Protective effect of Moringa oleifera Lam. leaf extract against carbon tetrachloride-induced neuroinflammation in a mouse model of hepatic encephalopathy

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Research

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Abstract

Background

Hepatic encephalopathy (HE) is a neuropsychiatric disorder associated with acute or chronic liver injury. Carbon tetrachloride (CCl₄) is usually used as an experimental model for HE. The present study aimed to assess the neuroprotective impacts of Moringa oleifera Lam. leaf ethanolic extract (MOLE) against neurotoxicity in CCl₄-induced mouse model of HE.

Methods

High-performance liquid chromatography (HPLC) analysis was used for the detection of marker compounds; rutin and β-sitosterol. Animals were divided into four groups; vehicle group, CCl₄ treated group, MOLE treated group, and (CCl₄ + MOLE) group treated with MOLE for 14 days before inducing neurotoxicity by CCl₄.

Results

Pretreatment with MOLE decreased alanine aminotransferase (ALT), aspartate aminotransferase (AST), corticosterone, and ammonia levels in serum as well as it improved the antioxidant status of CCl₄ treated mice in the tissue of hippocampus (HC) and cerebral cortex (CC). It reduced the expression of toll-like receptor (TLR)4, TLR2, myeloid differentiation primary response 88 (MYD88), and nuclear factor kappa B (NF-κB) genes and the protein levels of the pro-inflammatory cytokines in the selected brain regions. MOLE also exhibited anti-apoptotic effect as revealed by the reduced expression of caspase3, and prevented histological deteriorations caused by CCl₄ treatment. Furthermore, CCl₄-induced anxiety and depression-like behavioral changes were attenuated by MOLE preadministration.

Conclusions

Taken together, the current results suggest significant anxiolytic and antidepressant effects of MOLE via modulation of neuroinflammation, oxidative stress, TLR4/2-MyD88/NF-κB pathway, and apoptosis in HE experimental model.

Background

It is well known that people with liver disease suffer from neuropsychiatric disorders due to alteration of lipid peroxidative and antioxidative mechanisms in the brain along with severe hepatic encephalopathy (HE) propagation. HE is a neuropsychiatric disorder propagated as a result of acute or chronic hepatic
failure \(^2\). Carbon tetrachloride (CCl\(_4\)) is a toxic substance that is used to induce liver injury with concomitant brain disorders and can be used as an experimental model for HE \(^3\).

In the liver, CCl\(_4\) is metabolized to highly reactive free radicals which oxidize fatty acids in the phospholipids of cell membranes leading to structural and functional changes in these membranes \(^4\). Moreover, these free radicals along with CCl\(_4\) itself cause injuries in the endoplasmic reticulum with a consequent effect on protein synthesis and lead to lipid accumulation \(^4\). Meanwhile, CCl\(_4\) leads to the production of inflammatory mediators from the triggered macrophages in the liver with concomitant systemic inflammation exerting a critical role in aggravating neurological manifestations, possibly through triggering the brain predisposition to the associated hyperammonemia \(^5\).

Neuroinflammation and oxidative stress have been evidenced to be involved in the development of depression and anxiety \(^6\). Noteworthy, CCl\(_4\) has been reported to induce neuropsychiatric disorders mimicking what appear in patients with acute or chronic liver damage via targeting brain antioxidative system and inflammatory pathways such as toll-like receptor (TLR)4/nuclear factor kappa B (NF-κB) pathway \(^1\). Moreover, it has been reported that corticotrophin-releasing factor (CRF) hypersecretion in response to the release of pro-inflammatory cytokines is attributable to the modification of hypothalamic-pituitary-adrenal (HPA)-axis resulting in an elevated level of plasma corticosterone and depression symptoms’ exacerbation \(^7\).

Plants are well known for having therapeutic effects and have been used in this regard in traditional and modern medications. *Moringa oleifera* Lam. (MOL), family Moringaceae \(^8\) is widely known as “Miracle tree” as a result of its medicinal outcomes. Leaves, the most common and suitable part of the plant for medicinal commercial mass production \(^9\) can be consumed in different ways and can also be dried and stored for months without losing nutritional benefits \(^10\). The following compounds were recorded in our previous study in *Moringa oleifera* Lam. leaf ethanolic extract (MOLE); quercetagetin-7-O-glucoside, quercetin 3,5,7,3',4'-pentamethyl ether, and β-sitosterol along with other phytochemicals using gas chromatography-mass spectrometry (GC-MS) analysis \(^11\). Moreover, the total phenolic and the total flavonoid contents in MOLE were also measured \(^11\). Furthermore, it has been reported that MOL possesses pain relief, anti-depression, anti-inflammatory, immunomodulatory, and neuroprotective activities \(^12\).

