RESEARCH ARTICLE

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Effects of *Carissa opaca* fruits extracts on oxidative pulmonary damages and fibrosis in rats

Sumaira Sahreen^{1,2*}, Muhammad Rashid Khan² and Rahmat Ali Khan^{2,3*}

Abstract

Background: Carissa opaca is a Pakistani fruit, traditionally used in the treatment of various human ailments including asthma and pulmonary damage. The present study investigated the protective effects of Carissa opaca against CCl_{4} -induced oxidative stress in rat lungs.

Methods: To assess the protective effects of *Carissa opaca*, 42 Sprague–Dawley male rats (170–180 g) were randomly divided into 7 groups. Group I was untreated and group II received olive oil intraperitoneally (i.p.) and dimethyl sulfoxide orally. Groups III, IV, V, VI and VII were administered CCl₄, 3 ml/kg bodyweight (30% in olive oil i.p.). Group IV was administered 50 mg/kg bodyweight silymarin whereas groups V, VI and VII were treated with 200 mg/kg of various fractions of *Carissa opaca* after 48 h of CCl₄ treatment for eight weeks. Antioxidant profiles in lungs were evaluated by estimating the activities of antioxidant enzymes: catalase, peroxidase, superoxide dismutase, glutathione-S-transferase, glutathione reductase, glutathione peroxidase, quinone reductase and reduced glutathione. CCl₄-induced lipid peroxidation was determined by measuring the level of thiobarbituric acid reactive substances (TBARS) with conjugation of DNA damage and histopathology.

Results: Administration of CCl_4 for 8 weeks significantly reduced (p < 0.05) the activities of antioxidant enzymes and GSH concentration while increasing TBARS content and DNA damage. Co-treatment of various fractions of *Carissa opaca* and silymarin restored the activities of antioxidant enzymes and glutathione content. Changes in TBARS concentration and DNA fragmentation was significantly decreased (p < 0.05) following *Carissa opaca* and silymarin treatment in lung.

Conclusions: Histopathological changes in rat lungs induced by CCl₄ were significantly restored by co-treatment with *Carissa opaca* and silymarin.

Keyword: Carissa opaca, Lungs, CCl4, Antioxidant enzymes, TBARS, DNA

Background

Exposure to UV radiation, X-rays, environmental pollutants, toxic chemicals and excessive drug use causes the production of free radicals and reactive oxygen species that lead to oxidative damage in the kidney, liver and lungs [1]. The lung is the main organ of respiration and is exposed to higher oxygen stress compared with other tissues. Exogenous oxidants, cigarette smoke and asbestos fibers activate inflammatory cells to generate oxidative stress and lung fibrosis [1]. Several important reactive oxygen species (ROS) are generated endogenously in these circumstances [2], which include superoxide radicals, hydrogen peroxide (H_2O_2), and hydroxyl radicals. The major enzymes/reaction pathways that are activated to generate ROS in human lungs include nicotinamide adenine dinucleotide phosphate oxidases, myeloperoxidase, eosinophil peroxidase, mitochondrial electron transport chain, and possibly xanthine oxidase [3]. A balance between intracellular and extracellular oxidants and antioxidants is a prerequisite for normal lung homeostasis. The lung has highly specialized and compartmentalized antioxidant defenses to protect against ROS and reactive nitrogen species. Induction of these antioxidant enzymes and related proteins after pulmonary insults may protect the lung and promote normal repair. Conversely, impaired



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induction or inactivation/clearance of antioxidant enzymes may result in a sustained redox imbalance that may contribute to the progression of pulmonary fibrosis [4].

Plants are commonly used for the discovery of new therapeutic products. In recent years, there has been increasing interest in finding natural antioxidants from commonly available wild plants, fruits and vegetables [5] as well as their important role in detoxification of free radical-induced lung injuries and fibrosis in experimental animal models [6]. *Carissa opaca* (*C. opaca*) Stapf ex Hanes is a 2–3 meter tall evergreen shrub containing glabrous fruits widely found in Pakistan [7]. Traditionally this plant is used for the treatment of asthma [8], hepatitis [9], diarrhea [10] and renal dysfunction [11]. The present study examined the toxic effects of CCl_4 and investigated the beneficial effects of plant extracts on tissues from various CCl_4 -induced lung damage experimental groups.

