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RESEARCH ARTICLE

## Metabolic profile and hepatoprotective activity of the anthocyanin-rich extract of *Hibiscus sabdariffa* calyces

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### ABSTRACT

**Context:** *Hibiscus sabdariffa* L. (Malvaceae) is a common traditional tea that has many biological activities. **Objectives:** To evaluate the hepatoprotective effect and study the metabolic profile of the anthocyanin-rich extract of *H. sabdariffa* calyces (HSARE).

**Materials and methods:** The hepatoprotective activity of HSARE was assessed (100 mg/kg/d for 4 weeks) by examining the hepatic, inflammatory, oxidative stress markers and performing a histopathological examination in rats with thioacetamide (TAA)-induced hepatotoxicity. HSARE was analyzed using ultra-performance liquid chromatography-quadrupole-time-of-flight-photodiode array-mass spectrometry (UPLC-qTOF-PDA-MS).

**Results:** The UPLC-qTOF-PDA-MS analysis of HSARE enabled the identification of 25 compounds represented by delphinidin and its derivatives, cyanidin, kaempferol, quercetin, myricetin aglycones and glycosides, together with hibiscus lactone, hibiscus acid and caffeoylquinic acids. Compared to the TAA-intoxicated group, HSARE significantly reduced the serum levels of alanine aminotransferase, aspartate aminotransferase and hepatic malondialdehyde by 37.96, 42.74 and 45.31%, respectively. It also decreased hepatic inflammatory markers, including tumour necrosis factor alpha, interleukin-6 and interferon gamma (INF- $\gamma$ ), by 85.39, 14.96 and 70.87%, respectively. Moreover, it decreased the immunopositivity of nuclear factor kappa-B and CYP2E1 in liver tissue, with an increase in the effector apoptotic marker (caspase-3 positive cells), restoration of the altered hepatic architecture and increases in the activities of superoxide dismutase (SOD) and glutathione by 150.08 and 89.23%, respectively.

**Discussion and conclusion:** HSARE revealed pronounced antioxidant and anti-inflammatory potential where SOD and INF- $\gamma$  were significantly improved. HSARE possesses the added value of being more water-soluble and of natural origin with fewer side effects expected compared to silymarin.

### ARTICLE HISTORY

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### Introduction

Liver disorders, despite being a worldwide problem, still suffer from inadequate medical management. Liver cell injury can be induced by various toxicants, such as certain chemotherapeutic agents, carbon tetrachloride (CCl<sub>4</sub>), thioacetamide (TAA), chronic alcohol consumption and microbes (Agarwal 2001). Hepatotoxins initially damage the centrilobular region of liver, where there are high levels of cytochrome P450 oxidases that convert potential hepatotoxins to toxic intermediates, releasing reactive oxygen species (ROS) and inducing lipid peroxidation and the production of proinflammatory cytokines (Luster et al. 2000). Tumour necrosis factor alpha (TNF- $\alpha$ ) and interleukin-6 (IL-6) are considered the major hepatotoxic mediators in several experimental models of liver injuries (Zhang et al. 2004).

Corticosteroids and antiviral drugs are considered the best conventional remedies for the treatment of liver disease despite their known adverse effects (Yang et al. 2008). Silymarin is one of the oldest plant products used as a hepatoprotective (Pradhan & Girish 2006). The drug has poor water solubility, leading to a low bioavailability. It also possesses a laxative effect and causes bloating, dyspepsia, nausea and irregular stools due to increased

bile secretion and flow (Jacobs et al. 2002). To our knowledge, a successful and safe therapy to prevent the progression of the hepatic disease is not yet available. *Hibiscus sabdariffa*, as a member of the family Malvaceae (Castañeda-Ovando et al. 2009), shows antioxidant, antigenotoxic and biomodulatory effects on various cells in animals exposed to toxic agents (Ramirez-Rodriguez et al. 2011). The plant also has antimicrobial, diuretic and antihypertensive effects (Alarcon-Aguilar et al. 2007). Water-soluble extracts of *H. sabdariffa*, containing several antioxidants, such as a protocatechuic acid (Liu et al. 2002) and anthocyanins (Wang et al. 2000), have the power to prevent peroxidative liver damage (Tatsuzawa et al. 1994). The anthocyanin-rich extract of *H. sabdariffa* calyces (HSARE), which is known to be water-soluble, was reported to be useful against diseases for which oxidative stress plays a role in their aetiopathogenesis, probably through its antioxidant and free radical-defusing effects (Ozkol et al. 2015).

The growing interest in phytomedicine and herbal supplements drives the development of analytical methods for the quality control of multicomponent mixtures, e.g., 'plant-derived nutraceuticals'. Concerns regarding quality control have stimulated a search for ultra-performance analytical methods (Farang et al. 2014). Ultra-performance liquid chromatography (UPLC) has

offered new possibilities for improving analytical methods of a complex sample matrix compared to conventional high-performance liquid chromatography (HPLC). The mass analyzer quadrupole-time-of-flight mass spectrometry (qTOF/MS) combines the high performance of TOF analysis in both MS and tandem MS (MS<sup>2</sup>) modes, providing high sensitivity and high mass accuracy for both precursor and product ions (Motilva et al. 2013).

In this study, the advanced analytical platform UPLC-qTOF/PDA/MS was used to achieve a better separation and characterization of the phenolic compounds of HSARE. Moreover, the hepatoprotective, anti-inflammatory and antioxidant effects of HSARE were compared to those of the conventional hepatoprotective drug 'silymarin' in rats with TAA-induced hepatotoxicity.

## Materials and methods

### Chemicals and reagents

HPLC grade acetonitrile, trifluoroacetic acid (TFA) and formic acid (J. T. Baker; Deventer, Netherlands) were used, and Milli-Q water was used for the for UPLC analysis (Millipore, Merck, Darmstadt, Germany). Diaion HP-20 AG was used for the column chromatography (75–150 µm, Mitsubishi Chemical Industries Co., Ltd., Tokyo, Japan). Delphinidin, cyanidin, chlorogenic acid and quercetin were obtained from Indofine (Somerville, NJ). TAA and trichloroacetic acid (TCA) were obtained from Sigma-Aldrich Chemical Co. (St. Louis, MO) Thiobarbituric acid (TBA) was obtained from BDH Chemicals Ltd. (Poole, Dorset, United Kingdom). Silymarin 'Legalon<sup>®</sup>' was obtained from Chemical Industries Development (CID, Giza, Egypt). Alanine aminotransferase (ALT) and aspartate aminotransferase (AST) kits were obtained from Spectrum, MDSS (Hannover, Germany). TNF- $\alpha$ , IL-6 and interferon gamma (INF- $\gamma$ ), enzyme-linked immunosorbent assay (ELISA) complete test were obtained from Komabiotech (Seoul, South Korea), and superoxide dismutase (SOD) and glutathione (GSH) reduced kits were obtained from Biodiagnostic (Giza, Egypt).

### Plant material

Calyces of *H. sabdariffa* were obtained from Aswan in April 2013 and authenticated by Dr. Reem Samir Hamdy, Lecturer of Plant Taxonomy, Botany Department, Faculty of Science, Cairo University, Giza, Egypt. A voucher specimen (HS-2013-65) was submitted to the herbarium of the Department of Pharmacognosy, Faculty of Pharmacy, Cairo University.

### Preparation of HSARE

HSARE was prepared according to Hong and Wrolstad (1990). *H. sabdariffa* calyces (1 kg) were pulverized and extracted with 10 l of 0.1% TFA aqueous solution for 12 h at 20 °C, and the extract was filtered and applied to a Diaion HP column. The resin was washed with 3 l of water and then eluted with 50% ethanol solution containing 0.1% TFA. The eluate was dried under vacuum at 20 °C and then freeze-dried (LyoQuest-55 freeze dryer; Telstar Life Science Company, Spain).