The current study is the first-ever to evaluate the possible role of MOLE as a protective adjuvant against brain manifestations induced by CCl\(_4\) in HE mouse model. The alleviation potentials of MOLE were assessed by tracking its effect on TLR4/2-MyD88/NF-κB pathway, neuroinflammation, apoptosis, oxidative stress, anxiety and depression-like behavior, and histopathological changes in the hippocampus (HC) and cerebral cortex (CC) regions of the mouse brain.

**Methods**
Chemicals

CCl₄ was purchased from Sigma (St. Louis, MO, USA). All other chemicals and reagents used were of the highest analytical grade.

MOL source and identification

The source for the plant leaves was from Jazan city, KSA with latitude 16° 53' 12.59” N and longitude: 42° 33' 23.99” E coordinates according to degrees minutes seconds (DMS). The authentication of the plant was carried out by taking the herbarium specimens found at Jazan University Herbarium (JAZUH), KSA, as a reference.

Preparation of MOLE

MOL leaves were washed, dried, and finally ground. 96% ethanol was mixed with the ground leaves and the mixture was kept in the shaking incubator for 24 h at 37 °C. The obtained extract was then filtered and put in the rotary evaporator at 40 °C until complete evaporation of ethanol. Finally, a semi-solid extract was produced and stored at 4 °C until use.

High-performance liquid chromatography (HPLC) analysis

MOLE was analyzed using HPLC method for a qualitative analysis of two marker compounds. About 50 mg of the extract were dissolved in 25 mL methanol and injected into an HPLC (Agilent 1200 series, UV detector). For rutin, Agilent Eclipse XDB-C18 (150 x 4.6 mm, 5 µm), wavelength 254 nm, and flow rate of 1 mL/min. The mobile phase consisted of acetonitrile: water/0.1 formic acid with gradient increased from 5% to 95% over 15 min. For β-sitosterol, a waters symmetry shield C18 column (150 x 4.6, 5 µm) and wavelength 210 nm was used. The mobile phase consisted of methanol: acetonitrile with the ratio 30:70 (v/v), with a flow rate of 1.0 mL/min.

Experimental design

Adult healthy BALB/c male albino mice weighting 20 – 25 g (8 weeks old) were brought from the National Cancer Institute (NCI). Throughout the experiment, animals were kept in conventional cages at the standard conditions of temperature, humidity, and light/dark cycle. Animals had free access to the standard food and drink ad libitum. Animal experimentation protocols were carried out following the National Institutes of Health (NIH) guidelines for animal experimentation and approved by Cairo University Institutional Animal Care and Use Committee (CU-IACUC), Egypt, (permission number: CU/I/F/41/20).

The experiment lasted for 15 days following 1-week acclimatization. Animals were haphazardly divided into 4 groups with 8 mice in each group; Control group (group 1), CCl₄-treated group (group 2), MOLE-treated group (group 3), and CCl₄+MOLE-treated group (group 4). The first two groups received distilled water orally by gavage on daily basis for consecutive 14 days. The last two groups received MOLE (400
mg/kg body weight) orally by gavage daily for consecutive 14 days. Then on day 15, group 2 and group 4 were administered a single dose of CCl₄ (1 mL/kg body weight) prepared by dilution in olive oil; 1:1 (v/v), intraperitoneally (i.p.) ¹,¹⁴, while other groups (groups 1 & 3) received olive oil, (i.p). 24 hours later, behavioral tests were carried out in separate animal groups ¹. Euthanasia was conducted by decapitation under xylazine/ketamine anesthesia ¹⁵.

**Estimation of depression-like behavior by forced swimming test (FST) and tail suspension test (TST)**

FST was carried out as described by Porsolt et al. (1977) ¹⁶. Initially, each mouse was placed into water at a depth of 20 cm and a temperature of 23 ± 2 °C inside a transparent cylinder. Afterward, the mice individually were forced to swim for 6 min. The time of immobility was recorded by considering the halt of escape-oriented behavior during the last 5 min.

TST was also executed as reported by Steru et al. (1985) ¹⁷. For 6 min, each mouse was hung about 1 cm from the tip of the tail by using sticky tape on the edge of a rod at a height of 50 cm above the floor. The duration of immobility time was considered by recording the time during which each mouse was suspended without any activity or any motion in the last 5 min.

**Collection of blood and tissue samples**

The blood was collected and the serum was isolated by centrifugation of the blood at 2000 x g for 15 min at 4 °C for biochemical analysis.