Methods and Materials

Plant collection

C. opaca ripened fruits were collected in March-April 2011 from the Quaid-i-Azam University Islamabad, Pakistan. The plants were recognized by their local names and then validated by Dr. Mir Ajab Khan, Department of Plant Sciences, Quaid-i-Azam University, Islamabad. A voucher specimen with Accession No. 24561 (*C. opaca*) was deposited at the Herbarium of Pakistan Quaid-i-Azam University, Islamabad Pakistan.

Extract preparation

The collected fruits were cleaned to get rid of dust particles and then dried under shade for one to two weeks. Willy Mill of 60-mesh size was used to prepare powder of dried samples. 5 kg of powdered sample was extracted twice with 10 L of 95% methanol at 25°C for 48 h. For filtration Whatman No. 1 filter paper was used and then filtrate was concentrated through rotary evaporator (Panchun Scientific Co., Kaohsiung, Taiwan) under reduced pressure at 40°C. In order to resolve the compounds with escalating polarity, a part of the extract was suspended in distilled water and subjected to liquid-liquid partition by using solvents in a sequence of n-hexane and ethyl acetate. After fractioning, the solvent of respective fractions was also evaporated by rotary evaporator. Extract was dried and then stored at 4°C for further *in vivo* investigation.

Experimental plan

Six-week-old male Sprague Dawley rats weighing 180 ± 10 g were provided with food and water *ad libitum* and kept at 20-22°C on a 12-h light/dark cycle. All experimental procedures involving animals were conducted in accordance with the guidelines of National Institutes of Health (NIH guidelines). The study protocol were approved by Ethical committee of Quaid-i-Azam University

Islamabad. The rats were acclimatized to laboratory condition for 7 days before commencement of experiment. For chronic toxicity eight week experiment was designed. 42 male albino rats were randomly divided into seven groups (6 rats of each group). Administration of CCl₄ (0.5 ml/kg b.w., 20% CCl₄/olive oil) was intraperitoneally (i.p.) twice a week for eight weeks. At the same time, the rats were administered individually silymarin (50 mg/kg b.w.) and extract (200 mg/kg b.w.) orally twice a week for eight weeks.

Experimental protocol

Following dosing plan was adapted for the study.

Group I:
Normal control received only feed
Group II:
Olive oil (0.5 ml/kg b.w., i.p.) + DMSO (0.5 ml/kg
b.w. orally)
Group III:
CCl_4 twice a week (0.5 ml/kg b.w., i.p., 20% CCl_4 /
olive oil)
Group IV:
CCl_4 twice a week (0.5 ml/kg b.w., i.p.) + sylimarin
(50 mg/kg b.w., orally)
Group V:
CCl_4 twice a week (0.5 ml/kg b.w., i.p.) + n-heaxne
Carissa opaca fruit extract (HFC, 200 mg/kg
b.w., orally)
Group VI:
CCl_4 twice a week (0.5 ml/kg b.w., i.p.) + Ethyl
acetate Carissa opaca fruit extract (EFC 200 mg/kg
b.w., orally)
Group VII:
CCl_4 twice a week (0.5 ml/kg b.w., i.p.) + Methanol
Carissa opaca fruit extract (MFC 200 mg/kg
b.w., orally)

At the end of eight weeks, after 24 h of the last treatment, animals were given chloroform anesthesia and dissected from ventral side. All the animals were sacrificed; lungs were removed and washed in ice cold saline. Subsequently, half of the organs were treated with liquid nitrogen and stored at -80°C for further enzymatic and DNA damage analysis while the other portion was processed for histology.

Biochemical investigations

In order to evaluate the pharmacological effects of different fractions of *C. opaca* against the toxicity induced with CCl_4 in rats following assays had been carried out.