### Sample preparation for UPLC-PDA-MS analysis

HSARE (5 mg) was homogenized with 6 ml 100% methanol (MeOH) containing 5 µg/ml umbelliferone (internal standard) using a Turrax mixer (11,000 rpm) for five 20-s periods with 1 min intervals separating each period to prevent heating. The

extract was then vortexed vigorously and centrifuged at 3000 rpm for 30 min to remove debris. An aliquot of 500 µl was placed on a (500 mg) C<sub>18</sub> cartridge preconditioned with MeOH and water. The sample was eluted using 3 ml 100% MeOH, the eluent was evaporated under a nitrogen stream, and the obtained dry residue was resuspended in 300 µl MeOH. Three microlitres of the supernatant was used for UPLC-MS analysis.

### UPLC-ESI-low resolution full MS and MS<sup>n</sup> analysis

The UPLC coupled to electrospray ionization mass spectrometry (ESI) and ESI-MS<sup>n</sup> mass spectra were obtained from an LCQ Deca XP MAX ion trap system (Thermo Scientific, San Jose) equipped with an ESI source (electrospray voltage 4.0 kV, sheath gas: nitrogen; capillary temperature: 275 °C) in both positive and negative ionization modes. The Ion Trap MS system is coupled with an ACQUITY UPLC system (Waters) equipped with an HSS T3 column (100 × 1.0 mm, particle size 1.8 µm; Waters), applying the following binary gradient at a flow rate of 150 µl min<sup>-1</sup>: 0–1 min, isocratic 95% A (water/acetic acid, 99.8/0.2 [v/v]), 5% B (acetonitrile/acetic acid, 99.8/0.2 [v/v]); 1–16 min, linear from 5 to 95% B; 16–18 min, isocratic 95% B; 18–20 min, isocratic 5% B. The injection volume was 1.5 µl (partial loop with needle overfill injection). Eluted compounds were detected from *m/z* 100 to 1000. The MS<sup>n</sup> spectra were recorded using the following conditions: MS/MS<sup>n</sup> analysis with starting normalized collision energy of 45% and isolation width of ±2 D in data-dependent negative and positive ionization modes.

### High-resolution UPLC-MS analysis

A micrOTOF-Q hybrid qTOF-MS (Bruker Daltonics) was used for analyzing the metabolites after UPLC separation, using the Waters UHPLC setup described for the full-scan MS analysis with the same elution gradient and equipped with an Apollo II electrospray ion source in positive and negative ion modes. The instrument settings were as follows: nebulizer gas, nitrogen, 1.6 bar; dry gas, nitrogen, 6 l min<sup>-1</sup>, 190 °C; capillary, -5500 V (+4000 V); end plate offset, -500 V; funnel 1 RF, 200 Vpp; funnel 2 RF, 200 Vpp; in-source CID energy, 0 V; hexapole RF, 100 Vpp; quadrupole ion energy, 5 eV; collision gas, argon; collision energy, 10 eV; collision RF 200/400 Vpp (timing 50/50); transfer time, 70 µs; pre-pulse storage, 5 µs; pulser frequency, 10 kHz; spectra rate, 3 Hz. The internal mass calibration of each analysis was performed by an infusion of 20 µl of 10 mm lithium formate in isopropanol/water, 1/1 (v/v), at a gradient time of 18 min using a diverter valve. UPLC-MS files were converted to netCDF file format using the file converter tool in Xcalibur software and further processed using AMDIS software to assist in adjacent peak deconvolution and background subtraction (Halket et al. 1999).

## Biological study

### Animals

Male Sprague-Dawley rats were provided by the Schistosome Biological Supply Center, Theodor Bilharz Research Institute (TBRI), Giza, Egypt. The rats weighed 120–160 g at the beginning of the experiment and were maintained on a standard commercial pelleted diet (El-Kahira company for oils and soap, Egypt) in an air-conditioned animal house at 20–22 °C under 12 h light and dark cycles. All experiments followed international guidelines for animal ethics and were approved by the Institutional Review Board of TBRI.

### Induction of hepatic toxicity

A pilot study was conducted using 24 rats to evaluate the toxicity of TAA. Rats were randomly divided into three groups ( $n=8$ ), and each was injected with TAA intraperitoneally (IP) in doses of 100, 200 or 300 mg/kg once/week for 1 month to observe the mortality rate. At the end of the experiment, the animals were sacrificed, and the livers were extracted and maintained in 10% formalin for consequent histopathological examination to select the proper dose for the induction of hepatotoxicity.

### Animal groups

A total of 32 rats were divided equally into four groups of eight rats each. Group I (normal control) received the drug vehicle Cremophore-El. Groups II–IV were subjected to the induction of hepatotoxicity using an IP injection of TAA at a dose of 100 mg/kg once/week for four successive weeks. Group II was left without treatment as TAA-intoxicated rats, while groups III and IV concurrently received either silymarin or HSARE in a daily oral dose of 50 or 100 mg/kg for 5 d/week for four successive weeks, respectively. Twenty-four hours after the last dose, the animals were sacrificed, blood samples were collected and the livers were extracted and divided into two parts for histopathological examination and oxidative stress marker determination.

### Biochemical assays

Sera were used to examine biochemical markers expressing hepatic damage. AST, ALT (Reitman & Frankel 1957) and the pro-inflammatory cytokines TNF- $\alpha$ , IL-6 and INF- $\gamma$  were assayed using commercially available ELISA kits.

### Oxidative stress marker determination

One gram of the liver was homogenized in five volumes (w/v) of ice-cold 0.1 M potassium phosphate buffer (pH =7.4). The homogenate was centrifuged at 1000 rpm for 10 min to remove cell debris, the supernatant was centrifuged at 20,000 rpm and the pellet was discarded. The supernatant was collected and stored at  $-70^{\circ}\text{C}$  for the estimation of the liver GSH content, SOD activity and lipid peroxidation, expressed as malondialdehyde (MDA) formation.

Liver homogenates were deproteinized in 5% (w/v) TCA and centrifuged at 3000 rpm for 20 min, and the GSH content and SOD activity were estimated (Ellman 1959; Winterbourn et al. 1975). The MDA was assayed according to Ohkawa et al. (1979), and the absorbance was read, after cooling, at 530 nm against a blank containing all the reagents except the liver homogenate. The content of thiobarbituric acid reactive substances in the samples was calculated using the extinction coefficient of MDA ( $1.56 \times 10^5 \text{ M}^{-1} \text{ cm}^{-1}$ ), and the results were expressed as MDA equivalents in nmol/g liver.

### Histopathological and immunohistochemical examinations

The livers were fixed in 10% formalin, embedded in paraffin wax, sliced into 5  $\mu\text{m}$  thick sections on positively charged glass slides and stained with haematoxylin-eosin for examination under a Zeiss microscope with 200 $\times$  magnification power. For the immunohistochemical examination, three paraffin sections were deparaffinized and rehydrated. Endogenous peroxidase was inactivated by incubation in 0.3% hydrogen peroxide in absolute

MeOH for 30 min. The sections were incubated in 5% skim milk for 30 min at room temperature. Antigen retrieval was performed by microwave (700 W) treatment in 10 mm citrate buffer (pH =7.4) for 15 min. The sections were then incubated overnight at  $4^{\circ}\text{C}$  with anti-rat caspase 3, CYP2E1 and nuclear factor kappa-B (NF- $\kappa\text{B}$ ) primary antibodies (Abcam, Cambridge) at dilutions of 1:100 1:150 and 1:150, respectively. After washing with phosphate buffer saline (PBS), the sections were incubated at room temperature for 30 min in secondary antibodies (Abcam). A brown colour developed after the addition of 3-diaminobenzidine (DAB) for 2–4 min, followed by washing in distilled water and counterstaining with Mayer's haematoxylin for 1 min at room temperature. The per cent of positively stained brown nuclei (NF- $\kappa\text{B}$ ) or brown cytoplasm (caspase-3 and CYP2E1) in 10 successive fields at a magnification of 200 $\times$  were calculated.