Brains were dissected from the skull. For the histopathological investigation, one side of each brain was kept in 10% neutral buffered formalin for later use. HC and CC were excised from the other side and assigned into two portions. The first portion of each region was homogenized, centrifuged at 5000 x g, and the protein concentration was evaluated in the tissue supernatant according to the Bradford method by using Biorad assay kit ¹⁸. The analysis of oxidative stress parameters and pro-inflammatory cytokines in the supernatant was followed. The second portion from each brain region was collected in RNA lysis buffer for measuring gene expression of the inflammatory and apoptotic mediators. For the histopathological investigation, brains were kept in 10% neutral buffered formalin.

**Histopathological examination**

After washing the tissue samples, a series of diluted alcohol was used for dehydration, followed by clearance in xylene, infiltration in paraffin wax, and embedding in paraffin wax blocks. For the histopathological investigation, 5μm thickness coronal sections were obtained and stained with Ehrlich’s hematoxylin and eosin (H&E) as demonstrated by Bancroft and Gamble (2008) ¹⁹. The thickness of dentate gyrus (DG) in HC was measured in different groups using ImageJ software.

**Quantitative reverse transcription-polymerase chain reaction (RT-qPCR) assay**
Total RNA was isolated from the HC and CC tissues using SV Total RNA Isolation System (Promega Corporation, Madison, WI, USA) as previously described. RNA concentration and purity were analyzed using NanoDrop™ 2000/2000c Spectrophotometer (ThermoScientific, Lo, UK). Complementary DNA (cDNA) was then produced using SuperScript III First-Strand Synthesis System according to the manufacturer's instructions (Fermentas, Waltham, MA, USA). The cDNA yield was then used to detect the relative expression levels of TLR2, TLR4, myeloid differentiation primary response 88 (MYD88), and NF-κB genes. Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) housekeeping gene was used for data normalization. Table 1 shows the primers’ sequences of mice’s genes that have been used in the present study.

### Determination of proinflammatory cytokines in HC and CC

The protein level of tumor necrosis factor (TNF)-α and interleukin (IL)-6 was detected in HC and CC supernatant by enzyme-linked immunosorbent assay (ELISA) kits particularly for mice (Merck Millipore, San Francisco, California, USA) following the producer's protocol. Protein levels were measured using the microplate ELISA reader at 450 nm.

### Evaluation of biochemical parameters

#### Alanine aminotransferase (ALT) and aspartate aminotransferase (AST) levels in serum

Liver functions were checked by measuring ALT as described by Hafkenscheid and Dijt (1979) and AST according to Sampson et al. (1980) in serum using the enzymatic methods.

#### Detection of ammonia level in serum

Ammonia assay was used to assess the level of ammonia in serum, as described by Gutiérrez-de-Juan et al. (2017). The reaction of Nessler’s reagent is the key to detect ammonia production using ammonium chloride as a standard. The spectrophotometer was used at 425 nm and the results were presented in percentage.

#### Evaluation of corticosterone level in serum

Serum corticosterone concentration was determined in serum by using ELISA kits (ThermoScientific, Lo, UK) as per the manufacturer’s instructions.

#### Determination of malondialdehyde (MDA) level in the HC and CC

Measurement of lipid peroxidation (LPO) in the homogenates’ supernatant of each brain region was carried out based on thiobarbituric acid (TBA) reaction with MDA. The principle for the reaction is the formation of a product due to LPO of the membranes. After incubation, the spectrophotometer was used to record the absorbance at 532 nm (MDA Colorimetric/Fluorometric Assay kit, Biovision Inc., CA, USA).
Enzymatic and non-enzymatic antioxidants’ level in HC and CC

OxiSelect Superoxide dismutase (SOD) kit (CellBiolabs, Inc., CA, USA) was used for the detection of the activity of SOD as described by the producer’s protocol following the method reported by (Valentine and Hart, 2003) \(^\text{25}\). The absorbance was recorded spectrophotometrically at 540 nm.

Reduced glutathione (GSH) level was measured using the method modified by Jollow et al. (1974) \(^\text{26}\). The basis for the assay depends on the formation of yellow color as a result of the reaction between 5, 5-dithiobis-2 nitro benzoic acid (DTNB) and free thiol groups of GSH. The absorbance was assessed spectrophotometrically at 412 nm.

Statistical methods

Statistical analyses were performed using GraphPad PRISM (version 8.4.3 (686); Graph Pad Software, USA). Data were represented as mean ± SD. analyses were done using one-way analysis of variance (ANOVA) followed by Tukey’s multiple comparison test. Differences were considered significant at \(P < 0.05\).

Results

Rutin and \(\beta\)-sitosterol detection by HPLC

For HPLC analysis, the authentic reference standards of rutin and \(\beta\)-sitosterol were analyzed using an identical chromatographic method. By comparing peaks’ retention times, rutin and \(\beta\)-sitosterol were identified in MOLE (Fig. 1).