Assessment of antioxidant enzymes

10% tissue homogenate was prepared in 100 mM $\rm KH_2PO_4$ buffer containing 1 mM EDTA (pH 7.4) and centrifuged

at 12,000 \times g for 30 min at 4°C. The supernatant was collected and used for the following parameters as described below.

Catalase assay (CAT)

CAT activities were determined by the method of Chance and Maehly [12] with some modifications. The reaction solution of CAT activities contained: 2.5 ml of 50 mM phosphate buffer (pH 5.0), 0.4 ml of 5.9 mM H_2O_2 and 0.1 ml enzyme extract. Changes in absorbance of the reaction solution at 240 nm were determined after one minute. One unit of CAT activity was defined as an absorbance change of 0.01 as units/min.

Peroxidase assay (POD)

Activities of POD were determined by the method of Chance and Maehly [12] with some modifications. The POD reaction solution contained: 2.5 ml of 50 mM phosphate buffer (pH 5.0), 0.1 ml of 20 mM guaiacol, 0.3 ml of 40 mM H_2O_2 and 0.1 ml enzyme extract. Changes in absorbance of the reaction solution at 470 nm were determined after one minute. One unit of POD activity was defined as an absorbance change of 0.01 units/min.

Superoxide dismutase assay (SOD)

SOD activity was estimated by the method of Kakkar *et al.* [13]. Reaction mixture of this method contained: 0.1 ml of phenazine methosulphate (186 μ M), 1.2 ml of sodium pyrophosphate buffer (0.052 mM; pH 7.0), 0.3 ml of supernatant after centrifugation (1500 × g for 10 min followed by 10000 × g for 15 min) of lung homogenate was added to the reaction mixture. Enzyme reaction was initiated by adding 0.2 ml of NADH (780 μ M) and stopped after 1 min by adding 1 ml of glacial acetic acid. Amount of chromogen formed was measured by recording color intensity at 560 nm. Results are expressed in units/mg protein.

Glutathione-S-transferase assay (GST)

Glutathione-S-transferase activity was assayed by the method of Habig *et al.* [14]. The reaction mixture consisted of 1.475 ml phosphate buffer (0.1 mol, pH 6.5), 0.2 ml reduced glutathione (1 mM), 0.025 ml (CDNB) (1 mM) and 0.3 ml of homogenate in a total volume of 2.0 ml. The changes in the absorbance were recorded at 340 nm and enzymes activity was calculated as nM CDNB conjugate formed/min/mg protein using a molar extinction coefficient of $9.6 \times 10^3 \text{ M}^{-1} \text{ cm}^{-1}$.

Glutathione reductase assay (GR)

Glutathione reductase activity was determined by method of Carlberg and Mannervik [15]. The reaction mixture consisted of 1.65 ml phosphate buffer: (0.1 mol; pH 7.6), 0.1 ml EDTA (0.5 mM), 0.05 ml oxidized glutathione (1 mM), 0.1 ml NADPH (0.1 mmol) and 0.1 ml of homogenate

in a total volume of 2 ml. Enzyme activity was quantitated at 25 °C by measuring disappearance of NADPH at 340 nm and was calculated as nM NADPH oxidized/ min/mg protein using molar extinction coefficient of 6.22×10^3 M⁻¹ cm⁻¹.

Glutathione peroxidase assay (GPx)

Glutathione peroxidase activity was assayed by the method of Mohandas *et al.* [16]. The reaction mixture consisted of 1.49 ml phosphate buffer (0.1 M; pH 7.4), 0.1 ml EDTA (1 mM), 0.1 ml sodium azide (1 mM), 0.05 ml glutathione reductase (1 IU/ml), 0.05 ml GSH (1 mM), 0.1 ml NADPH (0.2 mM), 0.01 ml H₂O₂ (0.25 mM) and 0.1 ml of homogenate in a total volume of 2 ml. The disappearance of NADPH at 340 nm was recorded at 25 °C. Enzyme activity was calculated as nM NADPH oxidized/min/mg protein using molar extinction coefficient of $6.22 \times 10^3 \text{ M}^{-1} \text{ cm}^{-1}$.

