

Hepatoprotective Effect of Bioactive Fraction of *Lagerstroemia speciosa* (L.) Pers. Bark Against Monosodium Glutamate-Induced Liver Toxicity

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ABSTRACT

Background: The phenolics and flavanoid enriched bioactive fraction of *L. speciosa* bark were reported for its medicinal value in various illness however hepatoprotective activity against monosodium glutamate-induced liver toxicity yet to be reported. **Objective:** To evaluate the hepatoprotective and antioxidant potential of *L. speciosa* bark extract fraction against monosodium glutamate-induced liver toxicity. **Methods:** The phytochemical constituent of ethyl acetate fraction of *L. speciosa* bark extract (LSE) were identified by GC-MS analysis. The antioxidant activity of LSE were analyzed with *in-vitro* antioxidant assay and subjected to evaluate hepatoprotective activity against monosodium glutamate induced liver toxicity in rat. **Results:** LSE evaluated as rich in phenolics and flavonoid content along with potent hepatoprotective activity. GC-MS analysis of bioactive fraction exhibits Palmitic Acid, Octadecanoic acid, 5-methyluridine, catechine, epigallocatechin, and norgestrel as major biologically active phytochemicals. Oral administration of LSE (100 and 200 mg/kg.) declined the elevated levels of the biochemical marker as well as interleukins while enhanced the enzymatic antioxidant activity and reduced the increased level of stress marker (MDA) in monosodium glutamate-induced rats. It also restored the altered expression level of pro-apoptotic genes, but there is no significant change in the expression level of the anti-apoptotic gene. LSE improved histopathology of the liver through the improvement of hepatocellular architecture, inflammation, and attenuation of vascular and cellular degeneration. **Conclusion:** The bioactive fraction of *L. speciosa* bark was found to exhibit strong antioxidant and hepatoprotection in monosodium glutamate induced liver toxicity in rats.

Key words: *Lagerstroemia speciosa*, Monosodium glutamate, Superoxide dismutase, Apoptosis.

INTRODUCTION

Monosodium glutamate is derived from a non-essential amino acid, and it is used as a food additive to enhance the flavor and taste of the food. Monosodium glutamate (MSG) commonly used in Chinese cuisine and western diet¹. The average daily consumption of MSG in industrialized countries is 0.3–1.0 gram, but it can be too much occasionally, depending on the taste preferences and MSG content used for the preparation of individual food items². Regular oral administration of MSG (60 - 100 mg) for 30 days enhanced the possibility of fibrotic and fatty liver in male rats³. Administration of a high amount of MSG can alter the function of both renal and liver⁴. Flavoring agents enhance the consumption of foods ingredients that usually lack flavor; but, that does not have nutritional significance¹. The toxic effects of monosodium glutamate result in alterations in mitochondrial function, lipid peroxidation, damages in the hypothalamic neurons, impairment in memory retention, disturbed the cellular redox, and induce hyperphagia, which leads obesity. In this context, some studies have shown that MSG induces oxidative stress and hepatotoxicity in rats^{5-7,1}.

MSG also altered endocrine functions of neonatally treated rats, which reflect the altered metabolic rate of glucose and the declined antioxidant defense system⁸. The exact mechanism of glutamate toxicity remains unknown, but earlier reports point towards apoptosis due to the generation of reactive oxygen species (ROS)⁹. Antioxidants lower oxidative stress in cells either via preventing oxidation of substrate by ROS or by ameliorating free radicals oxidation via upregulation of enzymatic antioxidants in the cells like catalase (CAT) and superoxide dismutase (SOD)¹⁰. *Lagerstroemia speciosa* (L.) is commonly known as “Jarul” in India, and it is a member of the *Lythraceae* family. It is popularly known as Pride of India, and also known as Queen’s Flowers, or Queen Crape Myrtle in English. This plant is localized in the South East-Asian countries, Philippine and India. Pharmacologically active phytochemicals isolated from different parts of *L. speciosa* have revealed antibacterial, hypoglycemic, anti-inflammatory, hepatoprotective and antioxidant properties¹¹⁻¹⁶. This experimental study was designed to evaluate the effect of LSE in the prevention of hepatotoxicity from oxidative damage induced with MSG administration in Sprague Dawley rats and its possible mechanism of action.

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MATERIAL AND METHODS

Chemicals and kits

EDTA (Ethylenediaminetetraacetic acid) and Enhanced Avian HS RT-PCR kit procured from Sigma Aldrich (USA) while biochemical reagent kit was procured from Transasia Bio-medicals Ltd. (India). MSG (Sigma Aldrich USA) and ELISA kit procured from Elabscience. All other chemicals and reagents utilized were of analytical grades and procured from Merck and Himedia Pvt. Ltd. (India).

Preparation of extract and phytochemical profiling

The collection of *L. speciosa* bark was done from National Botanical Research Institute, Lucknow (India), and authenticated by Dr. A.K.S. Rawat, scientist, Pharmacognosy and Ethnopharmacology Division National Botanical Research Institute Lucknow, India. Herbarium (NBRI / CIF /256 / 2011) was deposited in the institute for future reference. Dried samples were extracted with (70% v/v) ethanol. The extract was concentrated in a rotary evaporator and freeze-dried in a lyophilizer. The lyophilized crude extract was fractionated with solvent (hexane, chloroform, and ethyl acetate) of different polarities. The most bioactive ethyl acetate fraction (LSE) was collected based on their potent antioxidant activity, which was screened initially. The total phenolic content (TPC) was estimated by using the method given by Ragazzi and Veronese (1973) and expressed as mg gallic acid equivalent (GAE)/gram LSE¹⁷. Total flavonoid content (TFC) was analyzed by the method of Oyaizu (1986) and expressed as mg quercetin equivalent (QE)/gram LSE¹⁸. The phytochemical characterization of the bioactive fraction of *L. speciosa* bark was done via GC-MS analysis using *N*-Methyl-*N*-(trimethylsilyl) trifluoroacetamide (MSTFA) as a derivatizing agent. The derivatized sample was investigated on the GC-MS instrument comprising of a gas chromatograph (Thermo Trace GC Ultra) and mass spectrometers (Thermo fisher DSQ II). The data was recorded by mass selective detector operating in the electron impact (EI) mode with 70 eV ionization energy at an ionization current of 2.0 mA and mass range 50–800 m/z. The resultant chromatographic and mass data were acquired using Xcalibur software. The software depicts the investigation of the m/z ratio values of each metabolite fragments detected in mass spectra using GC-MS spectral library databases such as WILLY and NIST. The GC-MS based phytochemical profiling was done by comparing the m/z ratio and LRI values of each metabolite. The relative concentration of detected metabolites was calculated as a percent peak area¹⁹.

