicals and shown concentration-dependent antioxidant activity *in vitro*.

3.6. Effect of MAEs on Liver GST and UGT2b1

NDEA's induction increased microsomal phase I metabolizing enzymes with a

Table 6. Effects of MAEs on TBARS and activities of related antioxidant enzymes in livers of NDEA-induced hepatocarcinogenesis rats.

Groups	TBARS (nmol/mg protein)	GSH (mg/g)	GSH-Px (U/mg)	SOD (U/mg)	CAT (U/mg)	MDA (nmol/mg)
Control	241.67 ± 11.61 ^c	4.38 ± 0.71 ^a	387.53 ± 23.72 ^a	172.09 ± 14.28 ^a	35.89 ± 8.58 ^a	4.73 ± 1.07 ^c
NDEA	454.74 ± 25.43 ^a	1.23 ± 0.34 ^e	293.02 ± 53.17 ^b	94.56 ± 20.70 ^c	14.56 ± 3.10 ^d	11.55 ± 3.66 ^a
C3G-150	277.36 ± 30.24 ^b	1.94 ± 1.03 ^d	363.57 ± 28.85 ^a	127.47 ± 24.65 ^b	23.26 ± 5.35 ^c	8.47 ± 2.18 ^b
MAEs-75	268.96 ± 32.90 ^{bc}	2.85 ± 0.70 ^c	364.78 ± 32.04 ^a	147.45 ± 36.13 ^b	28.21 ± 7.27 ^{bc}	5.95 ± 2.33 ^c
MAEs-150	252.30 ± 23.09 ^{bc}	3.61 ± 0.25 ^b	380.94 ± 37.81 ^a	170.77 ± 18.34 ^a	32.30 ± 5.55 ^{ab}	5.30 ± 1.00 ^c

The data are the mean ± SD from 8 samples for each group and at least three independent measurements. Values with different superscript letters (a, b, c, d) within cultivar are significantly different. TBARS, thiobarbituric acid-reactive substances; GSH, Glutathione; GSH-Px, glutathione peroxidase; SOD, superoxide dismutase; CAT, catalase; MDA, malondialdehyde.

simultaneous decrease in the Phase II detoxifying enzyme [6] [28]. GST and UGT2b1 play an important role in the detoxification and excretion of toxins, carcinogens, which may actively modulated by Nrf2 and the antioxidant response element [29]. In this study, administration of NDEA, the significant reduction in UGT2b1 and GST activity in livers of NDEA group were observed (**Figure 1(A)**, $P < 0.05$). Whereas the levels of GST and UGT2b1 in livers of MAEs-75 and MAEs-150 group rats were remarkably increased as compared to the NDEA group. What's more, the hepatic GST and UGT2b1 levels of MAEs-150 group were obviously higher than that of C3G-150 group ($P < 0.05$), and closed to the normal level. The result of the qRT-PCR also showed that NDEA induced GST and UGT2b1 mRNA expressions and MAEs repressed these alterations (**Figure 3(B)**). These results suggested MAEs can inhibit tumor development by stimulating the activity of phase II detoxification enzyme.

3.7. MAEs Activated the Nrf2 Signaling Pathway and Its Downstream Detoxification Enzymes

Nrf2-mediated antioxidant, detoxification enzymes and anti-inflammatory signaling are through Nrf2-ARE pathways to protect organisms against cellular damage caused by oxidative stress [28]. Yan *et al.* [15] reported that consumption of mulberry anthocyanin extract maintained Nrf2, HO-1, and p38 MAPK stimulation that were involved in oxidative stress modulation. The results of this study shown that Nrf2 activation in livers of rats was induced by 150 mg·kg⁻¹ MAEs, accompanied by the stimulation the mRNA expressions of NQO-1, Keap1, and HO-1 (**Figure 3(B)**, $P < 0.05$). Similar findings were observed in the measurements of Nrf2, Keap1, HO-1, and NQO-1 protein levels. Compared with the control and NDEA groups, treatment of 150 mg·kg⁻¹ MAEs caused significant changes in the protein expressions of Nrf2, Keap1, HO-1, and NQO-1 in the MAEs-150 group ($P < 0.05$), but less obvious changes were found in MAEs-75 group (**Figure 3(C)**). Combining these results, it was concluded that MAEs may activate the Nrf2 signaling pathway and induce Nrf2-mediated antioxidant enzymes, further activate the expression of downstream phase II detoxifying GST and UGT2b1, which then accelerated the elimination of the metabolites of NDEA, thereby achieving the goal of detoxification.