MOLE pretreatment reverses the increase in serum aminotransferases, ammonia, and corticosterone levels induced by \(\text{CCl}_4\)

\(\text{CCl}_4\) treatment for 24 hours remarkably increased ALT (\(P < 0.01\)) and AST (\(P < 0.001\)). These increases were markedly prevented in response to MOLE pretreatment at significances of (\(P < 0.01\)) and (\(P < 0.001\)) for ALT and AST, respectively (Fig. 2 A, B).

An obvious elevation in serum ammonia level was manifested in \(\text{CCl}_4\)-treated mice (\(P < 0.0001\)). However, MOLE pretreatment remarkably reversed this change (\(P < 0.0001\)), Fig. 2 (C).

Serum corticosterone level was measured to investigate the potential role of MOLE against the alteration in HPA-axis induced by \(\text{CCl}_4\) treatment. A significant increase in serum corticosterone level was found in \(\text{CCl}_4\)-treated mice (\(P < 0.01\)). However, this increase was markedly prevented upon MOLE pretreatment (\(P < 0.01\)), Fig. 2 (D).

MOLE pretreatment alleviates oxidative stress induced by \(\text{CCl}_4\)
The role of MOLE pretreatment against CCl₄ LPO was investigated by measuring MDA in the HC and CC brain tissues. CCl₄ markedly elevated LPO in both HC and CC as evidenced by the striking increase in MDA (P < 0.0001). This effect was remarkably attenuated in both tissues in the presence of MOLE pretreatment (P < 0.001), Fig. 3 (A, B).

To evaluate the potential antioxidant power of MOLE against CCl₄ neurotoxicity, SOD and GSH levels were measured in the HC and CC brain tissues. CCl₄ markedly enhanced the oxidative stress as evidenced by the significant decrease in SOD and GSH levels in the HC (P < 0.05) and CC (P < 0.01). However, MOLE administration before CCl₄ treatment significantly attenuated these decreases in both tissues (P < 0.05), Fig. 3 (C-F).

**The effect of MOLE pretreatment against inflammatory response initiated by CCl₄**

**MOLE pretreatment effect on TLR4/2-MyD88/NF-κB pathway**

To evaluate the role of MOLE pretreatment on the TLR4/2-MyD88/NF-κB pathway, RT-qPCR was used to measure the expression of TLR2, TLR4, MyD88, and NF-κB genes in the HC and CC brain tissues. CCl₄ markedly raised the gene expression of TLR2 (P < 0.001), TLR4 (P < 0.001) and MyD88 (P < 0.0001) in both brain regions. NF-κB gene expression also upregulated significantly in response to CCl₄ toxicity in the HC (P < 0.01) and CC (P < 0.05). Most of these increases were significantly alleviated in response to MOLE pretreatment. TLR2 gene expression was significantly decreased in both HC (P < 0.01) and CC (P < 0.001). TLR4 was significantly decreased in both tissues as well (P < 0.001). Marked restoration of MyD88 gene expression was also noticed (P < 0.0001). Although a decrease in NF-κB gene expression was noticed in both tissues, it was only significant in the HC (P < 0.01), (Table 2).

**MOLE pretreatment reverses alterations in TNF-α and IL-6 levels induced by CCl₄**

The effect of MOLE pretreatment on the inflammatory mediators activated by CCl₄ was evaluated by measuring TNF-α and IL-6 protein levels in the HC and CC brain tissues in different groups. CCl₄ significantly elevated the protein levels of TNF-α and IL-6 in both HC and CC (P < 0.01). Nevertheless, pretreatment with MOLE remarkably prevented these elevations in the HC (P < 0.05) and CC (P < 0.01), (Table 2).

**MOLE exhibits an antiapoptotic effect against CCl₄ neurotoxicity**

Caspase 3 gene expression was measured in the HC and CC brain tissues to assess the antiapoptotic effect of MOLE against CCl₄ neurotoxicity. A remarkable increase in caspase 3 gene expression was manifested in both brain tissues in response to CCl₄ toxicity (P < 0.0001). However, MOLE was found to have an antiapoptotic role against this effect as evidenced by the striking decrease in caspase 3 gene expression in both tissues (P < 0.0001), (Table 2).

**MOLE protects against depression-like behaviors induced by CCl₄**
FST and TST were performed to investigate the protective role of MOLE pretreatment against depression-like behaviors manifested by CCl₄-treated mice. Depression-like behaviors represented by mice immobility in seconds after FST and TST were recorded. It was found that CCl₄ markedly increased depression-like behaviors based on both FST (P < 0.01) and TST (P < 0.0001). Nevertheless, pretreatment with MOLE had a marked anxiolytic effect as evidenced by FST (P < 0.01) and TST (P < 0.001), (Fig. 4).