In-vitro antioxidant activity

The antioxidant potential of the extracts was evaluated with DPPH stable radical²⁰. The free radical scavenging potential of LSE was evaluated at A515 nm with a calibration curve and determined by linear regression. The results obtained are compared to those obtained for ascorbic acid, which is used as standard antioxidants. DPPH radical inhibition was evaluated according to the equation.

$$\text{DPPH}^{\cdot} \text{ radical inhibition} = \text{Control} - \text{Sample} / \text{Control} \times 100$$

Ferric Reducing power (RP) estimation was done by ferric reducing power assay and expressed as μg ascorbic acid equivalents (ASE) per mg LSE²¹.

Total antioxidant capacity was determined by using the spectrophotometric method²². Ascorbic acid was used as the standard, and the total antioxidant capacity is expressed as μg ascorbic acid equivalents (ASE) per mg LSE.

Experimental animal

Animal (Sprague Dawley adults male rats) were used in this study according to the regulations of the Institutional Animal Care

Committee, CPCSEA, India (Reg. No. 1732/GO/Re/s/13/CPCSEA). Acute toxicity studies were performed according to the guidelines (OECD Guideline 423)²³. Administration of LSE at doses of 300 mg/kg b.w. via oral gavage exhibits no mortality or any abnormal behavior in tested rats. Animals used in this study were divided into five groups, and each group has five animals. All animals in different groups except group-I (control) induced with MSG (35 mg/kg/day) orally up to 2 weeks. After induction, Group-III, and IV administered with LSE 100 and 200 mg/kg b.wt/day, p.o. respectfully. Group-V administered with vitamin E (100 mg/kg b.wt/day, p.o.). 2 mL Normal saline (vehicle) was given to group-I animal orally²⁴. After two weeks treatment period, rats were sacrificed on overnight fast by cervical dislocation. Blood was collected by heart puncture and centrifuge at 1000 rpm to separate serum for the estimation of biochemical markers and collected vital organs washed with phosphate buffer saline (PBS 1:9) and fixed in formalin for histopathology and remaining tissue stored at -20°C for enzymatic antioxidants and molecular analysis.

Determination of biochemical parameters

Biochemical kits (Transasia Bio-medicals Ltd) were used to evaluate Biochemical parameters from collected blood serum (AST, ALT, ALP, GGT, and total bilirubin) with auto chemistry analyzer (Csense 100).

Determination of antioxidant enzymes, and stress markers

Post mitochondrial supernatant (PMS) was obtained by Homogenized Liver tissue, 400 mg liver tissue was homogenized in phosphate buffer (10 mM, pH 7.4) having KCl (1.15%) and EDTA (1.15%, pH 7.4) followed by centrifugation at 12,000 \times g for 20 min at 4°C. The total protein content present in different samples was quantified by the method of Bradford (1976) at 595 nm (Spectramax 340PC, Molecular Devices, USA) by using BSA as a standard protein and expressed as mg/g FW²⁵.

Investigation of Superoxide dismutase (SOD) (EC 1.15.1.1) activity was done by the method of Kakkar *et al.* (1984)²⁶. The activity of catalase (CAT) (EC1.11.1.6) was estimated by the Aebi (1974) and exhibits as $\mu\text{mole H}_2\text{O}_2$ consumed /mg protein²⁷. Glutathione peroxidase (GPX) activity was investigated by using the method of Rotruck *et al.* (1973)²⁸. The evaluation of glutathione S-transferase (GST; EC 2.5.1.13) activity was done by using the Habig *et al.* (1974)²⁹. MDA test was done to analyze Lipid peroxidation by estimation of thiobarbituric acid reactive substances (TBARS), according to Hodges *et al.* (1999)³⁰. The reduced glutathione (GSH) content was determined by the method of Ellman (1961)³¹.

Estimation of cytokines in hepatic tissue

Interleukins (IL-2, IL-6) TNF- α , were investigated by kit protocol from liver tissue (Elabscience Biotech Co. Ltd. Wuhan, Hubei, China). Enzyme-linked immunosorbent assay (ELISA) kits were based on the principle of standard sandwich ELISA technology.

Gene expression analysis via quantitative real-time PCR (qRT-PCR)

RNA isolation accomplished by TRizol reagent. The quality and concentration of the isolated RNA were estimated at 260/280 nm using the NanoDrop instrument. Synthesis of cDNA was performed from isolated total RNA by the Enhanced Avian HS RT-PCR kit (Sigma Aldrich, USA). This cDNA worked as a template for qRT-PCR to estimate the total transcript level and using SYBR Green PCR Master Mix (Applied Biosystems, USA) in a StepOne real-time PCR system (Applied Biosystems, USA). The quantitative real-time expression of the genes was analyzed by the 2^{- $\Delta\Delta\text{Ct}$} method (Livak and Schmittgen, 2001)³². The primer sequences designed for each gene are depicted in

(Table S1).

Statistical analysis

All the estimated results are the mean of five replicates. The data were presented to Duncan's Multiple Range Test (DMRT) for the evaluation of the significant difference between the means ($p < 0.05$). SD was depicted using the average of the five replicates.

RESULTS

GC-MS analysis bioactive fraction of *L. speciosa*

3.1. GC-MS analysis of ethyl acetate fraction of *L. speciosa* bark extract carried out through mass spectrometry attached with GC. There are seven major biologically active phytochemicals that were identified from the bioactive fraction with their retention time, molecular formula, and peak area (%). The major phytochemical compounds among them were palmitic acid (1.16%), octadecanoic acid (1.94%), 5-methyluridine (29.48%), catechine (2.41%), epigallocatechin (40.06%), and norgestrel (2.59%) (Table 1, Figure 1). The compound prediction is based on a comparison of LSE chromatogram and mass spectra fragmentation patterns of computer inbuilt NIST library Databases.

Total phenolic and flavanoid content

The total phenolic and flavanoid contents in plant extract depend on the type of solvent used for their extraction procedure. Extraction yield of 70% ethanolic extract of *L. speciosa* bark was found 13.71% w/w. Total phenolic content in LSE was estimated, 180 ± 17.23 (mg GA eq/

gram LSE), and total flavonoid content was analyzed 140 ± 7.19 (mg RE eq./gram LSE).

In-vitro antioxidant studies

DPPH is a free radical that can be scavenged by accepting an electron from antioxidants and convert in a stable diamagnetic molecule. The free radical scavenging effect of the active fraction on DPPH radicals increases with increasing concentration. At 5, 10, 15, 20 and 25 $\mu\text{g/mL}$, the scavenging activities of DPPH radical were 21%, 34%, 46% 65% and 84% respectively. The calculated IC-50 value of LSE was 16 $\mu\text{g/mL}$, and reference standard ascorbic acid IC-50 evaluated 9 $\mu\text{g/mL}$. The reducing power of LSE exhibits the capability of Fe^{3+} to Fe^{2+} reduction by antioxidant. The analyzed reducing power of LSE was investigated 41.21 ± 2.43 μg ascorbic acid equivalents/mg LSE. Total antioxidant capacity was 32 ± 2.89 μg ascorbic acid equivalent/mg of LSE.