Quinone reductase assay (QR)

The activity of quinone reductase was determined by the method of Benson *et al.* [17]. The 3.0 ml reaction mixture consisted of 2.13 ml Tris-HCl buffer (25 mM; pH 7.4), 0.7 ml BSA, 0.1 ml FAD, 0.02 ml NADPH (0.1 mM), and 0.1 ml of homogenate. The reduction of dichlorophenolindophenol (DCPIP) was recorded at 600 nm and enzyme activity was calculated as nM of DCPIP reduced/min/mg protein using molar extinction coefficient of 2.1×10^4 M⁻¹ cm⁻¹

Reduced glutathione assay (GSH)

Reduced glutathione was estimated by the method of Jollow *et al.* [18]. 1.0 ml sample of homogenate was precipitated with 1.0 ml of (4%) sulfosalicylic acid. The samples were kept at 4°C for 1 h and then centrifuged at 1200 × g for 20 min at 4°C. The total volume of 3.0 ml assay mixture contained 0.1 ml filtered aliquot, 2.7 ml phosphate buffer (0.1 M; pH 7.4) and 0.2 ml DTNB (100 mM). The yellow color developed was read immediately at 412 nm on a SmartSpecTM plus Spectrophotometer. It was expressed as μ M GSH/g tissue.

Estimation of lipid peroxidation assay (TBARS/LPO)

The assay for lipid peroxidation was carried out following the modified method of Iqbal *et al.* [19]. The reaction mixture in a total volume of 1.0 ml contained 0.58 ml phosphate buffer (0.1 M; pH 7.4), 0.2 ml homogenate sample, 0.2 ml ascorbic acid (100 mM), and 0.02 ml ferric chloride (100 mM). The reaction mixture was incubated at 37°C in a shaking water bath for 1 h. The reaction was stopped by addition of 1.0 ml 10% trichloroacetic acid. Following addition of 1.0 ml 0.67% thiobarbituric acid, all the tubes were placed in boiling water bath for 20 min and then shifted to crushed ice-bath before centrifuging at 2500 × g for 10 min. The amount of TBARS formed in each of the samples was assessed by measuring optical density of the supernatant at 535 nm using spectrophotometer against a reagent blank. The results were expressed as nM TBARS/min/mg tissue at 37°C using molar extinction coefficient of 1.56×10^5 M⁻¹ cm⁻¹.

Hydrogen peroxide assay (H₂O₂)

Hydrogen peroxide (H_2O_2) was assayed by H_2O_2 -mediated horseradish peroxidase-dependent oxidation of phenol red by the method of Pick and Keisari [20]. 2.0 ml of homogenate sample was suspended in 1.0 ml of solution containing phenol red (0.28 nM), horse radish peroxidase (8.5 units), dextrose (5.5 nM) and phosphate buffer (0.05 M; pH 7.0) and were incubated at 37°C for 60 min. The reaction was stopped by the addition of 0.01 ml of NaOH (10 N) and then centrifuged at 800 × g for 5 min. The absorbance of the supernatant was recorded at 610 nm against a reagent blank. The quantity of H_2O_2 produced was expressed as nM $H_2O_2/min/mg$ tissue based on the standard curve of H_2O_2 oxidized phenol red.

Molecular studies

DNA had been isolated and its fragmentation percent was quantified in molecular studies of *in vivo* toxicity.

DNA fragmentation assay with diphenylamine reaction

DNA fragmentation from tissue extract was determined using the procedure of Wu *et al.* [21]. 100 mg tissue was homogenized in TTE solution. 0.1 ml of homogenate was labeled B, centrifuged at 200 × g at 4°C for 10 min, got supernatant labeled S. S tubes were centrifuged at 20,000 × g for 10 min at 4°C to separate intact chromatin, was labeled T. 1.0 ml of 25% TCA was added in all tubes T, B, S and incubated over night at 4°C. After incubation precipitated DNA was recovered by pelleting for 10 min at 18,000 × g at 4°C. 160 µl of 5% TCA was added to each pellet and heated for 15 min at 90°C then 320 µl of freshly prepared DPA solution was added, vortexed and incubated for 4 hr 37°C. Optical density was read at 600 nm with a spectrophotometer (Smart spec[™] Plus, catalog # 170-2525).