### Statistical analysis

The data are expressed as the mean  $\pm$  SEM. Statistical analysis of the data was performed using the one-way ANOVA test followed by Tukey's *post hoc* test to determine the differences among the mean values of different groups. The data were analyzed using the SPSS software package, version 16.0 (Chicago, IL). *p* Values  $<0.05$  were considered statistically significant.

## Results

### Identification and quantification of metabolites via UPLC-MS

Metabolites were characterized by their UV-VIS spectra (220–600 nm), retention times relative to external standards, mass spectra and comparison to a phytochemical dictionary of natural products database (Wiley, CRC) and other reference literature. The analytical method applied enabled the characterization of 25 compounds. The compounds identified using UPLC-MS analysis is listed in Table 1 and Figures 1 and 2.

### Induction of hepatic toxicity

No mortality was observed with all tested doses of TAA; nevertheless, the dose of 100 mg/kg once per week for 1 month caused maximal severe focal hepatic necrosis and infiltration of chronic inflammatory cells when compared to the doses of 200 and 300 mg/kg (Figure 3).

### Hepatoprotective activity study

The treatment of the TAA-intoxicated rats with HSARE at a dose of 100 mg/kg produced a significant hepatoprotective effect (Table 2). This was expressed as a significant reduction in the activities of ALT and AST (37.96 and 42.74%, respectively) compared to the TAA group. This effect was comparable to that of silymarin at a dose of 50 mg/kg.

### Anti-inflammatory activity study

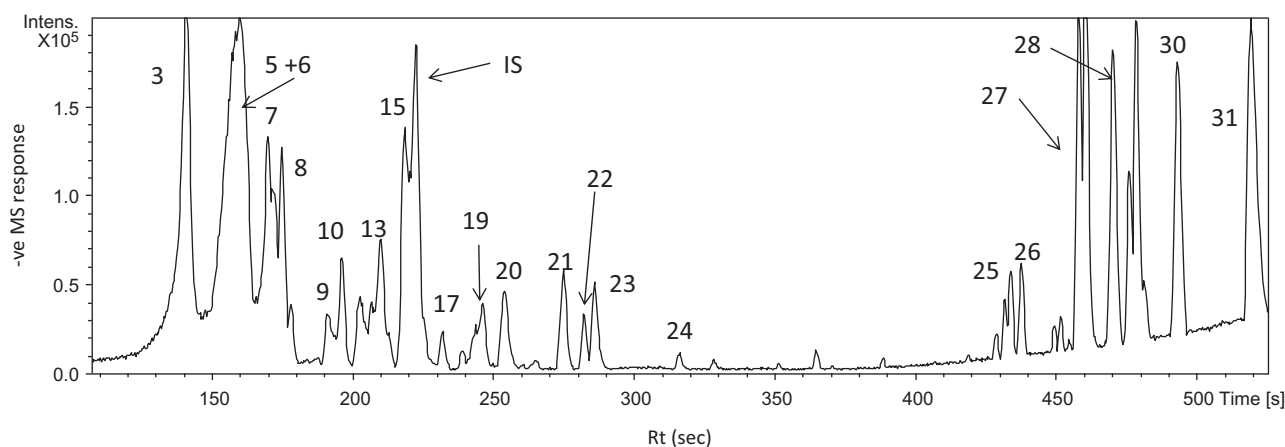
The level of TNF- $\alpha$  was significantly decreased in the groups treated with either HSARE or silymarin (Table 2) compared to the TAA-intoxicated rats. The level of IL-6 tended to be normal, while INF- $\gamma$  was normalized in the HSARE-treated group. It is worthy to note that the effect of HSARE on TNF- $\alpha$  and INF- $\gamma$  was more pronounced than that of silymarin.



**Table 1.** Peak assignments of metabolites in HSARE using UPLC-PDA-MS in negative ionization mode.

Peak (ppm)	$t_R$ (s)	$\lambda$ max (nm)	$[M-H]^-$	Molecular formula	Error	Identification	$MS^2$
1	22	Nd	189.0057	$C_6H_5O_7$	-0.8	Hibiscus lactone (Hydroxy citric acid lactone)	129
2	25	Nd	207.015	$C_6H_7O_8$	1.8	Hibiscus acid (Hydroxy citric acid)	189, 129
3	141	290 shd, 323	353.0939	$C_{16}H_{17}O_9$	16.3	O-Caffeoyl quinic acid	191, 179
4	148	277, 330, 500	681.1285	$C_{29}H_{29}O_{19}$	3.5	Delphinidin hexosyl pentosyl malonate	595, 301
5	150	Nd	613.1416	$C_{26}H_{29}O_{17}$	0.9	Unknown delphinidin derivative	467, 301
6	152	280,334, 440 shd, 500	595.1311	$C_{26}H_{27}O_{16}$	-1.1	Delphinidin 3-sambubioside (Hibiscin)	301
7	167	275, 330, 441 shd, 500	579.1354	$C_{26}H_{27}O_{15}$	0.2	Cyanidin 3-sambubioside	285 -ve
8	170	282, 325	353.0941	$C_{16}H_{17}O_9$	17.9	O-Caffeoyl quinic acid	191.0594, 36.0606
9	189	Nd	611.1255	$C_{26}H_{27}O_{17}$	-0.2	Myricetin 3-arabinogalactose	317, 315
10	196	Nd	611.1355	$C_{26}H_{27}O_{17}$	16.4	Myricetin pentosylhexoside isomer	317
11	199	276, 333, 444 shd, 500	625.1394	$C_{27}H_{29}O_{17}$	2.5	Delphinidin 3-gentiobioside	301
12	200	Nd	479.0837	$C_{21}H_{19}O_{13}$	-1.3	Gossypetin 3-glucoside (Gossytrin)	317
13	204	277, 345	595.1411	$C_{26}H_{27}O_{16}$	-0.1	Quercetin 3-sambubioside	463, 481, 301 neg
14	210	Nd	335.0835	$C_{16}H_{15}O_8$	2.5	O-Caffeoylshikimic acid	161
15	212	Nd	609.1465	$C_{27}H_{29}O_{16}$	-0.7	Delphinidin 3-neohesperidoside	301
16	217	276, 500	463.0884	$C_{21}H_{19}O_{12}$	-0.4	Delphinidin 3-galactoside	301
17	232	263, 348	593.1613	$C_{27}H_{29}O_{15}$	0	Kaempferol 3-O-rutinoside	285
18	239	263, 348	447	$C_{21}H_{19}O_{11}$	1.3	Kaempferol hexoside	285
19	246	Nd	533.1391	$C_{25}H_{25}O_{13}$	0.6	Dimethyl-delphinidin-glucosyl acetate	371, 329
20	253	Nd	317.0358	$C_{15}H_9O_8$	17.5	Myricetin	248.9644, 189.0064
21	275	Nd	312.1293	$C_{18}H_{18}NO_4$	-0.4	N-Feruloyltyramine	179
22	276	Nd	301.0357	$C_{15}H_9O_7$	-1	Delphinidin	179, 151
23	293	Nd	301.0405	$C_{15}H_9O_7$	17	Quercetin	-
24	311	Nd	285.0763	$C_{15}H_9O_6$	0	Cyanidin	-
30	494	Nd	279.2378	$C_{18}H_{31}O_2$	17.4	Linoleic acid	-
31	520	Nd	255.2369	$C_{16}H_{31}O_2$	15.6	Palmitic acid	112.988

Nd, not detected; shd, shoulder.