Effect of LSE on Biochemical parameters

MSG induced animals exhibit a significant increase in serum biochemical parameters at the end of the induction period. The increased biochemical parameters reverted toward normal by LSE administration. The group-II exhibits (107.24 ± 4.86 , 225.57 ± 12.76 , 289.67 ± 13.64 , 112.45 ± 4.46 , and 1.95 ± 0.9 for ALT, AST, ALP, GGT, and TBL respectively) increased level of biochemical parameters as compared to group-I (41.24 ± 4.46 , 68.43 ± 7.37 , 67.36 ± 5.55 , 38.24 ± 3.54 , and 0.424 ± 0.05). All these increased biochemical parameters in other groups (III, IV, V) were restored toward normal range, in group-III (85.45 ± 4.25 , 141.26 ± 8.7 , 175.75 ± 9.77 , 73.53 ± 3.86 , and $1.15 \pm$

Table 1: Phytochemicals Identified in Ethyl acetate fraction of *L. speciosa* bark extract through GC-MS analysis.

RT	Metabolites	Molecular formula	%Area
12.64	Glycerol, 3TMS	$\text{C}_{12}\text{H}_{32}\text{O}_3\text{Si}_3$	0.99
30.78	Palmitic Acid, TMS derivative	$\text{C}_{19}\text{H}_{40}\text{O}_2\text{Si}$	1.16
35.47	Octadecanoic acid, trimethylsilyl ester	$\text{C}_{18}\text{H}_{36}\text{O}_2$	1.94
44.17	5-Methyluridine, tris(trimethylsilyl) derivative	$\text{C}_{10}\text{H}_{14}\text{N}_2\text{O}_6$	29.48
46.15	Catechine (2R-E)-, 5TMS derivative	$\text{C}_{30}\text{H}_{54}\text{O}_6\text{Si}_5$	2.41
46.50	Epigallocatechin (6TMS)	$\text{C}_{33}\text{H}_{62}\text{O}_7\text{Si}_6$	40.06
48.21	Norgestrel, bis(trimethylsilyl) derivative	$\text{C}_{27}\text{H}_{44}\text{O}_2\text{Si}_2$	2.59

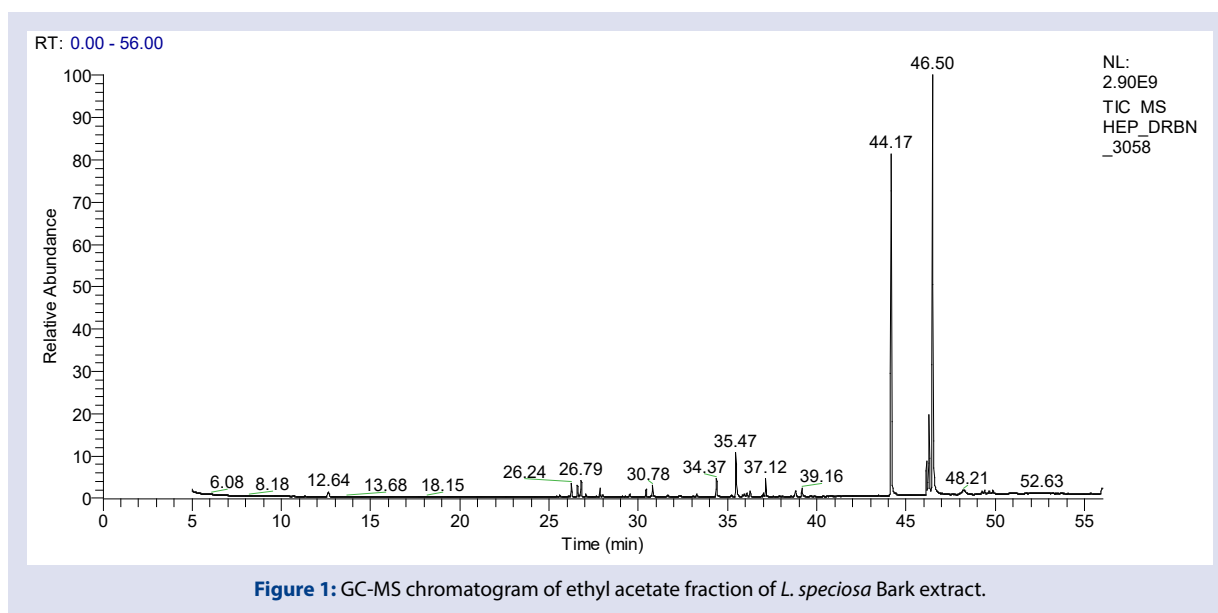


Figure 1: GC-MS chromatogram of ethyl acetate fraction of *L. speciosa* Bark extract.

0.08), group- IV (59.43 ± 3.14 , 73.87 ± 6.95 , 74.34 ± 9.43 , 51.45 ± 3.08 , and 0.624 ± 0.07) and group- V (45.46 ± 2.46 , 72.55 ± 5.48 , 69.63 ± 8.55 , 46.65 ± 2.95 , and 0.543 ± 0.06) respectively. Effective restoration of these biochemical serum markers was observed in the group- IV, which is near to the vitamin E administered group. Thus, these biochemical markers were restored toward normal range by LSE treatment in the group- III and group- IV (Figure 2).

LSE treatment improves antioxidant enzymes activity and lowers lipid peroxidation

The increased activity of SOD (Figure 3A) was found in *L. speciosa* treated group-III (69.94 ± 5.10 U/mg Protein) and group-IV (110.17 ± 6.10 U/mg Protein) as compared to the group-II (31.83 ± 3.08 U/mg Protein) and closed to the group- V (114.38 ± 5.90 U/mg Protein).

The elevated level of catalase activity was found in LSE, and vitamin E treated groups as compared to the group-II (11.65 ± 1.03 U/mg Protein). The catalase activity was improved in LSE treated group- IV (31.82 ± 1.36 U/mg Protein) and group- III (23.85 ± 1.25 U/mg Protein) near to group- V (34.08 ± 1.71 U/mg Protein) (Figure 3B).

Glutathione peroxidase (GPX) activity was found to be increased in all treated groups as compared to the group- II (1.984 ± 0.238 U/mg

Protein). The increased activity was shown in the LSE administered group- IV (5.122 ± 0.243 U/mg Protein), and group- III (3.25 ± 0.178 U/mg Protein) which was almost equal to the group- V (5.74 ± 0.256 U/mg Protein) (Figure 3C).