3.8. MAEs Attenuated the NDEA-Induced Inflammatory Response

Studies showed that the main cause of NDEA-induced liver cancer is inducing chronic inflammatory response and abnormal repair after liver injury [30] [31]. Hassimotto *et al.* studies showed that administration of wild mulberry extract suppressed carrageenan-induced acute inflammation [7]. As shown in Fig 3A, relative to the NDEA group (85.26 pg·mL⁻¹, 97.60 pg·mL⁻¹), the IL-6 and TNF- α levels were remarkably inhibited by MAEs-75 (46.24 pg·mL⁻¹, 50.08 pg·mL⁻¹), MAEs-150 (37.93 pg·mL⁻¹, 36.75 pg·mL⁻¹), and C3G-150 (62.14 pg·mL⁻¹, 56.62 pg·mL⁻¹). The inhibition of IL-6 and TNF- α showed a dose-dependent relationship with MAEs (75 mg·kg⁻¹ and 150 mg·kg⁻¹). Besides, MAEs treatments

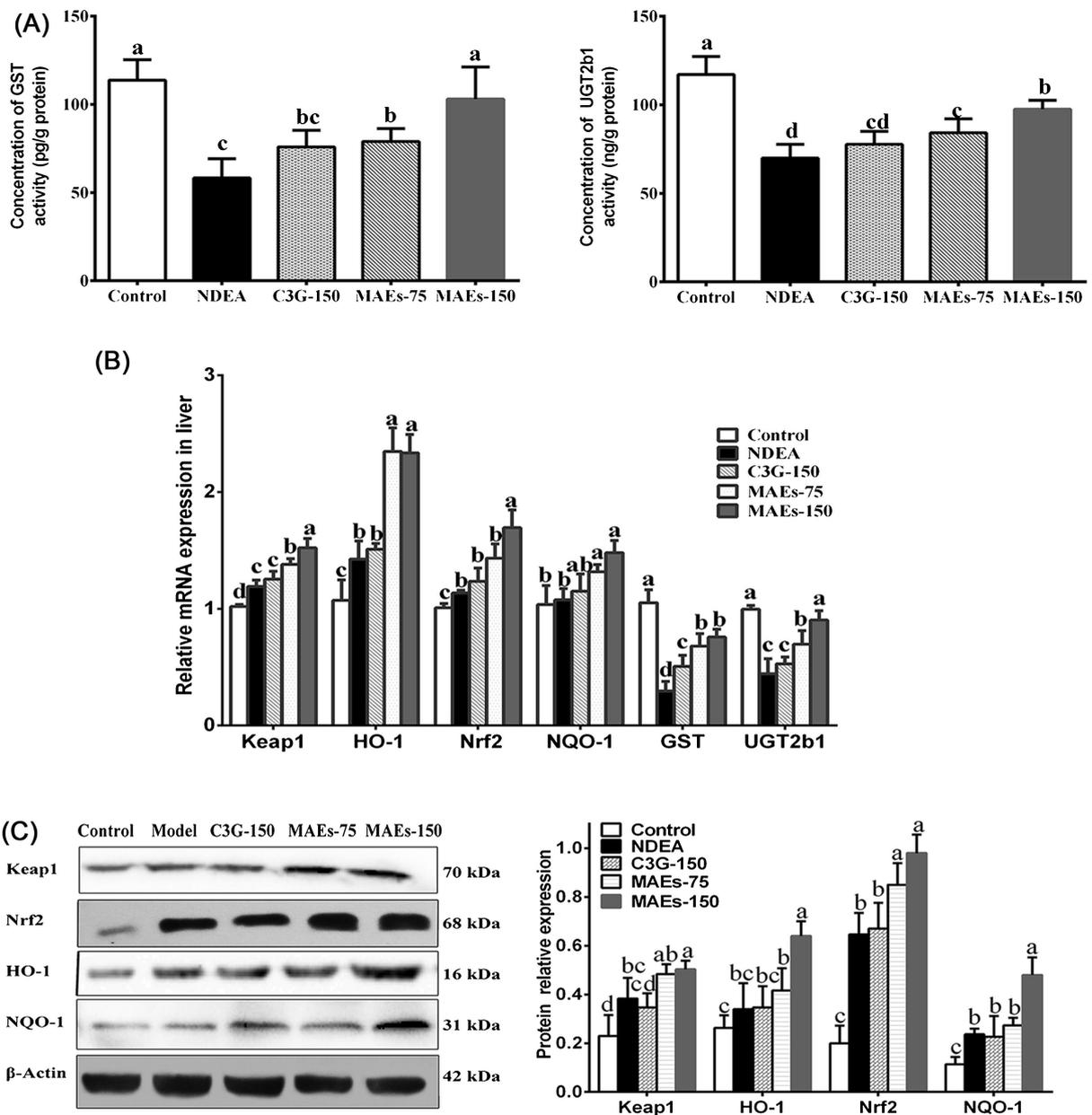


Figure 3. Effects of MAEs on the expression of Nrf2 and its downstream detoxification enzymes. (A) Effects of MAEs on GST and UGT2b1 levels in liver microsomes of rats exposed to NDEA. Values are expressed as concentration of the hepatic microsomal GST and UGT2b1 activity. GST, glutathione-S-transferase; UGT2b1, UDP glucuronosyl-transferase UGT2b1. (B) The mRNA relative expression levels of GST, UGT2b1, Keap1, HO-1, Nrf2, and NQO-1 in control and treatment groups. GAPDH was used as an internal control. (C) The representative immunoblots and protein levels of Keap1, HO-1, Nrf2, and NQO-1 in liver tissues in the rats of each group. Values with different superscript letters (a, b, c, d) within cultivar are significantly different among groups ($P < 0.05$).

increased anti-inflammatory cytokines IL-10 and IFN- γ in serum after NDEA treatment (Figure 4(A)).

NF- κ B, TNF- α , and COX-2 plays an important role in the development of inflammation [13] [24]. Thus, we further determined whether MAEs can reverse the effects on inflammation-related gene (TNF- α , NF- κ B, and COX-2) expression by qRT-PCR and Western blot. The result of qRT-PCR analysis revealed