**Figure 1.** A representative UPLC-negative ionization MS trace of methanol extract of *H. subdariffa* flowers, peak numbers follow those listed in Table 1 for metabolite identification using UPLC-PDA-MS. IS, internal standard (umbelliferone);  $R_t$ , retention time.

### Oxidative stress and lipid peroxidation studies

The administration of silymarin (50 mg/kg) and HSARE (100 mg/kg) produced a marked increase in GSH content (130.36 and 89.28%, respectively), improved the activity of SOD (99.41 and 150.08%, respectively) and reduced the MDA level (32.07 and 45.31%, respectively) relative to the TAA-intoxicated group (Table 2). Moreover, HSARE was more effective in improving the MDA level and SOD activity when compared with silymarin (Table 2).

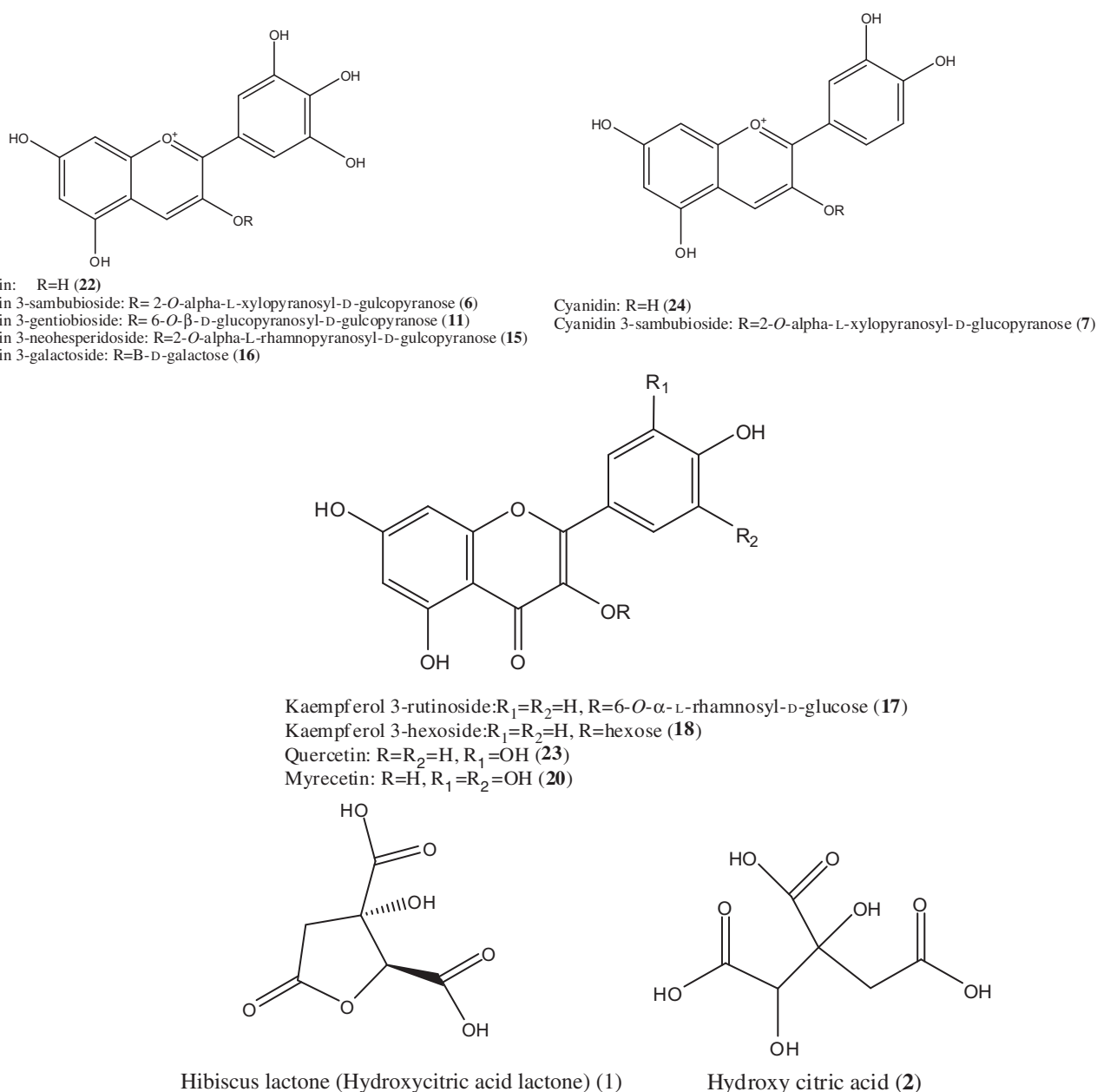
### Histopathological and immunohistochemical examinations

When compared to the normal rats (Figure 4(a)), TAA-intoxicated rats (Figure 4(b)) showed severe disturbances of the normal architecture of the hepatic tissue and portal tract inflammation with severe lymphocyte infiltration and focal spotty necrosis. On the other hand, the HSARE and silymarin-treated rats (Figure 4

(d) and (c), respectively) showed almost normal hepatic architectural patterns with only minor chronic inflammatory cell infiltration. The immunohistochemical examination of caspase-3 showed no significant difference in the TAA-intoxicated group compared to normal, while in the silymarin and HSARE-treated groups, the number of positive cells increased by 4.4- and 6.4-fold, respectively, when compared to the normal control group. In TAA-intoxicated rats, the percentages of positive cells of NF- $\kappa$ B and CYP2E1 were significantly elevated. Meanwhile, treatment with either silymarin or HSARE resulted in significant reductions in both NF- $\kappa$ B and CYP2E1 (4.3- and 3.9- and 3.4- and 4.5-fold, respectively) compared to the TAA-intoxicated group (Figures 5 and 6).

### Discussion

To our knowledge, this current study is the first report of the metabolic profiling of HSARE using UPLC-qTOF-PDA-MS.

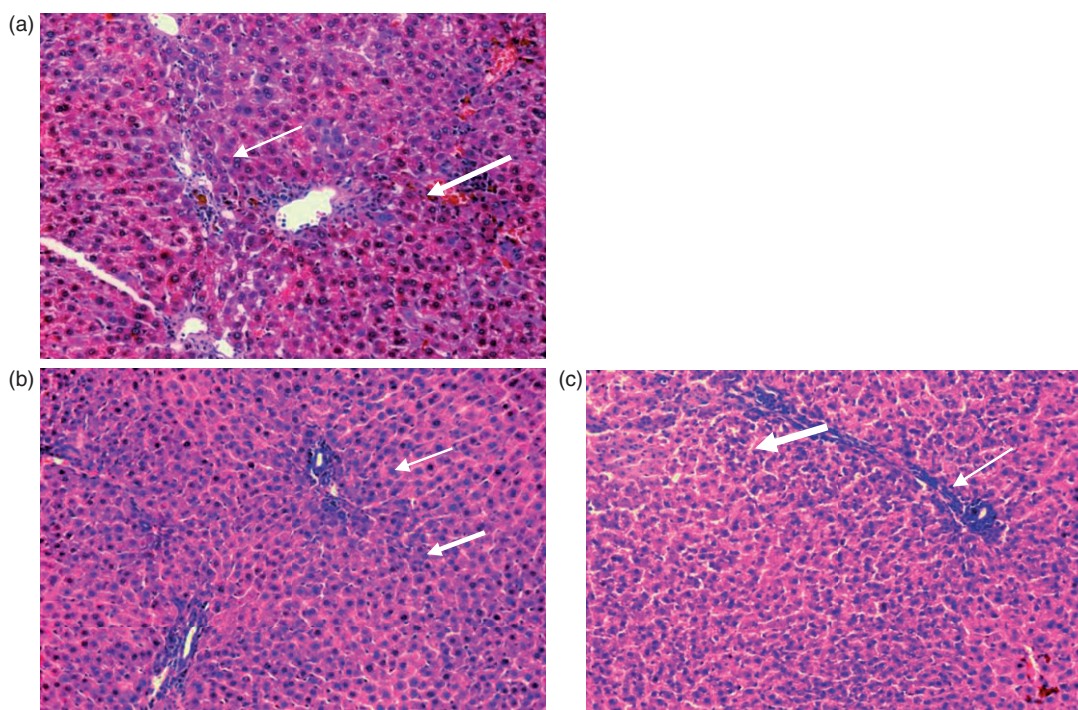


**Figure 2.** Structures of some identified compounds detected via UPLC-qTOF-PDA-MS analysis of *H. subdarifa* anthocyanin-rich extract (HSARE).