GST (glutathione-S-transferase) activity was found to be elevated in all treatment groups as compared to the group- II (0.735 ± 0.06 U/mg Protein). The elevated activity was shown 1.978 ± 0.102 U/mg Protein in the group- IV as compared to the group- III (1.374 ± 0.097 U/mg Protein), while the activity of GST in the group- IV is closed to the group- V (2.178 ± 0.091 U/mg Protein) (Figure 3D).

Reduced glutathione (GSH) was found to be increased in all LSE treated groups as compared to the group-II (37.757 ± 2.26 μ mol/ mg Protein). The elevation was shown 69.34 ± 2.23 μ mol/ mg Protein in group-IV as compared to the group- III (54.654 ± 3.58 μ mol/ mg Protein) and closed to group-V (75.46 ± 4.55 μ mol/ mg Protein) (Figure 3E).

The level of MDA was estimated as LPO marker; the declining level of MDA was found in the treatment group as compared to the group-II (5.143 ± 0.23 nmoles/mg Protein), which showed the declined lipid peroxidation potential of LSE in MSG induced increase in stress marker (Figure 3F).

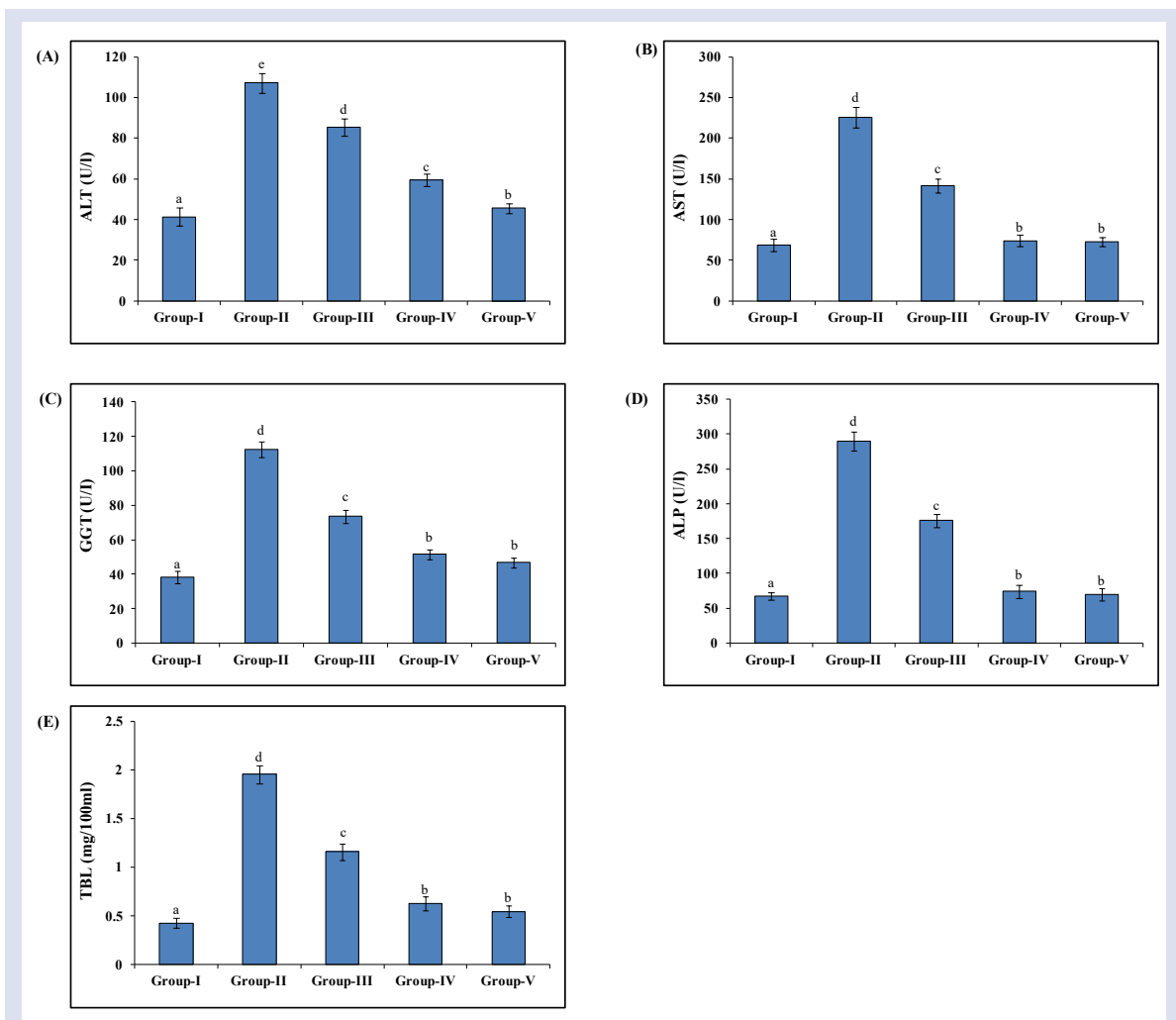
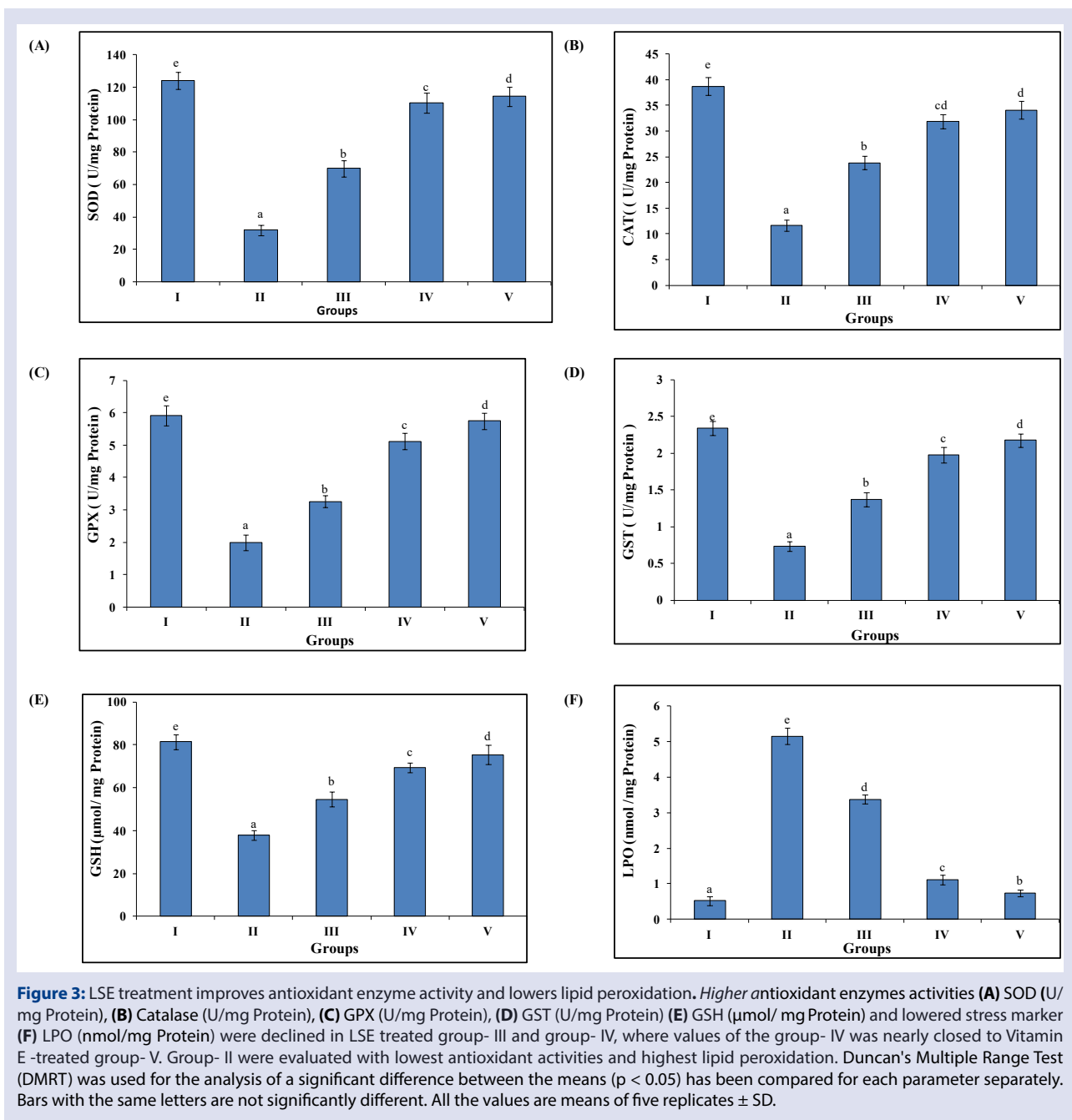