**MOLE pretreatment protects against histopathological changes induced in brain regions by CCl₄**

Coronal sections in the HC and CC were stained with H&E to assess the neuroprotective effect of MOLE against histopathological alterations induced by CCl₄. CCl₄ induced histopathological changes including thinning of the DG region in the HC (Fig. 5) and neuron degeneration (Nd) in the CC (Fig. 6). However, MOLE pretreatment alleviates these changes significantly.

**Discussion**

HE is ascribed with hyperammonemia in the bloodstream that can pass via the blood-brain barrier (BBB) and causes damage to the brain tissue. The mechanism for brain injury associated with HE was reported to be through oxidative stress, inflammatory response, and dysfunction of energy metabolism. It was previously documented that CCl₄ treatment can successfully stimulate liver injury along with HE associated brain tissue damage. For confirmation, the current experiment revealed that serum levels of liver enzymes; ALT and AST, elevated significantly in CCl₄-induced mice when compared with the control group. Pretreatment with MOLE halts the increase of ALT and AST levels in CCl₄-treated mice.

In the present study, CCl₄ induced oxidative stress whereas MOLE pretreatment protected brain tissue from oxidative stress damage. An increase of MDA and a decrease of the level of antioxidants were observed in HC and CC following CCl₄ injection in the current study. Hepatocyte damage is initiated by CYP450- mediated bioactivation of CCl₄ into reactive free radicals; trichloromethyl (CCl₃) and trichloromethyl peroxy radical (CCl₃OO). This activates the release of reactive oxygen species (ROS) which leads to LPO. The severity of LPO of cell membranes can be monitored via assessing the formed MDA in brain tissue. Moreover, the decline of GSH and SOD levels was reportedly due to ROS release in the CCl₄ mouse model of encephalopathy. The dysregulated antioxidant mechanism in the current model was in line with previous reports, in which an elevated MDA and reduced antioxidant mechanism in different brain regions were observed following CCl₄ intoxication. MOLE pretreatment successfully restored the antioxidative power in HC and CC through the reduction of MDA and restoration of antioxidative mechanisms which was in agreement with Idoga et al., (2018). The reversed alterations of antioxidant power in MOLE/CCl₄ treated mice indicate the neuroprotective effect of MOLE and its prominent antioxidant capacity protecting against LPO.

It was recorded that Inflammation is a crucial inducer for HE. Liver damage was associated with peripheral inflammation with cytokine storming that can cross BBB with ascribed neuroinflammation.
disorders \(^{32}\). To evaluate the protective effects of MOLE against neuroinflammation consequences in HC and CC regions of CCl\(_4\) injected mice, relative expressions of TLR2, TLR4, MyD88, and NF-\(\kappa\)B genes as well as the protein levels of TNF-\(\alpha\) and IL-6 were measured.

Toll-like receptors (TLRs) play major roles in inflammatory responses \(^{33}\). TLR2 and TLR4 are considered as neuroinflammatory receptors that are residential in neurons, astrocytes, and microglia \(^{34}\). The adaptor protein of almost all TLRs is MyD88 and acts as a link between the receptors and the downstream signaling components with subsequent activation of transcription and inflammatory responses \(^{34}\). In the current experiment, up-regulation of TLR4, TLR2, and MyD88 gene expressions were detected in CCl\(_4\) treated mice. Our findings were consistent with previous studies \(^{32,35}\), in which upregulated TLR4, TLR2, and MyD88 gene expressions were noticed in HE rat model. MOLE interferes with this cascade as evidenced by the downregulation of TLR4, TLR2, and MyD88 gene expressions manifested by CCl\(_4\)/MOLE treated group.

Gene expression of NF-\(\kappa\)B was also elevated in the present CCl\(_4\) treated mice, which was downregulated by MOLE pretreatment. It has been reported that activating TLR4/2-MyD88 dependent signaling pathway leads to NF-\(\kappa\)B transcription \(^{35}\). Consequently, TLR4/2-MyD88/NF-\(\kappa\)B signaling pathway might be targeted by MOLE in the mitigation of CCl\(_4\) toxicity.