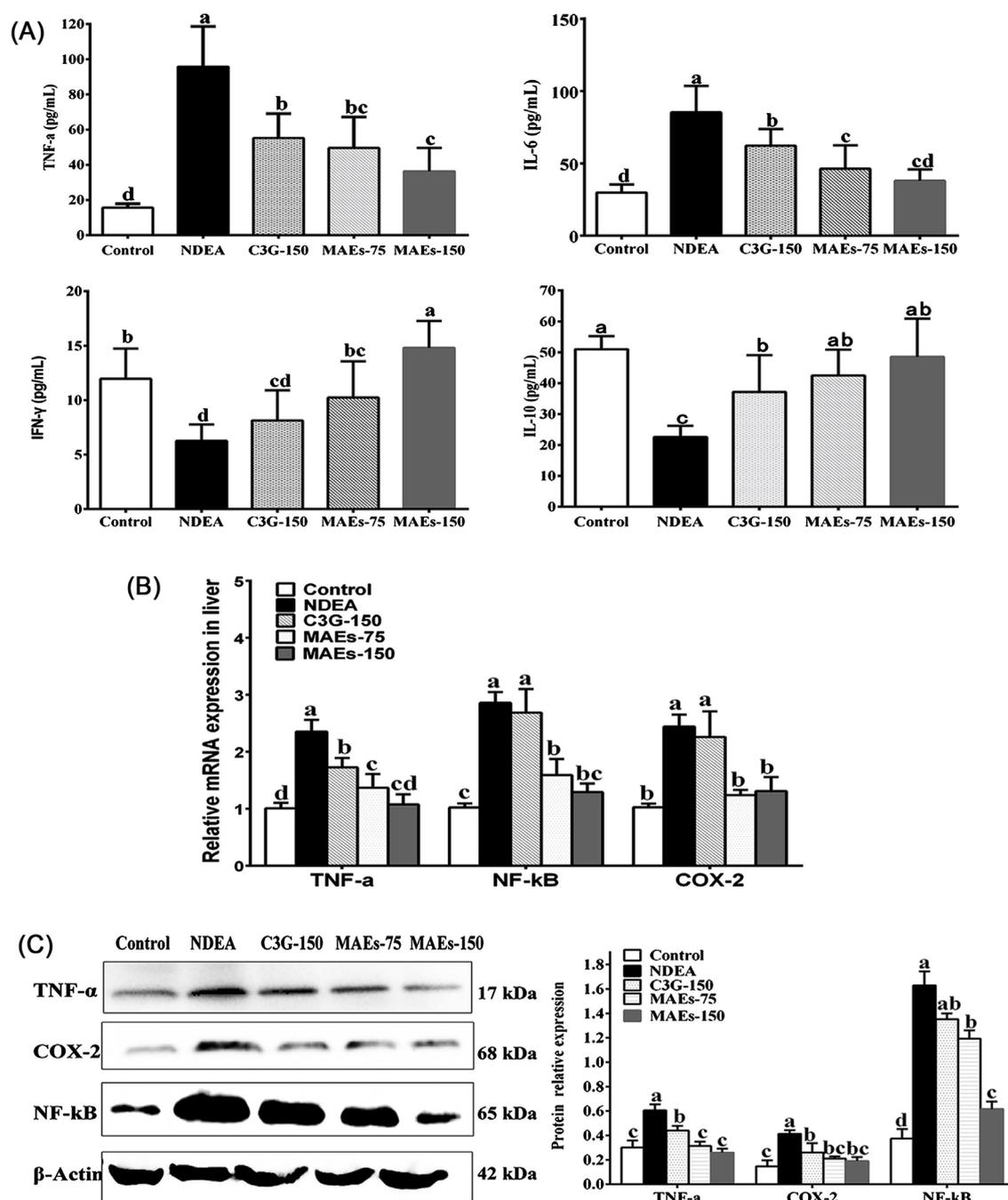


Figure 4. Effects of MAEs on markers of inflammation in NDEA-induced hepatocarcinogenesis rats. (A) The markers of pro-and anti-inflammatory cytokines in serum of each group rats. (B) The mRNA expression of TNF- α , NF- κ B, and COX-2 in livers of each group rats. GAPDH was used as an internal control; (C) Representative immunoblots of TNF- α , NF- κ B, and COX-2 and its protein levels in livers of each group rats. β -actin was used as an internal control. Values expressed as mean \pm SD. Values with different superscript letters (a, b, c, d) within cultivar are significantly different.

that, MAEs obviously decreased the TNF- α , NF- κ B, and COX-2 mRNA expression in the livers of MAEs-75 and MAEs-150 group rats, all of which were remarkably higher than that of C3G treatment rats (Figure 4(B)). These results indicate that MAEs reduced NDEA-induced TNF- α , NF- κ B, and COX-2 mRNA

expression in a dose-dependent manner. Similar findings were observed in the measurements of the TNF- α , NF- κ B, and COX-2 protein levels. Treatment with NDEA improved the protein expression levels of TNF- α , NF- κ B, and COX-2 by 2.01-, 4.35-, and 2.81-fold in NDEA group when compared with the control group, respectively (Figure 4(C)). However, administration of MAEs significantly reduced the levels of TNF- α , NF- κ B, and COX-2 proteins in MAEs-75 and MAEs-150 group rats (Figure 4(C), $P < 0.05$). Moreover, immunohistochemistry of NDEA-treated liver sections exhibited TNF- α , NF- κ B, and COX-2 positive cells in NDEA group. The amount of TNF- α , NF- κ B, and COX-2 positive cells in the liver, which was approximately 6.16-, 4.53-, and 15.92-fold higher in NDEA group rats than in control group rats ($P < 0.001$), reduced remarkably after MAEs (75 and 150 mg·kg⁻¹) treatment (Figure 5, $P < 0.01$).

The present study indicated that the NDEA-induced liver cancer may be related to the inflammatory reaction in rats, and MAEs may exert anti-inflammation by regulating the levels of inflammatory cytokines and attenuating the inflammatory