Figure 2: LSE maintained the reduced levels of various biochemical parameters in the group- III and group- IV. (A) ALT (IU/L), (B) AST (IU/L), (C) GGT (IU/L), (D) ALP (IU/L), (E) TBL (mg/100ml). All biochemical parameters were reversed by the test drug (LSE) administration in the group- III and group- IV nearly closed to Vitamin E treated group- V, while group- II showed elevated levels. Duncan's Multiple Range Test (DMRT) was used for the analysis of a significant difference between the means ($p < 0.05$) has been compared for each parameter separately. Bars with the same letters are not significantly different. All the values are means of five replicates \pm SD.



Elisa quantified interleukins and TNF-α

Inflammatory cytokines such as IL-2, IL-6, TNF-α, play a significant role in hepatic disease. The increased tissue concentration of IL-2 was observed in the group- II (18.75 ± 1.04 ng/mL) as compared to the group- I (5.24 ± 0.68 ng/mL), which was significantly declined in the treatment group- III (12.34 ± 0.43 ng/mL) and Group- IV (8.47 ± 0.56 ng/mL) which was closed to the group- V (6.39 ± 0.79 ng/mL) (Figure 4A).

The treatment of test drugs was found to be declined elevated IL-6 in the group- IV group (7.57 ± 0.847 ng/mL), which was almost closed to the group- V (6.40 ± 0.79 ng/ mL) and most significant than the group- III (13.44 ± 1.068 ng/ mL) while the declined concentration in the group-II was estimated 22.07 ± 1.30 ng/mL (Figure 4B).

The level of TNF alpha was found to be elevated in all experimental groups except group-I (509.54 ± 60.75 pg/mL). The level of TNF alpha

was reduced in LSE administered group- III (1736.57 ± 107.46 pg/mL) and group- IV (935.098 ± 68.67 pg/mL) as compared to group-II (2470.899 ± 143.66 pg/mL) but closed to group-V (723.546 ± 73.67 pg/ mL) (Figure 4C).

Quantitative real-time polymerase chain reaction (qRT-PCR) investigation

Gene expression level analysis of Bcl-2, p53, Bax, caspase-3, and caspase-9 performed by qRT-PCR. The expression level of the anti-apoptotic gene Bcl-2 showed no significant changes in all the groups while the expression level of the pro-apoptotic gene (p53, Bax, caspases-3, and caspase-9) was upregulated in group-II. These expression levels of pro-apoptotic genes reverted back toward normal in the LSE administered group-III and group-IV. The efficacy of 200 mg/kg LSE administration was nearest to vitamin E treated group-V, and 200mg/kg LSE administration was found to be most significant

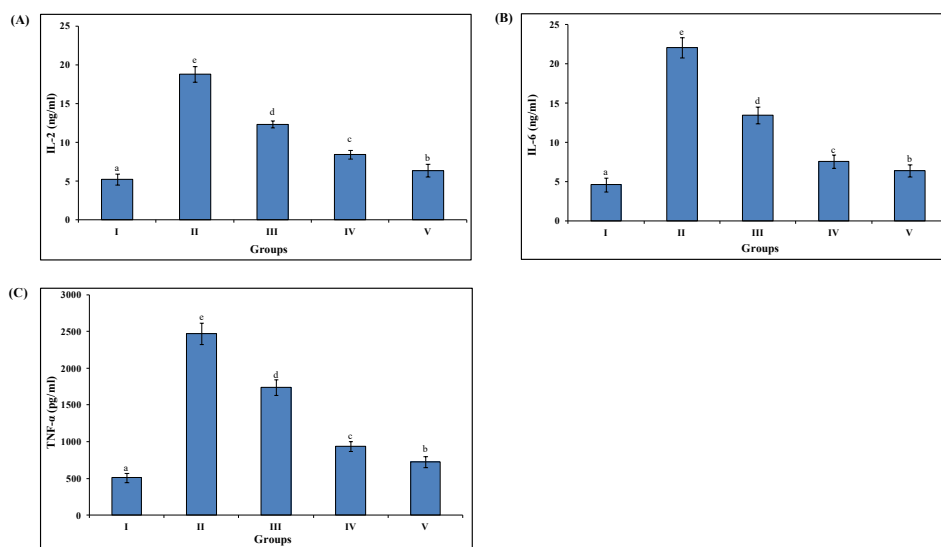


Figure 4: LSE treatment reduced the levels of interleukins and TNF-α. LSE treatment reduced the level of interleukins, (A) IL-2 (ng/ml), (B) IL-6 (ng/ml), and (C) TNF-α (pg/ml) in group-III and group-IV. All values of the group-IV were nearly closed to the Vitamin E-treated group-V. Group -II has elevated levels of all parameters. Duncan's Multiple Range Test (DMRT) was used for the analysis of a significant difference between the means ($p < 0.05$) has been compared for each parameter separately. Bars with the same letters are not significantly different. All the values are means of five replicates \pm SD.