DNA Isolations and ladder assay

DNA was isolated by using the methods of Wu *et al.* [21]. 100 mg of tissue in a petri dish was washed with DNA Buffer and homogenized in 1 ml lysis buffer. 100 μ l of proteinase K (10 mg/ml) and 240 μ l 10% SDS, shaked gently, and incubate overnight at 45°C in a water bath then 0.4 ml of phenol, was added shaked for 5-10 min, and centrifuge at 3000 rpm for 5 min at 10°C. Supernatant was mixed with 1.2 ml phenol, 1.2 ml chloroform/isoamyl alcohol (24:1); shaked for 5-10 min, and centrifuged at 3000 rpm for 5 min at 10°C. 25 μ l of 3 M sodium acetate (pH 5.2) and 5 ml ethanol was added with supernatant,

shake until DNA was precipitated. DNA was washed with 70% ethanol, dried, dissolved in TE buffer and its concentration checked at 260 and 280 nm.5 μ g of total DNA and 0.5 μ g DNA standard per well were loaded on 1.5% agarose gel containing ethidium bromide. Electrophoresis was performed for 45 min with 100 V batteries, and DNA was observed under digital gel doc system and photographed.

Histopathological study of tissue

After weighting the portion specifies for histology small pieces of each tissue was fixed for 3-4 h in fixative sera followed by dehydration with ascending grades of alcohol (80%, 90%, and 100%) and transferred in cedar wood oil. When tissue becomes clear then all tissues were embedded in paraplast and prepared blocks for further microtomy. 3-4 μ m thin slides were prepared with microtome; wax was removed, stained with hemotoxilin-eosin and photographed under light microscope at 10× and 40×.

Statistical analysis

To find the different treatment effects, one way analysis of variance was carried by computer software SPSS 13.0. Level of significance among the various treatments was determined by LSD at 0.05% level of probability.

Results

Effects of *C. opaca* fruit against CCl₄ induced pulmonary toxicity in rat

 CCl_4 is a powerful pulmonary toxin that induces acute and chronic lung toxicity. This study induced pulmonary toxicity by CCl_4 and determined the therapeutic effect of *C. opaca* fruit in CCl_4 administered rats. Parameters studied included antioxidant enzymatic levels, genotoxicity and characteristic histological findings of lungs.

Effects of C. opaca fruit on enzymatic antioxidant levels

Oxidative stress can affect almost every organ including the lungs. ROS production is caused by oxidative stress that directly damages lipids, proteins and DNA. In reaction to this, the body has its own defense system that consists of antioxidant enzymes that limit the level of damage. The protective effects of different fractions of *C. opaca* fruit against CCl₄-induced pulmonary changes of tissue proteins and antioxidant levels is shown in Table 1. CCl₄ treatment considerably (p < 0.05) decreased tissue protein, catalase, peroxidase, and superoxide dismutase levels compared with the control group. Co-administration of EFC, HFC and MFC significantly (p < 0.05) increased antioxidant enzyme activity.