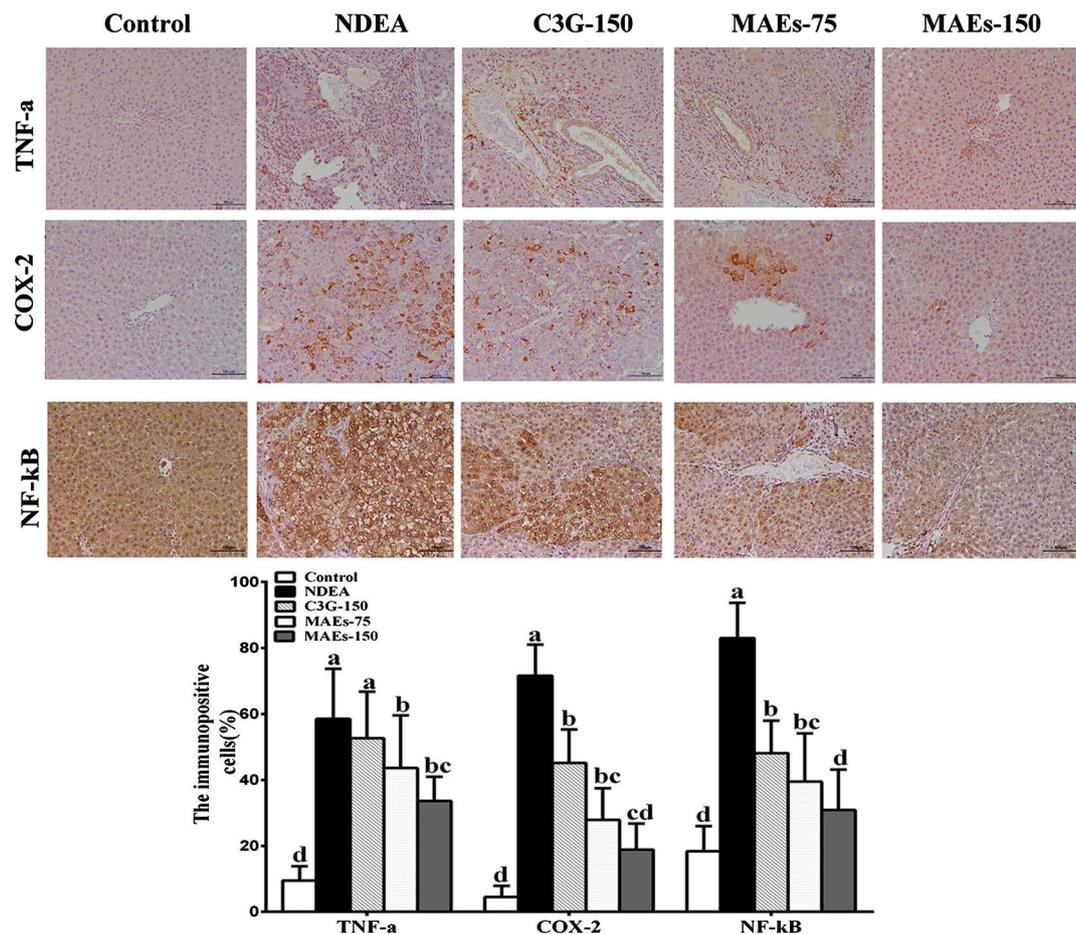


Figure 5. Representative immunohistochemical staining of TNF- α , COX-2, and NF- κ B in the liver of control and experimental animals at $\times 200$ original magnification. Bar graph represents average percentage of positive stained cells of TNF- α , COX-2, and NF- κ B in the liver of control and experimental groups. Results are expressed as mean \pm SD. Values with different superscript letters (a, b, c, d) within cultivar are significantly different.

mediators COX-2 through NF- κ B inhibition pathway.

4. Conclusion

In vivo studies, we present evidence that MAEs of mulberry variety PR-01 fruits exhibit prevention of NDEA-induced liver damage, fibrosis and hepatocellular carcinoma via reducing liver function enzymes, decreasing lipid peroxidation, promoting antioxidant enzymes, and increasing the hepatic phase II detoxifying enzymes GST and UGT2b1 activity as well as inhibiting the inflammatory responses by blocking NF- κ B activation and the release of pro-inflammatory mediators. These findings provided that a promising usefulness for MAEs in the prevention of the liver cancer.

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Author Contributions

JG Zheng and SF Liao conceived and designed the experiments. SF Liao and JH Liu performed the experiments, analyzed the data. SF Liao drafted the manuscript. JG Zheng and M Xu revised the manuscript.

Conflicts of Interest

There is no conflict of interest in this article. All the authors reviewed the paper and approved the final version.

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Appendix A. Supplementary Data

This document file contains Supplementary **Table A1**.

Evaluation of the liver cancer prevention of anthocyanin extracts from mulberry (*Morus alba* L.) variety PR-01.

Table A1. Structural identification of the mulberry anthocyanin monomer.

Peak	Compound	Code	[M-H] ⁺ (m/z)	Fragment (m/z)
1	pelargonidin 3-glucoside chloride	P3G	433.09	271
2	Cyanidin-3-O-glucoside chloride	C3G	449.08	287
3	Cyanidin 3-rutinoside chloride	C3R	595.05	287,449