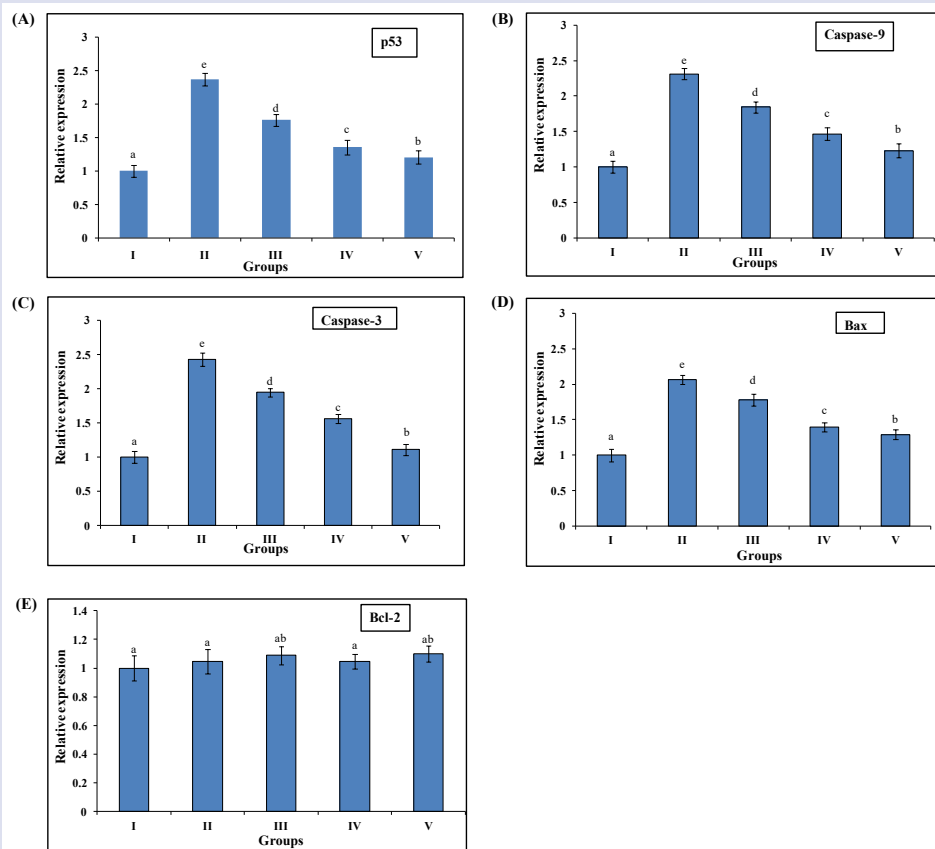


Figure 5: Relative expression of Antiapoptotic and Pro-apoptotic genes by qRT-PCR under MSG induced liver toxicity. The relative expression of pro-apoptotic genes (A) p53, (B) caspase-9 (C) Caspase-3, (D) Bax, were downregulated in all treatment groups as compared to the group- II. Antiapoptotic gene Bcl-2 has no significant change in their expression level (E). Duncan's Multiple Range Test (DMRT) was used for the analysis of a significant difference between the means ($p < 0.05$) has been compared for each parameter separately. Bars with the same letters are not significantly different. All the values are means of five replicates \pm SD.

than 100 mg/kg. The relative expression of the pro-apoptotic and anti-apoptotic gene was observed by using gene-specific primers. GAPDH gene primer was taken as an endogenous control. The results analysis depicted the levels of p53, Bax, caspase-3, and caspase-9 were found to be highly expressed in Group-II as compared to the group-I .while all treatment group exhibits significant restoration of the expression level of all genes (Figure 5).

Histopathological studies

Microscopic examination of the liver section was done at (40X); H & E stain of a group -I exhibit healthy cells having well preserved granulated cytoplasm and sinusoidal spaces with prominent nucleus and nucleolus (Figure 6A). The liver section from MSG treated rats exhibited disarrangement of normal hepatic cells with centrilobular necrosis, vascular and cellular degeneration, and inflammation (Figure 6B). The reticuloendothelial system was much more improved in Vit-E, and LSE treated groups. The administration of LSE in group-III and IV exhibited a reduction of necrosis with decreased inflammation and improved the

architecture of hepatocytes. The level of improvement in the group -IV was much better than group-III and close to group -V (Figure 6C,D,E).

DISCUSSION

This study was targeted to evaluate the hepatoprotective effect of LSE in Rat induced with monosodium glutamate oxidative stress and hepatic toxicity. Most probably monosodium glutamate caused alterations in the liver because liver are mainly accountable for the detoxification of drug and chemical compounds in the body³³. Monosodium glutamate caused elevation of serum biochemical (ALT, AST, ALP, GGT, and TBL) in the blood which might be due to cellular leakage of hepatic cell membranes, on the other hand, reduced levels antioxidant defence system, Such results were consistent with previous study^{6,34,35}, the conducted experiment revealed LSE treatment was found to be reduced liver injury marker (ALT, AST, ALP, GGT and TBL) and ROS with enhanced antioxidant enzyme and improved histopathology of liver. GCMS analysis of LSE depicts the presence of Palmitic Acid, Octadecanoic acid, 5-Methyluridine, Catechine, Epigallocatechin, and Norgestrel as major