The analytical method applied enabled the characterization of 25 compounds. For determining the structures of these compounds, the UV-VIS spectra and HPLC retention characteristics were first considered. The compounds associated with peaks 4, 6, 11, 15 and 22 showed  $\lambda$  maxes at 277, 330 and 500 nm, which suggests that they might be delphinidin derivatives (Giusti et al. 1999). Peak 6 showed a molecular ion at  $m/z$  595.1311 that produced a unique MS<sup>2</sup> fragment at  $m/z$  301 (delphinidin). The direct loss of 294 amu [ $-(162 + 132)$ ] indicates the separation of a disaccharide (hexose + pentose) residue (Giusti et al. 1999). This suggestion is also supported by the existence of a shoulder approximately 440 nm in the UV-VIS spectrum of this peak, characterized by 3-glycosylated anthocyanidins (Santos-Buelga et al. 2003). The sugar moiety was identified as sambubiose (2-O- $\beta$ -D-xylosyl-D-glucose), which is most frequently found as an anthocyanin substituent (Mazza & Miniati 1993). Peak 6, identified as delphinidin-3-sambubioside (hibiscin) (C<sub>26</sub>H<sub>27</sub>O<sub>16</sub>), was previously identified in the calyx of *H. subdariffa*

(Williamson et al. 2013). Mass data of Peak 4 showed a molecular ion peak at  $m/z$  681.1285 and MS<sup>2</sup> at  $m/z$  595 (M-86) and 301 [(M-86)-294]. These were indicative of a delphinidin with a malonic acid moiety (86 amu) and hexose and pentose moieties (294 amu). The UV-VIS spectrum of this peak was characteristic of 3-substituted anthocyanidin. The loss of the malonic acid followed by the two sugars together may indicate that the acyl and sugar residues are attached to the same position. This compound was tentatively identified as delphinidin hexosyl pentosyl malonate (C<sub>29</sub>H<sub>29</sub>O<sub>19</sub>).

Peak 11 showed a molecular ion at  $m/z$  625.1394 and a MS<sup>2</sup> at  $m/z$  301 with a loss of 324 amu (162 + 162), corresponding to two hexose molecules; this compound was regarded as delphinidin-3-gentiobioside (C<sub>27</sub>H<sub>29</sub>O<sub>17</sub>). Delphinidin-3-neohesperidoside was assigned to peak 15 based on the MS, showing a molecular ion at  $m/z$  609.1465 for the molecular formula C<sub>27</sub>H<sub>29</sub>O<sub>16</sub> and MS<sup>2</sup> at  $m/z$  301 [M-(162 of glucose + 146 of rhamnose)]. These data were in agreement with the fragmentation pattern of the



**Figure 3.** (a) Rats liver section injected with a single intraperitoneal dose (100 mg/kg) of TAA once per week for 1 month showing portal tract infiltration with chronic inflammatory cells (thin arrow), focal spotty necrosis (inflammatory cells in between hepatocytes, thick arrow), (b and c) rats injected with a single intraperitoneal dose (200 mg/kg or 300 mg/kg, respectively) of TAA once per week for 1 month showed portal tracts area infiltrated by moderate number of chronic inflammatory cells (thin arrow), focal spotty necrosis (thick arrow), inflammatory cells in between hepatocytes (H&E  $\times$  200).

**Table 2.** Effect of HSARE on hepatic marker enzymes, anti-inflammatory markers, oxidative stress and lipid peroxidation after 4 weeks of thioacetamide (TAA) intoxication in rats.

	Normal	TAA	Silymarin (50 mg/kg/d)	HSARE (100 mg/kg/d)
ALT (U/l)	61.89 $\pm$ 4.38	138.42 $\pm$ 8.14 <sup>a</sup>	80.49 $\pm$ 3.08 <sup>b</sup> (-41.85)	85.87 $\pm$ 8.37 <sup>b</sup> (-37.96)
AST (U/l)	115.01 $\pm$ 2.67	236.46 $\pm$ 16.35 <sup>a</sup>	125.24 $\pm$ 5.52 <sup>b</sup> (-47.03)	135.38 $\pm$ 3.21 <sup>b</sup> (-42.74)
TNF- $\alpha$ (pg/ml)	245.17 $\pm$ 24.13	1775.09 $\pm$ 136.24 <sup>a</sup>	279.25 $\pm$ 15.59 <sup>b</sup> (-84.27)	259.38 $\pm$ 17.54 <sup>b</sup> (-85.39)
IL-6 (pg/ml)	278.33 $\pm$ 4.59	380.00 $\pm$ 5.35 <sup>a</sup>	267.50 $\pm$ 10.00 <sup>b</sup> (-29.6)	323.13 $\pm$ 3.89 <sup>a,b,c</sup> (-14.96)
INF- $\gamma$ (pg/ml)	23.09 $\pm$ 1.38	82.60 $\pm$ 3.82 <sup>a</sup>	38.52 $\pm$ 3.28 <sup>a,b</sup> (-53.36)	24.06 $\pm$ 1.05 <sup>b,c</sup> (-70.87)
GSH (mmol/g liver)	2.66 $\pm$ 0.23	1.12 $\pm$ 0.15 <sup>a</sup>	2.58 $\pm$ 0.14 <sup>b</sup> (+130.36)	2.12 $\pm$ 0.04 <sup>b</sup> (+89.28)
SOD (U/g liver)	385.42 $\pm$ 21.63	124.25 $\pm$ 4.27 <sup>a</sup>	247.77 $\pm$ 17.33 <sup>a,b</sup> (+99.41)	310.73 $\pm$ 19.81 <sup>b</sup> (+150.08)
MDA (nmol/g liver)	6.36 $\pm$ 0.34	12.69 $\pm$ 0.15 <sup>a</sup>	8.62 $\pm$ 0.34 <sup>a,b</sup> (-32.07)	6.94 $\pm$ 0.21 <sup>b,c</sup> (-45.31)

Values presented as mean of 6–8 rats  $\pm$  SEM.

Numbers between parentheses represent percentage change compared to the TAA-intoxicated group.

<sup>a</sup>Significantly different from normal control at  $p < 0.05$ .

<sup>b</sup>Significantly different from TAA at  $p < 0.05$ .