TLR4, a member of TLRs, has been evidenced to play a significant role in initiating the inflammatory response after brain damage \(^{36}\). In the present study, we found an increase in proinflammatory cytokines in the CCl\(_4\) group which was prevented in CCl\(_4\)+MOLE group. CCl\(_4\) toxicity increases the levels of proinflammatory cytokines produced by Kupffer cells. Consequently, liver stromal cells are recruited to assist in intensifying the inflammatory response via the production of cytokines and chemokines \(^{37}\). This peripheral intensification in inflammatory response promotes the activation of microglia and TLR4. Microglial activation is ascribed with the secretion of pro-inflammatory cytokines such as IL-6 and TNF-\(\alpha\) that were associated with the brain deteriorations observed in CCl\(_4\) treated mice \(^{38}\). Whereas, activation of TLR4 exacerbates the inflammatory reactions by inducing NF-\(\kappa\)B pathway leading to the generation of proinflammatory factors \(^{39}\).

It has been established that oxidative stress and neuroinflammation are associated with neurobehavioral changes \(^{40}\). For instance, proinflammatory cytokines such as IL-1\(\beta\), IL-6, and TNF-\(\alpha\) have been demonstrated to be elevated in depression and anxiety, implying immune dysregulation \(^{41}\). Moreover, symptoms of depression were proven to be aggravated by pro-inflammatory cytokines which lead to disturbance of HPA-axis as a result of hypersecretion of CRF with elevated corticosterone level in plasma and subsequent induction of depression symptoms \(^{1,7}\). Subsequently, the neuroprotective effect of MOLE against CCl\(_4\)-induced anxiety and depression-like behavioral changes via FST and TST was assessed. CCl\(_4\)-induced anxiety and depression-like behavioral changes observed in the current study agree with a previous report \(^1\). However, pretreatment with MOLE significantly improved the behavioral status of mice.
This may indicate that MOLE contains anxiolytic and anti-depression phytochemicals. These phytochemicals could be acting on serotonergic, dopaminergic, and/or noradrenergic neurotransmitter systems.  

Besides, the elevated level of corticosterone in CCl₄-treated mice was alleviated by MOLE pretreatment in the current study confirming the previously reported antidepressant effect of MOLE.  

An increased level of ammonia in serum was observed in CCl₄ group in agreement with previous findings. Ammonia exists in biological solutions in two forms; NH₃ and NH₄⁺. CCl₄-induced liver damage results in hyperammonemia represented by increased levels of circulating ammonia. Hyperammonemia has been implicated in neurological disorders through activating brain oxidative stress and neuroinflammation. Hyperammonemia results from liver injury and ammonia can easily pass through BBB causing neurotoxicity.  

In the current study, the antiapoptotic effect of MOLE pretreatment against CCl₄-induced apoptosis was evaluated by assessing relative caspase 3 mRNA level. A decrease in caspase 3 expression in the presence of MOLE confirms the antidepressant effect of MOLE as suggested by Khan et al. (2019). Apoptosis can be triggered via inflammatory response or ROS generated by CCl₄. It can also be explained by the pro-apoptotic effect of corticosteroids on the brain regions especially HC. Consequently, neuroinflammation, oxidative stress, along with apoptosis in the brain cells, can cause anxiety and depression-like behaviors. However, our results proved the anxiolytic and anti-depression properties of MOLE.  

To further explore the mechanisms behind the neuroprotective effect of MOLE against CCl₄-induced neurotoxicity, histopathology of CC and HC was assessed using H&E. The thickness of the DG cellular layer in HC region in CCl₄-challenged mice was found to be markedly thinner than that of the control mice. This agrees with a previous study. Nd was also manifested in the CC of CCl₄-treated mice in agreement with Shaalan et al. (2017). However, MOLE evidently attenuated these alterations. Histopathological results were consistent with the biochemical findings.  

**Conclusions**  
The present study demonstrates, for the first time, the neuroprotective role of MOLE against CCl₄-induced neurotoxicity through biochemical, molecular, behavioral, and histological examinations. MOLE attenuated neuroinflammation, brain oxidative stress, apoptosis, biochemical alterations, and histopathological changes in HC and CC. Furthermore, MOLE significantly improved anxiety and depression-like behaviors. Accordingly, MOLE may be used in the prevention of HE related brain dysfunctions.  

**Declarations**
Ethics approval and consent to participate

Ethical approval was received from Cairo University Institutional Animal Care and Use Committee (CU-IACUC), Egypt, (permission number: CU/I/F/41/20). For consent to participate, it is not applicable.

Consent for publication

Not applicable.

Availability of data and materials

The authors declare that the datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

Competing interests

The authors declare that they have no competing interests.

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Authors’ contributions

SMF and MSMM suggested the research point of the study, designed the experimental protocol, involved in the implementation of the overall study, performed the statistical analyses of the study, researched the data, and wrote the manuscript. Both authors contributed to the critical revision of the manuscript.