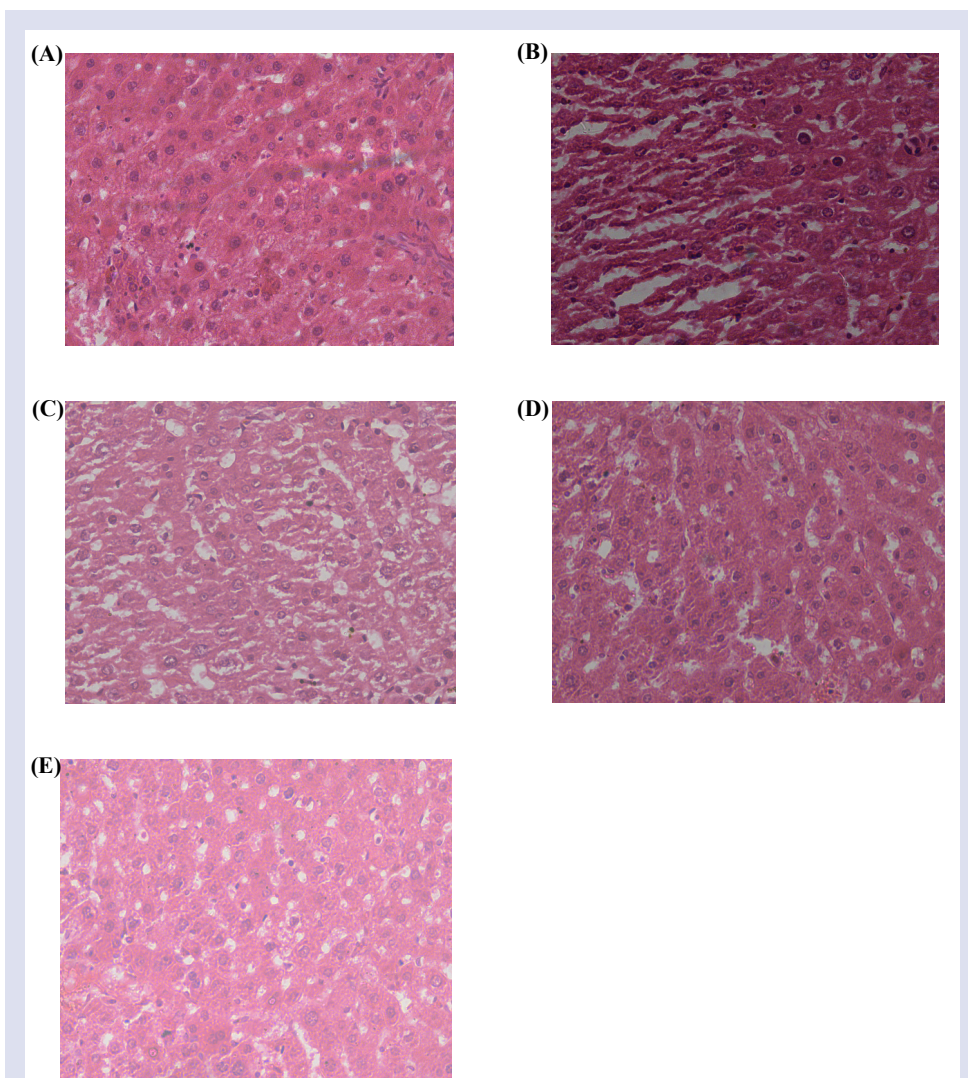


Figure 6: LSE improved the architecture of hepatocytes' lesion and inflammation. (A) (40X); H & E stain of the group - I exhibit normal cells having well preserved granulated cytoplasm and sinusoidal spaces with prominent nucleus and nucleolus. (B) Liver section from MSG treated rats exhibited disarrangement of normal hepatic cells with centrilobular necrosis, vascular and cellular degeneration, with inflammation. (C, D & E) Administration of LSE in group-III and IV exhibited a reduction of necrosis with decreased inflammation and improved the architecture of hepatocytes, the level of improvement in the group-IV was much better than the group-III and close to the group-V.

biologically active phytochemicals. Palmitic Acid is explored as an anti-inflammatory agent in liver toxicity³⁶. Catechins and epicatechins are the sources of several tannins. It is a condensed tannin like friedelin; it reduces gastric hypersecretion and exhibits strong antioxidant and anti-inflammatory activity^{37,38}. Methyluridine was reported as antimicrobial and anticancer compound³⁹. The liver is an essential vital organ that metabolizes various toxins and nutrients to keep the healthy human body with normal biochemical and physiological action; it should always function properly. The mechanism of xenobiotics to induce liver toxicity is still unclear, although cytochrome 450 involved in the biotransformation of drug and metabolite. The reactive chemicals may enhance reactive oxygen species either by GSH depletion or redox cycling⁴⁰. Phenolic and flavonoid are potential therapeutic components of many plants and exhibits its role in the prevention and cure of many diseases because these compounds possess major hydroxyls group which are responsible for the prevention of reactive oxygen species^{41,42}. Liver injury is strongly associated with free radical stress, which may be ameliorated partially by antioxidant entities. The DPPH radical (DPPH[•]) is a stable radical and can gain an electron or hydrogen radical and form a stable diamagnetic compound producing a change in color. The percent color change of DPPH has been widely used to estimate the radical scavenging capacity⁴³. The presence of reducing agents (i.e., antioxidants) in active fraction results in the reduction of the Fe³⁺/ferricyanide complex to the ferrous form. Therefore, measuring the formation of Prussian blue at 700 nm can monitor the Fe²⁺ concentration. The reducing ability of the LSE increased with an increase in concentration⁴⁴. After dissociation of MSG, L-glutamate converts into glutamine and accumulated in hepatocytes and caused its damage. MSG caused alteration in hepatic parenchyma, with pyknotic form dilated sinusoids, and inflammatory cells⁴⁵. Moreover, MSG administration initially attacked the peripheral hepatocytes in the central lobules of the liver tissues leading to hepatocellular degeneration⁴⁶. Accordingly, MSG induction depicted the elevation of ALT, AST, ALP, GGT, and TBL, which may be reflected hepatic leakage and instability of cell membrane in the liver in case of hepatocellular injury⁴⁷. In this study, MSG consumption induces oxidative stress in hepatocytes which react with the polyunsaturated fatty acids of cell membrane caused mutilation of plasma membranes and mitochondrial stability, which is ensured by significant increased MDA level in liver and the declined level of antioxidant enzymes and molecule, these results supported with some earlier experimental studies⁶. Moreover, increased intracellular calcium can induce free radical generation through glutamate receptors to result in augmented GSH levels that could enhance ROS-induced liver cell damage. GSH have their crucial role in endogenous and exogenous defenses against tissue damage by free radicals⁴⁸. Various biological processes regulated by GSH like xenobiotics detoxification, synthesis of nucleic acid and protein, sulfate transport, conjugation of metabolites, signal transduction in the synthesis of protein, enzyme regulation, and induction of stress-responsive genes on ROS stress⁴⁹. The ratio of reduced and oxidized GSH imbalance and shifted toward oxidized GSH (GSSG) in the MSG induced animals, which is an indicator of tissue damage^{50,51}. Superoxide dismutase (SOD) protects against superoxide free-radicals and, it converts superoxide radicals to H₂O₂ and O₂, further CAT and GPX catalyzed H₂O₂ to H₂O. In this way, these enzymatic defense system works against oxidative stress generated by MSG induction⁴⁸. GPX is an intracellular selenium-containing enzymatic protein located in the cytosol and mitochondrial matrix protect against oxidative stress generated by MSG induction. GSH reduces lipid and nonlipid hydroperoxides with double molecules of GSH and declines the level of the detrimental molecule⁵². GST catalyzed thiol group conjugation with electrophilic xenobiotics to eliminate or conversion in mercapturic acid⁵³. Antioxidant-rich fraction competes with xenobiotic activation and metabolizes activated xenobiotics in non-toxic molecules⁵⁴.