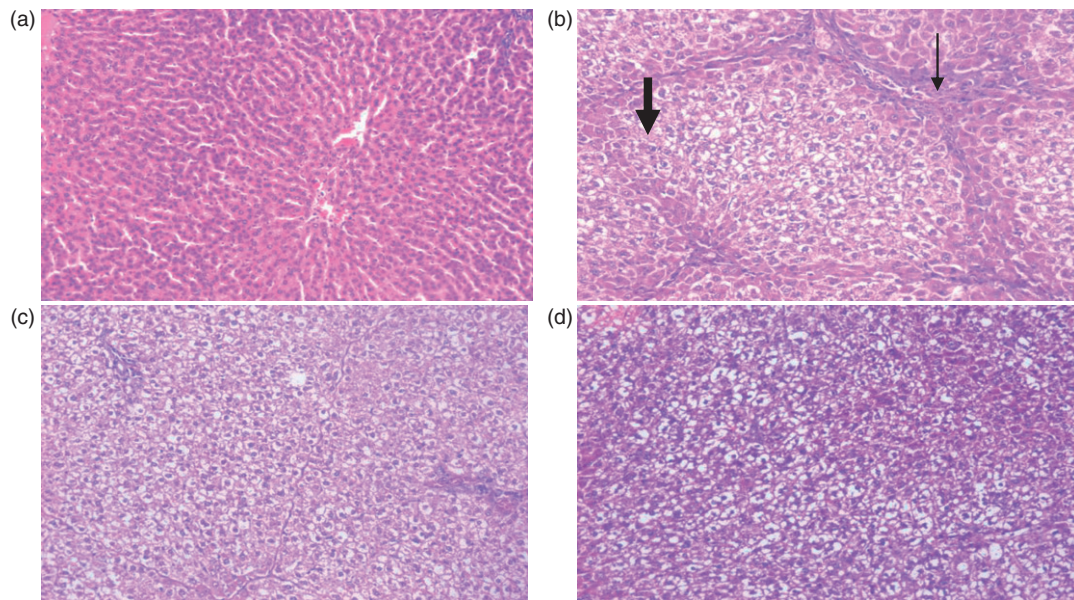
<sup>c</sup>Significantly different from silymarin at  $p < 0.05$ .

compound. Peak **16** with  $m/z$  463.0884 was characterized as delphinidin-3-galactoside ( $C_{21}H_{19}O_{12}$ ), as it showed a base peak at  $m/z$  301 [M-162], which corresponds to the loss of a hexose molecule. Peak **19** was identified as dimethyl-delphinidin-glucosyl acetate ( $C_{25}H_{25}O_{13}$ ), with a base peak  $m/z$  595.1311 corresponding to [M-H]<sup>-</sup> and MS<sup>2</sup> at  $m/z$  371 (M-162) and 329 [(M-162)-CH<sub>3</sub>CO], assigned to the sequential loss of glucose then an acetyl group. Peak **22**, with a base peak [M-H]<sup>-</sup> at  $m/z$  301.0357, corresponds to the molecular formula  $C_{15}H_9O_7$  and was identified as delphinidin (aglycone); MS<sup>2</sup> at [(M-H)-122] and [(M-H)-122]-28] were assigned for retro Diels–Alder fission and the loss of CO, respectively. Delphinidin was previously identified in the flowers of *H. sabdariffa* (Mahadevan et al. 2009). In addition, an unknown delphinidin derivative at peak **5** with a molecular ion at  $m/z$  613.1416 corresponds to the formula  $C_{26}H_{29}O_{17}$ . Another group of anthocyanins were cyanidin and its derivatives, which were detected as peak **7**, exhibited a fragmentation pattern, absorption spectrum and sugar identification similar to those of peak **6**. This compound showed a molecular ion at  $m/z$  579.1354

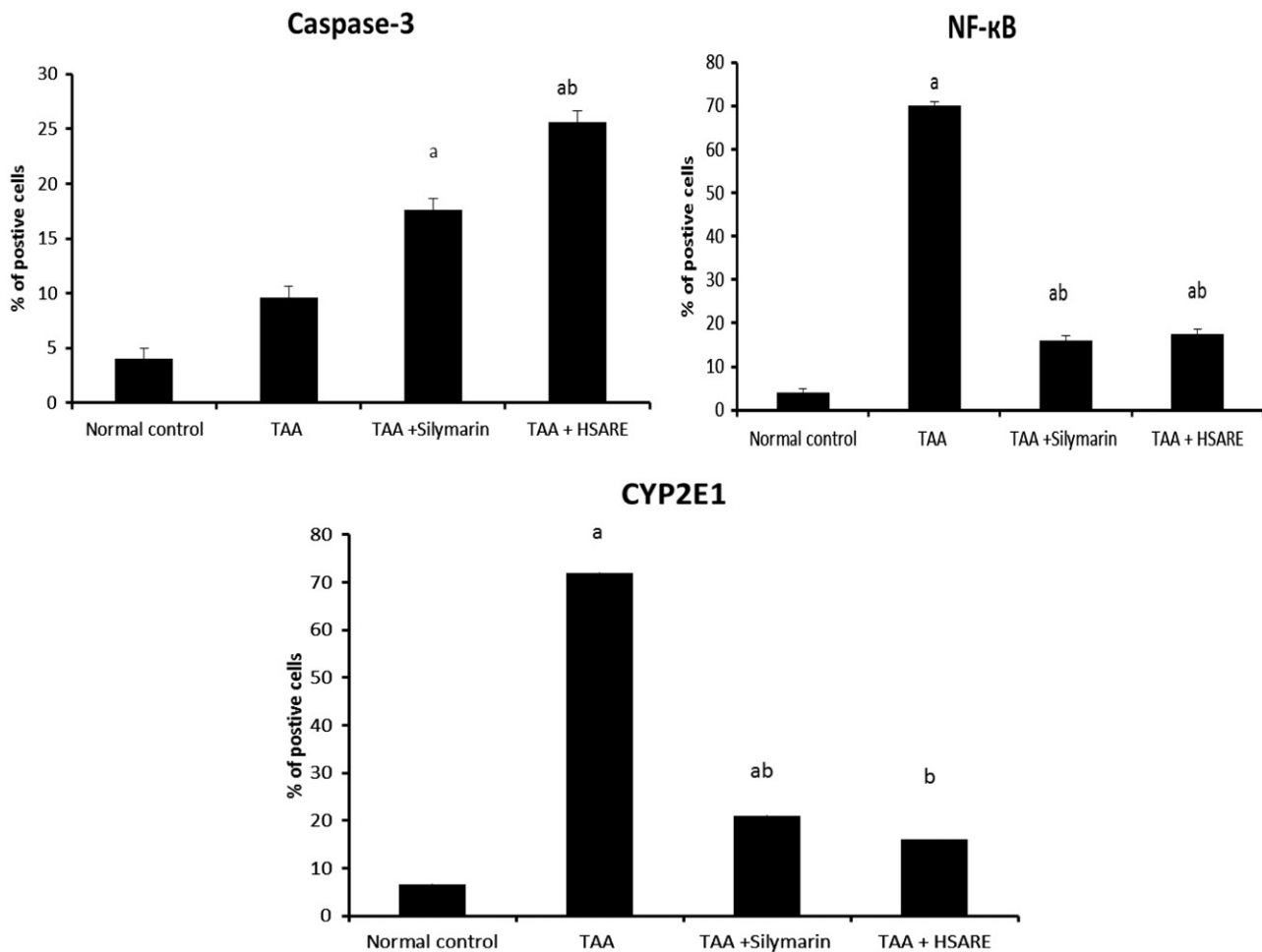
that produced an MS<sup>2</sup> fragmentation pattern, permitted a fragment ion to be distinguished at 285, corresponding to cyanidin due to the loss of 294 amu, which corresponded to glucose–xylose loss. Thus, this compound was characterized as cyanidin-3-sambubioside. This compound was previously reported in the calyx of *H. sabdariffa* (Herranz-Lopez et al. 2012). Peak **24** shows a molecular ion at  $m/z$  285.0763 [M-H]<sup>-</sup>, corresponding to the molecular formula  $C_{15}H_9O_6$  and assigned to cyanidin (aglycone).