Effects of C. opaca fruit on TBARS and H₂O₂

The effects of *C. opaca* fruit on TBARS and H_2O_2 levels are shown in Table 2. Increased TBARS and H_2O_2 levels were due to CCl_4 toxicity. Post-administration of various

1.12 ± 0.02a

 $2.59 \pm 0.10d$

 $1.46 \pm 0.10b$

1.51 ± 0.08b

 $2.20 \pm 0.13c$

	-	-	-	
Group	Protein (µg/mg tissue)	CAT (U/min)	POD (U/min)	SOD (U/mg protein)
Control	2.01 ± 0.03f	4.11 ± 0.21d	$10.42 \pm 0.53d$	3.40 ± 0.23e
Oil + DMSO	$1.96 \pm 0.03 f$	$4.09 \pm 0.30d$	$11.00 \pm 0.20d$	3.25 ± 0.21e

 $2.09 \pm 0.24a$

 $3.60 \pm 0.11c$

 $2.60 \pm 0.16b$

 $3.02 \pm 0.07c$

3.30 ± 0.10c

Table 1 Effects of various fractions of C. opaca fruit on tissue proteins and antioxidant enzyme levels

Values are Mean ± SD (06 number). Sily = Silymarin.

CCl₄

 $Sily + CCl_4$

HFC + CCl₄

EFC + CCl₄

MFC + CCl₄

a-f (Means with different letters) indicate significance at p < 0.05.

0.74 ± 0.01a

 $1.49 \pm 0.02e$

0.89 ± 0.03b

1.05 ± 0.06c

 $1.35 \pm 0.078d$

fractions of *C. opaca* fruits appreciably (p < 0.05) ameliorated the toxic effects of CCl₄ in lung tissues in contrast to the control group. Similar ameliorating effects were observed with silymarin administration, which had levels similar to the control group.

Effects of C. opaca fruit on glutathione enzymes

Phase II antioxidant metabolizing enzymes play a key role in the detoxification of oxidative stress. CCl₄ insults significantly (p < 0.05) reduced the activities of glutathione enzymes such as glutathione S-transferase (GST), glutathione peroxidase (GPx) and glutathione reductase (GR) when compared with the control group (Table 3). Various fractions of *C. opaca* fruits significantly (p < 0.05) ameliorated toxicity by increasing the activity of phase II antioxidant enzymes towards control group levels. Silymarin administration to rats significantly (p < 0.05) reduced toxicity and restored the activities of GST, GPx and GR in lung tissues.

Effects of *C. opaca* fruit on GSH, QR and DNA fragmentation Glutathiones and DNA play important roles in free radicalinduced detoxification. Administration of CCl_4 significantly reduced (p < 0.05) both GSH and QR content, and significantly (p < 0.05) increased DNA fragmentation (Table 4).

Table	2 Effects	of various	fractions	of C.	opaca fruit or	Ì
tissue	e TBARS a	nd H ₂ O ₂				

Group	TBARS (nM/min/mg protein)	H ₂ O ₂ (nM/min/mg tissue)
Control	3.13 ± 0.18c	1.21 ± 0.074f
Oil + DMSO	3.20 ± 0.13c	$1.17 \pm 0.051 f$
CCl ₄	5.41 ± 0.68a	$2.64 \pm 0.062a$
$Sily + CCl_4$	$4.02 \pm 0.47 b$	1.53 ± 0.058e
$HFC + CCI_4$	4.91 ± 0.11a	2.24 ± 0.035b
$EFC + CCI_4$	4.89 ± 0.15a	2.21 ± 0.043b
$MFC + CCI_4$	4.07 ± 0.23b	1.99 ± 0.094c

Values are Mean ± SD (06 number). Sily = Silymarin.

a-f (Means with different letters) indicate significance at p < 0.05.

Treatment of rats with various fractions of *C. opaca* significantly (p < 0.05) restored DNA fragmentation and GSH and QR activity compared with the CCl₄ group. Silymarin also showed significant protection against CCl₄.

5.74 ± 0.42a

 $9.64 \pm 0.23c$

6.53 ± 0.32b

6.89 ± 0.17b

7.80 ± 0.17b

Effects of C. opaca fruit on DNA damages (ladder assay)

 CCl_4 is a genotoxic chemical and causes DNA damages. The protective effects of various fractions of *C. opaca* fruit against CCl_4 induced DNA damages in rats are shown by DNA ladder assay in Figure 1. Intact genomic DNA was revealed by ladder assay of control group while, CCl_4 group showed marked DNA damages. Co-treatment of silymarin and various fractions viz; HFC, EFC and MFC proved the recovering effects by DNA band pattern showing similarity with control group.