The increased activity of GST induced higher coupling of GSH with electrophilic intermediate in the treatment group as compared to the group-II. The interleukins (IL-2, IL-6) and TNF-alpha were elevated in MSG administered animals and seems to stimulate inflammation in hepatic tissue^{55,56}. LSE administration exhibits the significant reduction of interleukins (IL-2, IL-6) and TNF-alpha but the most potent reduction was shown in Vit-E treated animal (figure. 4). The gene expression study revealed an increase in the expression of p⁵³, Bax, Caspase-9, and Caspase-3, while Bcl-2 has no change in MSG administered groups; hence MSG accelerated the process of apoptosis, in this pathway, BAX protein interacts with Cyt-C released from mitochondria and further binds with procaspase-9 and Apaf-1 to form an apoptosome⁵⁷. Caspase-3 and Caspase-9 are involved in apoptosis through the process of apoptosome formation. In this study, the LSE and Vit-E were found to restore altered gene expression effectively⁵⁸. Hence, the outcome of this study exhibit valuable restoration effect of LSE on different altered liver injury markers but less effective than Vit-E.

CONCLUSION

The results of the present study have shown that the slight increase in MSG above the recorded safe limit is capable of producing alterations in the liver functions by inducing oxidative stress and inflammation in the liver. On the other side, LSE has been shown to suppress liver toxicity via inhibiting oxidative damage. So, foods containing MSG can be fortified with minute quantities of LSE to overcome their adverse effects. The declined biochemical marker and enhance enzymatic antioxidant by LSE treatment revealed that LSE improved MSG induced hepatotoxicity and oxidative stress. LSE treatment declined the increased level of IL-6, IL-2, and TNF alpha, suggesting LSE attenuated monosodium glutamate-induced inflammatory cascade in the liver and brought it back toward normal. The molecular studies indicate the MSG administration induces the progression of apoptotic pathways, which was reversed by LSE significantly. These outcomes support the existence of pharmacologically active phytochemical in LSE fraction with their potent antioxidant, anti-inflammatory, and anti-apoptotic activities. Detail clinical studies have to be required to confirm the safety and benefits of LSE before it used in human beings.

DECLARATION OF INTEREST

All the authors have no conflicts of interest.

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AUTHOR'S CONTRIBUTION

Conceptualization: LCP & Ch.VR, Formal analysis: LCP and AK, Funding acquisition: Ch.VR, Investigation: LCP & AK, Methodology: LCP & AK, Project administration: Ch.VR, Supervision: Ch.VR, and VP, Validation: LCP, Writing- review & editing: Ch.VR & LCP.

ABBREVIATIONS

ALT = alanine aminotransferase, EDTA = Ethylene diamine tetracetic acid, GSH = reduced glutathione, SOD = superoxide dismutase, MDA = Malonaldehyde, AST = aspartate aminotransferase, GC-MS = Gas chromatography mass spectrometry, = DPPH = 2,2-diphenyl-1-picrylhydrazyl, RP = reducing power, CEA = Carcinoembryonic antigen, IL-2 = interleukin-2 IL-6 = interleukin-6, TNF- α = tumor necrosis factor-alpha, SOD = superoxide dismutase, GPx = glutathione peroxidase, GSH = glutathione, GST = glutathione transferase, ELISA = enzyme-linked immunosorbent assay.

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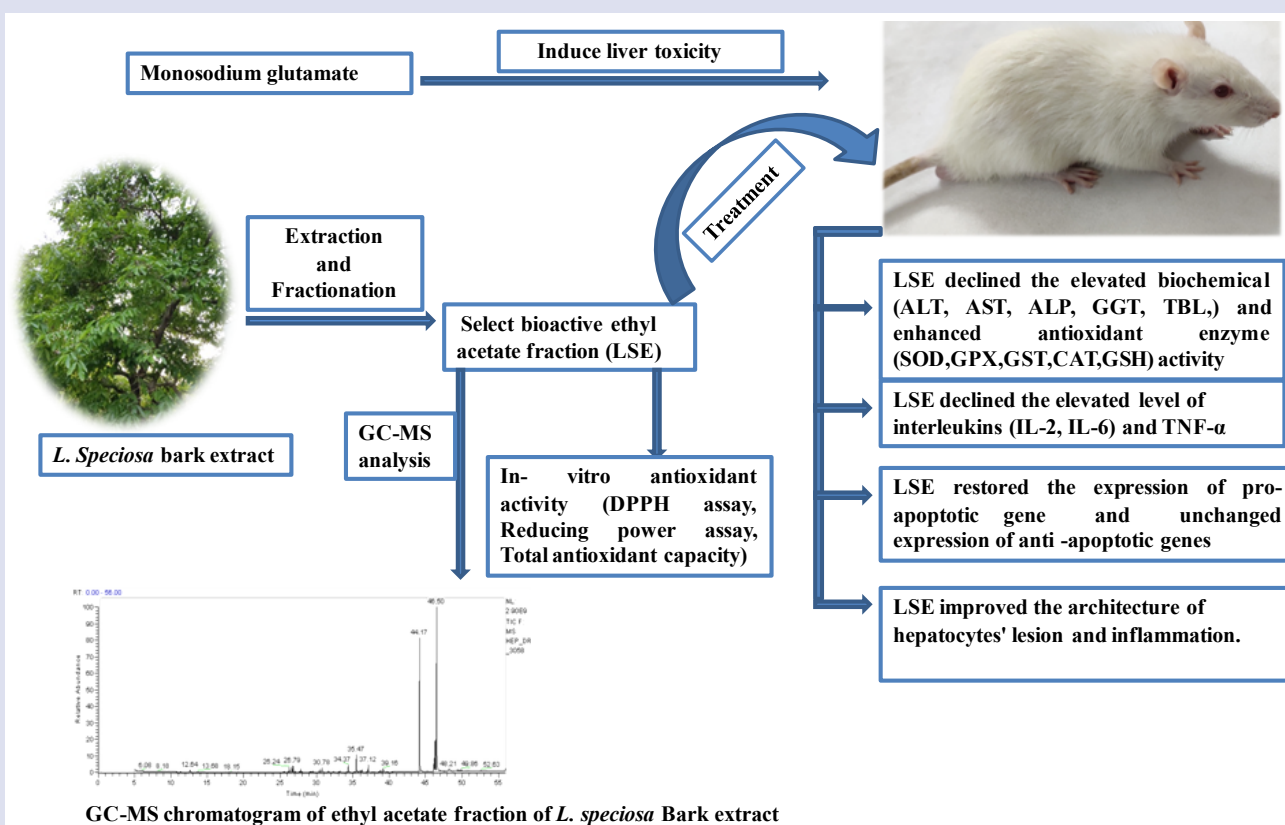
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GRAPHICAL ABSTRACT



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