Flavonoids were also detected and were represented by seven compounds. Peaks **9** and **10** exhibited the same molecular ion at  $m/z$  611.1255 in MS spectra, characterized as myricetin-3-arabinogalactose ( $C_{26}H_{27}O_{17}$ ). The examination of their respective MS<sup>2</sup> fragmentation patterns allowed one main fragment ion to be distinguished at 317, which corresponds to the aglycone myricetin and the loss of an arabinogalactose moiety. Thus, these compounds were characterized as two isomers and were previously reported in *H. sabdariffa* (Herranz-Lopez et al. 2012). Peak **12** was identified as gossypetin-3-glucoside (gossytrin) ( $C_{21}H_{19}O_{13}$ ) with a molecular ion peak at  $m/z$  479.0837 based on



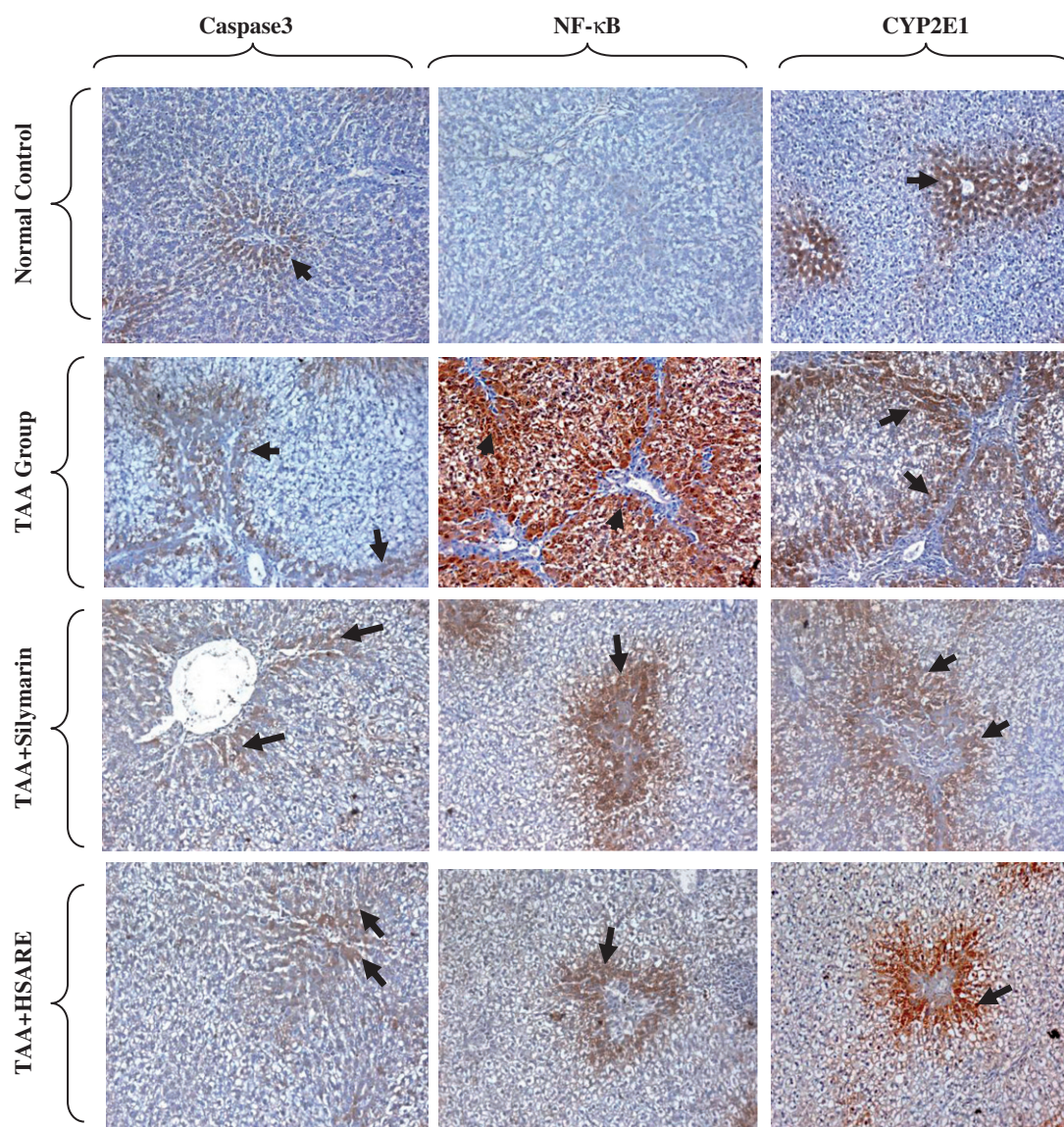


**Figure 4.** Histopathological examination of liver sections from normal untreated rats (a) showing normal hepatic architecture, TAA-intoxicated (100 mg/kg) rats (b) showing severely disturbed hepatic architecture and portal tract inflammation with severe lymphocyte infiltration (thick arrow) and focal spotty necrosis (thin arrow), silymarin-treated (50 mg/kg) rats (c) showing almost normal hepatic architectural pattern with mild ballooning of hepatocytes, mild dilation of sinusoids and lymphocyte infiltration and HSARE-treated (100 mg/kg) rats (d) showing normal hepatic architecture with mild ballooning of hepatocytes and mild number of chronic inflammatory cells infiltration (H&E  $\times$  200).



**Figure 5.** Bar graphs represent liver ( $n=6$ ) per cent of hepatocytes with positively stained brown cytoplasm (caspase-3 and CYP2E1) or brown nuclei (NF-κB), in 10 successive fields. <sup>a</sup>Significantly different from normal control at  $p < 0.05$ . <sup>b</sup>Significantly different from TAA at  $p < 0.05$ .





**Figure 6.** Immunohistochemical staining of rat liver sections for caspase-3, NF- $\kappa$ B and CYP2E1, showing normal control rats with few number of positive hepatocytes for caspase-3, NF- $\kappa$ B and CYP2E1 appearing as brownish stain (black arrows), TAA-intoxicated (100 mg/kg) rats showing few positive hepatocytes (centrilobular and periportal areas) with mild density for caspase-3, and many positive hepatocytes (centrilobular and periportal areas) with severe density for NF- $\kappa$ B and CYP2E1 (black arrows), and silymarin and HSARE-treated rats (50 and 100 mg/kg, respectively) showing moderate number of positive hepatocytes (periportal areas) with moderate density (periportal areas) for caspase-3 NF- $\kappa$ B and CYP2E1 (black arrows), (IHC, DAB,  $\times$  200).

its MS<sup>2</sup> fragmentation pattern and the published data. Gossypetin-3-glucoside (gossytrin) was previously identified in the calyx of *H. sabdariffa* (Williamson et al. 2013). Peak 13 showed [M-H]<sup>-</sup> at  $m/z$  595.1411 with a formula C<sub>26</sub>H<sub>27</sub>O<sub>16</sub> characterized as quercetin 3-sambubioside. The MS<sup>2</sup> of this ion revealed fragments at  $m/z$  463, indicating the loss of xylose (132 amu) and 301, indicating the loss of glucose (162 amu), which was also confirmed by its UV spectral data. Quercetin had already been identified in *H. sabdariffa* (Herranz-Lopez et al. 2012). Peak 17 was identified as kaempferol-3-O-rutinoside (C<sub>27</sub>H<sub>29</sub>O<sub>15</sub>), with a molecular ion peak at  $m/z$  593.1613. Its MS<sup>2</sup> pattern gave rise to a fragment ion at  $m/z$  285 assigned to the sequential loss of rhamnose and glucose. Peak 18 showed a molecular ion of  $m/z$  447 (C<sub>21</sub>H<sub>19</sub>O<sub>11</sub>), which yielded a daughter fragment of  $m/z$  285, assigned to kaempferol aglycone with a loss of 162 amu (a hexose moiety), and was tentatively identified as kaempferol-hexoside. Finally, the non-polar peaks 20 and 23 showed molecular ions at  $m/z$  317.0358 and 301.0405, referred to myricetin and quercetin, respectively.