Effects of C. opaca fruit on lung histoarchitecture

The effects of various fractions of C. opaca fruits against CCl₄ induced lung injury and histological changes were observed. Histological assessments of lung tissues of control and DMSO group confirmed the typical cellular architecture with distinct alveolar septa and bronchioles, structured Clara cells and fibroblasts as shown in Figure 2A and B, respectively. Aggregation of fibroblasts, collagen fibres, ruptured alveolar bronchioles and walls, disorganized Clara cells showing pulmonary edema and interstitial hemorrhage were found in CCl₄ intoxicated rats (Figure 2C). The lung sections of rats treated with various fractions of C. opaca fruits viz; EFC and MFC intoxicated with CCl₄ (Figure 2E-F), showed normal structure of alveolar bronchioles and less degenerative changes with various degrees in case of each group, supplementing the protective effects of the plant samples. Post-administrations of silymarin reduced the toxic effects of CCl₄ and reversed the histopathology towards the control group (Figure 2D). Present histological observations are in agreement with the results of pulmonary oxidative stress level.

Discussion

Lungs are the site of respiration and are exposed to the action of toxic chemicals, drugs or smoking. Toxin inhalation

Group	GST (nM/min/mg protein)	GPx (nM/min/mgprotein)	GR (nM/min/mg protein)
Control	170.29±4.28 h	114.21 ± 3.29 g	247.01 ± 6.00 g
Oil + DMSO	177.43 ± 4.31 h	108.48 ± 3.12 g	236.23 ± 5.73 g
CCl ₄	92.13 ± 3.48a	65.08 ± 3.32a	136.24 ± 3.72a
$Sily + CCl_4$	161.34 ± 3.22 g	101.48 ± 2.81f	192.41 ± 4.07e
$HFC + CCI_4$	99.61 ± 2.03b	73.40 ± 1.45b	149.32 ± 4.56b
$EFC + CCI_4$	105.34 ± 2.51c	72.02 ± 1.81b	154.66 ± 4.39b
MFC + CCl ₄	130.80 ± 3.73e	83.52 ± 2.40d	177.46 ± 4.29d

Table 3 Effects of various fractions of C. opaca fruit on GST, GPx and GSH

Values are Mean \pm SD (06 number). Sily = Silymarin.

a-f (Means with different letters) indicate significance at p < 0.05.

is often injurious to health and can cause pulmonary disease. Lungs are often injured by oxidative stress. A biochemical study suggested CCl₄ free radicals could have the same consequences [2]. Pulmonary toxicity causes notable pathological effects such as fibrosis, inflammatory changes and degeneration of epithelial cells. Symptoms of pulmonary toxicity include fibrosis, inflammatory response, and degeneration of epithelial cells. Therefore, this study investigated the pharmacological outcomes of different plant extracts and their fractions on CCl₄-induced oxidative damage in rat lungs [22]. The potency of natural antioxidants depends on their chain breaking capacity, declining hydrogen peroxide levels, scavenging superoxides and chelating transitory metal ions [23]. Food scientists and nutrition specialists suggest that plants are useful as a source of natural antioxidants that contribute to reducing risks of certain diseases, such as cancer and cardiovascular disease. In this context, intake of plants might prevent the onset of these diseases [24]. Food deterioration is brought about by lipid peroxidation. The use of synthetic drugs as preservative agents to inhibit lipid peroxidation are strictly prohibited as they are potent carcinogens [25]. Consequently, these chemical therapeutics/synthetic drugs should be replaced with naturally occurring agents having no or very few side-effects. A previous study verified that a diet high in vegetables and fruits is associated with a decline in

Table 4 Effects of various fractions of C. opaca fruit onGSH, QR and DNA fragmentation