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Abbreviations

ALT: alanine aminotransferase; ANOVA: one-way analysis of variance; AST: aspartate aminotransferase; BBB: blood-brain barrier; CCl₃: trichloromethyl; CCl₃OO: trichloromethyl peroxy radical; CCl₄: carbon tetrachloride; CC: cerebral cortex; cDNA: complementary DNA; CRF: corticotrophin-releasing factor; CU-IACUC: Cairo University Institutional Animal Care and Use Committee; DG: dentate gyrus; DMS: degrees minutes seconds; DTNB: 5, 5-dithiobis-2 nitro benzoic acid; ELISA: enzyme-linked immunosorbent assay; FST: forced swimming test; GAPDH: Glyceraldehyde 3-phosphate dehydrogenase; GSH: reduced glutathione; HE: hepatic encephalopathy; HC: hippocampus; H&E: hematoxylin and eosin; HPLC: High-performance liquid chromatography; HPA: hypothalamic-pituitary-adrenal; IL: interleukin; i.p.:
intraperitoneally; JAZUH: Jazan University Herbarium; LPO: lipid peroxidation; MDA: malondialdehyde; MOLE: *Moringa oleifera* Lam. leaf ethanolic extract; MYD88: myeloid differentiation primary response 88; NCI: National Cancer Institute; Nd: neuron degeneration; NF-κB: nuclear factor kappa B; NIH: National Institutes of Health; ROS: reactive oxygen species; RT-qPCR: quantitative reverse transcription-polymerase chain reaction; SOD: superoxide dismutase; TBA: thiobarbituric acid; TLR: toll-like receptor; TNF: tumor necrosis factor; TST: tail suspension test.

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**Table**

**Table 1:** Sequence of primers used for RT-qPCR

| Genes  | Forward primer (5’-3’)                           | Reverse primer (5’-3’)                          |
|--------|--------------------------------------------------|--------------------------------------------------|
| TLR2   | 5′ CAA ATG GAT CAT TGA CAA CAT CATC 3’           | 5′ TTC GTA CTI GCA CCA CTC GC 3’                 |
| TLR4   | 5′ GCTTGAATCCCTGCATAGAGGTAG 3’                   | 5′ TCTTCAAGGGGTGAAGCAGTCAG 3’                   |
| MYD88  | 5′ GGAACAGACCAACTATCGGC 3’                       | 5′ GAGACAACCACCTCAGCCG 3’                       |
| NF-κB  | 5′ AAGGATGTCTCCACACCAGCTG 3’                     | 5′ CACTGTCTGCTCTCGCTGC 3’                       |
| Caspase3| 5′ CCTCAGAGACATTCATGG 3’                         | 5′ GCAGTAGTCGCTCTGAAGA 3’                       |
| GAPDH  | 5′ CATCAACGGAAGGCCATC 3’                         | 5′ CTCGTGGTCACACCCATC 3’                       |

**Table 2:** MOLE pretreatment effect on the relative expression of TLR2, TLR4, MyD88, NF-κB, and caspase 3 as well as the protein levels of TNF-α and IL6 in the hippocampus (HC) and cerebral cortex (CC) of CCl4-injected mice.
### Relative mRNA expression (fold change over control)

|       |       | Control          | CCl₄          | MOLE          | CCl₄+MOLE      |
|-------|-------|------------------|---------------|---------------|---------------|
|       |       |                  | 1.01±0.007071 | 5.000±0.1414  | 1.090±0.007071 | 2.100±0.4950  |
| TLR2  | HC    | 1.01±0.007071    | 5.000±0.1414  | 1.090±0.007071 | 2.100±0.4950  |
|       | CC    | 1.01±0.007071    | 6.000±0.1414  | 0.9800±0.007071 | 2.800±0.4950  |
| TLR4  | HC    | 1.030±0.01414    | 7.000±0.4243  | 1.010±0.02121  | 3.500±0.3465  |
|       | CC    | 1.010±0.01414    | 7.000±0.4243  | 1.020±0.02121  | 3.010±0.3465  |
| MyD88 | HC    | 1.010±0.03512    | 5.000±0.07071 | 1.020±0.01414  | 1.900±0.1131  |
|       | CC    | 1.010±0.02121    | 5.000±0.07071 | 1.000±0.01414  | 2.060±0.1131  |
| NF-κB | HC    | 1.080±0.03512    | 10.00±1.701   | 1.010±0.007071 | 4.010±0.4050  |
|       | CC    | 1.010±0.03512    | 6.000±1.701   | 1.050±0.007071 | 3.200±0.4050  |
| Caspase 3 | HC | 1.000±0.02121    | 6.000±0.2121  | 1.100±0.01414  | 2.900±0.1414  |
|       | CC    | 1.030±0.02121    | 6.000±0.2121  | 1.200±0.02121  | 2.700±0.1414  |

