Investigation of the effect of watercress via nuclear factor-E2-related factor-2 pathway on 7,12-dimethylbenz[a] anthracene-induced oxidative damage in rat liver

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ABSTRACT

Aim: Nuclear factor E2-related factor-2 (Nrf2) regulates many cytoprotective antioxidants, and detoxifying enzymes, has been a hopeful approach for chemoprevention of cancer. We researched the chemopreventive impact of watercress on the DMBA-induced oxidative damage in the rat liver via the Nrf2/ARE signalling pathway in the present study.

Material and Method: Female Wistar albino rats have categorized into three groups. The control was the first group, rats in the second and third groups were administered 7,12-dimethylbenz[a]anthracene (DMBA) (20 mg/kg b.w., i.p.) and rats in the third group were given daily doses of watercress (250 mg/kg b.w. Nasturtium officinale R.Br. for 4 weeks by oral gavage). Antioxidant and phase II detoxification enzyme activities, Nrf2 transcription factor level in liver, and serum ALT were determined. Also, histopathological analysis of the liver was performed.

Results: We observed that watercress induces DNA-binding of Nrf2. It was related to increased enzyme activities of phase II detoxifying and the antioxidant. Our results also demonstrated that watercress ameliorated liver injury.

Conclusion: Our data ensured considerable evidence that the dietary watercress ameliorates DMBA induced liver toxicity via regulating the activation of the Nrf2/ARE pathway which increases the expression of cytoprotective enzymes.

Keywords: Watercress, Nrf2, DMBA, xenobiotic enzymes, hepatoprotective effect

INTRODUCTION

Cancerincidence of the world and its mortality is associated with lifestyle and environmental agents like food and chemicals. Sustained exposure to chemicals directly or indirectly leads to increase cancer risk by participating in all aspects of carcinogenesis (1). Polycyclic aromatic hydrocarbons (PAHs) are xenobiotic compounds that are environmental contaminants. PAHs are lipophilic and metabolized by xenobiotic-metabolizing enzymes (XMEs) (2). Firstly, enzymes of cytochrome P 450 family 1 that includes CYP1A1, CYP1A2, and CYP1B1 oxidize lipophilic compounds and generates reactive metabolites that interact with protein and DNA. Secondly, highly reactive compounds are converted to more hydrophilic species and/or inactivated by enzymes of phase II detoxification such as glutathione-S-transferase (GST), UDP -glucuronosyltransferases, heme oxygenases-1

(HO-1), NAD(P)H: quinine oxidoreductase 1 (NQO1), and sulfotransferases (3).

DMBA is one of the PAHs that is mainly found in automobile exhaust, cigarette smoke, grilled foods (4) which leads to the generation of reactive oxygen species (ROS), causing oxidative damage and decreased antioxidant defense enzymes, and increases carcinogenicity and toxicity (5). DMBA causes liver toxicity through the released carcinogenic metabolites as a result of xenobiotic metabolism (6). Normally Nrf2, is sensitive to redox changes, exists in an inactive form with a cytosolic inhibitor called Kelch-like epichlorohydrinassociated protein 1 (Keap1). Oxidative stress generating factors including electrophiles, xenobiotics, heavy metals, UV radiation, and physiological stress lead to cleavage of Nrf2 from the Nrf2-Keap1 complex. Nrf2 translocates to



the nucleus and binds to the antioxidant response element (ARE). This causes the transcription of genes of phase II enzymes including NQO1 and GST (7). Excessive ROS generation due to the xenobiotic metabolism of DMBA causes oxidative stress, liver damage, and carcinogenesis. The Nrf2/ARE binding results in the induction of detoxification enzymes, antioxidative stress proteins, and other defense mechanisms (8).

Watercress belongs to the Cruciferae, also called Brassicaceae, family and is one of the perennial freshwater macrohydrophytes that has ecological and economic importance. It has been used by humans for many years both for food and medical purposes (9). This plant has some medicinal properties such as a diuretic (10), antihypertensive (11), antidiabetic (12), antihyperlipidemic (13), antiasthmatic, antituberculosis (14), and anticarcinogenic (15). Furthermore, it contains vitamins (Vitamin A and C), phenolic compounds, minerals, and glucosinolates (16). Glucosinolates are converted to isothiocyanates (ITC) non-enzymatically by physical factors or enzymatically by myrosinase during food preparation, cooking, and chewing (17). Phenethylisothiocyanate (PEITC) in watercress, is an ITC, has been extensively investigated as the most promising chemopreventive compound (18).

In this study, we hypothesized that watercress-mediated Nrf2 activation is a novel hepatoprotective pathway. The goal of this study was to evaluate the chemopreventive effect of watercress grown in the Malatya region of Turkey on oxidative liver injury induced by DMBA via the Nrf2/ARE pathway.