Group	GSH (µM/g tissue)	QR (nM/min/mg protein)	DNA damages %
Control	27.07 ± 1.41e	119.18±3.03 g	12.82 ± 1.38e
Oil + DMSO	24.23 ± 1.50e	124.27 ± 3.68 g	13.53 ± 1.98e
CCI_4	10.23 ± 0.65a	60.74 ± 2.13a	61.42 ± 2.70a
$Sily + CCl_4$	20.53 ± 1.41d	104.47 ± 3.22f	19.19 ± 1.48d
$HFC + CCI_4$	11.65 ± 0.46b	66.18 ± 2.87b	33.26 ± 0.72b
$EFC + CCI_4$	11.90 ± 0.57b	73.30 ± 3.17c	33.47 ± 0.48b
$MFC + CCI_4$	15.79 ± 1.00c	91.26 ± 2.32e	20.33 ± 1.54d

Values are Mean \pm SD (06 number). Sily = Silymarin.

a-h (Means with different letters) indicate significance at p < 0.05.

degenerative diseases [26], thus natural antioxidants, flavonoids and phenolic compounds have gained considerable attention. Structurally, phenolic compounds contain a conjugated ring with a hydroxyl group, and can act as an antioxidant by preventing free radical-mediated diseases. Plant antioxidant capacity is characterized by quenching free radicals such as lipid peroxy radicals, singlet oxygen and superoxide anions [27]. Combinatorial methods are required to explore the antioxidant tendency of natural resources because no single assay can reflect all antioxidants in a mixed assay or the complex nature of phytochemicals. Adedapo et al. [28] reported the scavenging abilities of plant extracts against free radicals in a complex assay system to eradicate the radical-related pathological diseases. Toxic compounds of drugs and xenobiotics are metabolized by the glutathione system (reduced glutathione, glutathione reductase, glutathione peroxidase and glutathione-S-transferase). Administration of C. opaca reduced CCl₄ toxicity, thereby increasing the activity of GST, GSR, GSH-Px and QR [29,30]. Similar observations were reported by Khan et al. [31], when administering melatonin against CCl₄-induced oxidative stress. Free radicals cause lipid peroxidation, elevate TBARS and deplete tissue GSH contents [32]. In the present study, low levels of GSH were accompanied by elevated levels of TBARS and H₂O₂ compared with the control group. Fruit extracts of C. opaca were characterized by the high expression level of GSH contents with low level of TBARS and H₂O₂. Similar observations were reported during co-treatment of plant extracts against CCl₄-induced damage in rats [33]. Lipid peroxidation induced by CCl₄ disturbs protein synthesis but can also diffuse into the nucleus, causing DNA fragmentation [34,35] that can lead to pulmonary damages. In the present study, CCl₄-induced DNA damage was significantly ameliorated by C. opaca as reported previously by Khan et al. [36]. Extensive variations were observed during histopathological study of rat lungs. CCl₄ damage of the alveolar septa and mobbing of blood capillaries resulted in the accumulation of blood cells and collagen fibers at various places causing an endemic condition. Similar observations were found in rat lungs in previous studies during CCl₄

Sahreen et al. BMC Complementary and Alternative Medicine 2014, **14**:40 http://www.biomedcentral.com/1472-6882/14/40





Conclusion

The present results revealed that *C. opaca* comprised of bioactive compounds; presenting protective effects against CCl_4 induced toxic effects in lungs of rat. Further studies of isolation and purification of these constituents are in progress in our lab.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

SS made significant contribution to acquisition of data, analysis, conception, design of the manuscript. MRK and RAK (ORCID ID: 0000-0003-0453-2090) made significant contribution to acquisition of data, analysis, drafting and conception. All the authors read and approved the final manuscript.

Acknowledgements

We are very thankful to Higher Education Commission (HEC) Pakistan for provision research funds.

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Received: 15 February 2013 Accepted: 28 January 2014 Published: 30 January 2014

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doi:10.1186/1472-6882-14-40

Cite this article as: Sahreen *et al.*: **Effects of** *Carissa opaca* **fruits extracts on oxidative pulmonary damages and fibrosis in rats**. *BMC Complementary and Alternative Medicine* 2014 **14**:40.

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