### Protein level (pg/mg)

|       |       | Control          | CCl₄          | MOLE          | CCl₄+MOLE      |
|-------|-------|------------------|---------------|---------------|---------------|
| TNF-α | HC    | 28.60±3.606      | 118.0±6.010   | 26.50±6.000   | 68.40±18.03   |
|       | CC    | 33.70±3.606      | 126.0±6.010   | 35.10±6.000   | 42.90±18.03   |
| IL-6  | HC    | 52.10±8.910      | 141.0±18.17   | 36.10±1.000   | 82.30±12.59   |
|       | CC    | 39.50±8.910      | 167.0±18.17   | 38.40±1.000   | 64.50±12.59   |

TLR: toll-like receptor, MYD88: myeloid differentiation primary response 88, NF-κB: nuclear factor kappa B, TNF-α: tumor necrosis factor- α, IL-6: interleukin 6. All data are represented as mean ± SD, n=8. a indicates P < 0.05, b indicates P < 0.01, c indicates P < 0.001, and d indicates P < 0.0001 compared to control group. w indicates P < 0.05, x indicates P < 0.01, y indicates P < 0.001, and z indicates P < 0.0001 compared to CCl₄ group.

### Figures

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HPLC chromatograms (λ 254 nm) of (A) rutin reference standard (retention time 1.3 min) and (B) ethanolic extract of MOL leaves; HPLC chromatograms (λ 210 nm) of (C) β-sitosterol reference standard (retention time 0.8 min) and (D) ethanolic extract of MOL leaves.

Figure 1

High-performance liquid chromatography (HPLC) analysis of MOLE.
Fig. 2

MOLE pretreatment protects against the increase in the levels of alanine aminotransferase (ALT) (A), aspartate aminotransferase (AST) (B), ammonia (C), and corticosterone (D) induced by CCl₄. All data are represented as mean ± SD, n=8. *P < 0.05, **P < 0.01, ***P < 0.001 and ****P < 0.0001 compared to control group. ##P < 0.01, ###P < 0.001, and ####P < 0.0001 compared to CCl₄ group.

Figure 2

Protective effect of MOLE pretreatment on serum aminotransferases, ammonia, and corticosterone levels.
MOLE pretreatment attenuates elevated LPO, represented by malondialdehyde (MDA), by CCl₄ toxicity in the HC (A) and CC (B). It also restores antioxidant mechanisms, represented by superoxide dismutase (SOD) (C, D) and reduced glutathione (GSH) (E, F) affected by CCl₄ toxicity. All data are represented as mean ± SD, n=8. *P < 0.05, **P < 0.01, and ****P < 0.0001 compared to control group. #P < 0.05, ##P < 0.01, ###P < 0.001, and ####P < 0.0001 compared to CCl₄ group.

Figure 3

MOLE pretreatment impacts lipid peroxidation (LPO) and antioxidant mechanisms in the hippocampus (HC) and cerebral cortex (CC).
MOLE pretreatment protects against the exhibition of depression-like behaviors induced by CCl4 based on forced swimming test (FST) (A) and tail suspension test (TST) (B). All data are represented as mean ± SD, n=8. **P < 0.01, ***P < 0.001, and ****P < 0.0001 compared to control group. ##P < 0.01, ###P < 0.001, and ####P < 0.0001 compared to CCl4 group.

**Figure 4**

MOLE pretreatment influence on the depression-like behaviors induced by CCl4.
Fig. 5

Micrographs (A, B, C, and D) show coronal sections in the HC stained with H&E (40x magnification, scale bar=200µm) of control, CCL₄, MOLE, and CCL₄+MOLE groups, respectively. CA1, CA2, and CA3 denote different regions of cornu ammonis (CA). DG denotes dentate gyrus. Micrograph (E) shows that MOLE pretreatment protects against the reduction in DG thickness induced by CCL₄. All data are represented as mean ± SD, n=8. **P < 0.01 compared to control group.

**Figure 5**

MOLE pretreatment effect on the induced histopathological changes in the hippocampus (HC) by CCL₄ administration.
Micrographs (A, B, C, and D) show coronal sections in the CC (100x magnification, scale bar=50μm) of control, CCl₄, MOLE, and CCl₄+MOLE groups, respectively. Neuron degeneration (Nd) is manifested in the CCl₄ group but MOLE pretreatment protected the neurons against this effect.

**Figure 6**

MOLE pretreatment effect on the induced histopathological changes in the cerebral cortex (CC) by CCl₄ administration.

**Supplementary Files**

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