The phenolic acids mainly included caffeic acid derivatives and were detected as two peaks 3 and 8 characterized by an ion at  $m/z$  353.0939 with a formula C<sub>16</sub>H<sub>17</sub>O<sub>9</sub> and identified as *O*-caffeoylquinic acid (Borrás-Linares et al. 2015). The MS<sup>2</sup> of the two compounds were 191.0594, corresponding to the loss of a caffeoyl moiety (162 amu) and the presence of a quinic acid moiety (191 amu). This is in accordance with the fragmentation pattern of *O*-caffeoyl quinic acid. Peak 14 corresponds to *O*-caffeoyl shikimic acid, identified by an ion at  $m/z$  335.0835 with a formula C<sub>16</sub>H<sub>15</sub>O<sub>8</sub>. The MS<sup>2</sup> of the compound was at 161 amu, corresponding to the loss of shikimic acid (174 amu), and it was previously reported in *H. sabdariffa* (Herranz-Lopez et al. 2012). Organic acids, such as citric acid derivatives, were represented by peak 1, corresponding to hibiscus lactone (hydroxy citric acid lactone) with an ion peak at  $m/z$  189.0057 and a formula (C<sub>6</sub>H<sub>5</sub>O<sub>7</sub>). Peak 2 with an ion at  $m/z$  207.015 with a formula C<sub>6</sub>H<sub>7</sub>O<sub>8</sub> corresponds to the [M-H]<sup>-</sup> characterized for hibiscus acid (hydroxy citric acid). The MS<sup>2</sup> of the compound revealed two peaks at 189 and 129. These two acids were previously

reported in the calyces of *H. sabdariffa* (Herranz-Lopez et al. 2012; Borrás-Linares et al. 2015).

A tyramine derivative was also detected and represented as peak 21 with  $[M-H]^-$  at  $m/z$  312.1293, corresponding to the formula ( $C_{18}H_{18}NO_4$ ) characterized for *N*-feruloyltyramine. The  $MS^2$  of the compound exhibited an ion peak at 179 and was previously isolated from the stem of *H. tiliaceus* (Chen et al. 2006a). Peaks 32 and 33, corresponding to fatty acids, were identified as linoleic and palmitic acids, with ions at  $m/z$  279.2378 and 255.2369, respectively.

In this study, the hepatoprotective activity of HSARE was examined in rats with TAA-induced hepatotoxicity. The TAA hepatotoxicity was more evident for the dose of 100 mg/kg once/week for 4 weeks than with higher doses of 200 or 300 mg/kg. The lesser hepatotoxicity with the higher doses may be related to the possible competition between TAA in high doses and its intermediate metabolite, thioacetamide *S,S*-dioxide (TASO), hindering its progression to the second reactive metabolite,  $TASO_2$ , which is capable of reacting with proteins, leading to the impairment of liver function (Hajovsky et al. 2012).

The dose of HSARE (100 mg/kg) selected was based on the earlier studies of Wang et al. (2000) and Adetutu and Owoade (2013), who demonstrated its effective antioxidant and hepatoprotective activities. In this work, HSARE demonstrated a significant hepatoprotective effect that was expressed as reductions in ALT and AST. Its effect was comparable to that of silymarin at a dose of 50 mg/kg. Significant reductions in the levels of these enzymes were recorded upon treatment with HSARE in mice with  $CCl_4$ -induced hepatotoxicity (Usoh et al. 2012).

Regarding the antioxidant activities or free radical scavenging, leading to the prevention and treatment of acute and chronic liver diseases (Bruck et al. 2004), silymarin (50 mg/kg) and HSARE (100 mg/kg) were found to significantly reduce lipid peroxidation and increase endogenous antioxidant activity, with a marked increase in hepatic GSH, improvement in SOD and reduction in the MDA level recorded. These enzymes (GSH and SOD) were significantly reduced with an increased level of MDA in TAA-intoxicated rats. Silymarin was reported to significantly reduce lipid peroxidation and increase endogenous antioxidants of SOD, CAT and GSH (Kiruthiga et al. 2007). Although the antioxidant effects of silymarin and HSARE were comparable with respect to GSH and MDA, HSARE showed significant improvement with respect to SOD when compared to silymarin.

With respect to inflammatory cytokines, the results revealed normalized levels of almost all examined cytokines (TNF- $\alpha$  and IL-6) in silymarin and HSARE-treated rats, apart from INF- $\gamma$ , which was not normalized in silymarin-treated rats. Both HSARE (Okoko & Ere 2012) and silymarin (Gazák et al. 2007; Saller et al. 2008) were reported to significantly reduce the levels of inflammatory cytokines. The significant improvement in all examined inflammatory cytokines as a result of HSARE could be related to an immunoprotective effect, in addition to its known anti-inflammatory activity reported by Okoko and Ere (2012), who recorded reductions in nitric oxide and catalase activities that are essential for the activation of macrophages. The normalization of inflammatory cytokines was supported by histopathological and immunohistochemical examinations, where both silymarin and HSARE resulted in positive histopathological changes as a result of TAA intoxication, which was revealed as fewer chronic inflammatory cells and diminished hepatic necrosis. The NF- $\kappa$ B and CYP2E1 levels were also significantly less, indicating almost the same mechanism of action for HSARE and silymarin to suppress hepatic injuries caused by TAA. Luster et al. (2000) demonstrated

that silymarin has a cytoprotective effect, leading to the inhibition of the transcription of NF- $\kappa$ B, responsible for the regulation of various genes involved in the inflammatory process.

Concerning apoptosis, both HSARE and silymarin increased caspase-3 activity, which may cause the upregulation of apoptosis or the renewal of liver tissue to remove the necrotic tissue as a result of TAA intoxication. TAA-induced toxicity was reported to cause the upregulation of the Bax protein and the downregulation of the antiapoptotic protein Bcl-2 and its translocation into the mitochondria, causing apoptosis (Chen et al. 2006b), while other studies suggested that the release of ROS as a result of TAA biotransformation may cause centrilobular necrosis (Sarkar & Sil 2007). Some plant extracts, such as *Curcuma longa* extract (Salama et al. 2013a), and some antioxidants, such as  $\alpha$ -lipoic acid (Moungjaroen et al. 2006), curcumin, curcuminoids (Wang et al. 2012), *Boesenbergia rotunda* and silymarin (Salama et al. 2013b), were shown to induce apoptosis.

The effects of HSARE and silymarin were comparable with respect to their hepatoprotective, antioxidant and anti-inflammatory activities, with only minute differences in SOD and INF- $\gamma$ , which were not significantly improved in silymarin-treated rats. The hepatoprotective effect of HSARE could be attributed to its delphinidin and cyanidin derivatives, reported to possess antioxidative effects (Wawer 2001). In this study, UPLC-qTOF/PDA/MS as a method of separation permitted better separation and characterization, which allows room for an increase in the dose with expected higher hepatoprotective effects.

## Conclusion

The findings in this study revealed a higher sensitivity and mass accuracy upon the use of UPLC-qTOF-PDA-MS, where 25 compounds were critically identified. Concerning the hepatoprotective and anti-inflammatory potential, both silymarin and HSARE showed comparable effects, although HSARE showed significantly higher improvements in both SOD and INF- $\gamma$ . HSARE also possessed the added value of being more water-soluble and of natural origin with fewer expected side effects, contrary to silymarin, which is a semi-synthetic preparation and known to possess side effects, e.g., diarrhoea, bloating, dyspepsia, nausea and irregular stool, due to increased bile secretion and bile flow. Future work will involve testing higher doses of HSARE.

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