MATERIAL AND METHOD

Animals and Experimental Design

All experimental procedures of animals were approved by the Local Ethics Committee of the Animal Experiments of İnönü University (Date: 20/02/2015, Decision No: 2015/A-25). Twenty-seven female Wistar albino rats (weighing 150-250 g) accommodated in a room kept at 22°C with a 12-hr light/dark cycle and had free access to food. Rats were separated randomly into three groups (9 rats per group): Rats in group I served as control and were injected intraperitoneally with corn oil. Rats in group II (DMBA) and group III (DMBA+watercress) were injected intraperitoneally with DMBA (20 mg/kg body-weight in corn oil) (Sigma-Aldrich, USA) (19). Rats in group III were given daily doses of watercress (Nasturtium officinale R.Br) (250 mg/kg b.w) for 4 weeks by oral gavage before intraperitoneal injection of DMBA (20 mg/kg bodyweight in corn oil). After 30 days, under xylazine/ketamine anesthesia blood of rats in all groups was collected and then liver tissues of the rats were resected, washed in phosphate buffer, and rapidly frozen at -80°C.

Plant Material and Extraction

Nasturtium officinale R.Br were obtained from the pond in Arguvan country, city of Malatya, Turkey. A voucher specimen (No: 1001) was deposited into a herbarium at the Faculty of Pharmacy of İnönü University in Malatya, Turkey (**Figure 1**). Samples were freeze-dried and were powdered. The powdered sample was extracted with 70% methanol by heating at 70°C for 30 minutes and filtered. This procedure was carried out two more times and then centrifuged at 4000 rpm for 10 min. The solvent was removed with a rotary evaporator (Heidolph Laborota 4011-digital) at 90 rpm, and 45°C for 2 h and dried.



Figure 1. The aerial parts of Nasturtium officinale R.Br. were collected from the pond in Arguvan country, city of Malatya, Turkey (Fresh plant and a voucher specimen (No: 1001)).

Determination of PEITC Content of the Watercress

The PEITC content of the watercress was determined by HPLC. PEITC standards were prepared with acetonitrile at concentrations of 0.312, 0.625, 1.25, 2.5, 5, and 10 ppm. 100 μL of the cyclocondensation working reagent (10 mM 1,2 benzene dithiol isopropanol and pH 8.5 phosphate buffer) was mixed and incubated at 65°C for 2 h. For the determination of PEITC amount is needed the conversion of glucosinolates to isothiocyanates and derivatization of the isothiocyanates by a cyclocondensation reaction. 2 mg of dry watercress powder was solved in 2 mL deionized water and added 1 mL myrosinase. It was incubated at 37°C for 2 h, and centrifugated at 875 g for 15 min. It was derivatized by cyclocondensation reaction before HPLC analysis (20). For the chromatographic separation method in HPLC (Agilent 1100, Germany), a C18 column (150×4.6 mm i.d., 5 µm-ACE columns) and a mobile phase (Methanol-water (90:10, v/v) were used. It flowed at 1.1 mL/ min and was measured at 365 nm. The amount of PEITC was calculated according to the calibration curve of PEITC standards. The correlation coefficient was found to be 0,999 for PEITC (Figure 2). The amount of PEITC in watercress was expressed as µg PEITC per mg of watercress.



Figure 2. Calibration curve of standard solutions of PEITC.

Biochemical Analysis

Determination of protein content of liver tissues: The protein levels in the liver tissues were measured by the method of Bradford using BSA as the standard (21). Protein levels were expressed as milligram protein.

Determination of alanine aminotransferase (ALT): Rat ALT ELISA kit (Elabscience) was used to measure ALT level in the rat serum, according to the manufacturers' instructions. ALT level was expressed as ng/mL.

Determination of superoxide dismutase (SOD) activity: The SOD activity was measured using the Nitroblue tetrazolium (NBT) assay as described by Beauchamp and Fridovich (22). NBT-formazan was spectrophotometrically determined at 560 nm. SOD activity was expressed as U/mg protein. **Determination of glutathione peroxidase (GPx) activity:** GPx activity was determined by the method of Lawrence and Burk (23). The absorbance at 340 nm was recorded for 1 min. GPx activity was expressed as U/mg protein.

Determination of catalase (CAT) activity: CAT activity was determined by the method of Luck (24). The absorption was detected spectrophotometrically at 240 nm for 1 min. CAT activity was expressed as U/mg protein.

Determination of malondialdehyde (MDA): Amount of MDA was measured using the method of Mihara and Uchiyama (25). The absorption at 532 nm was recorded using 1,1,3,3-tetramethoxypropane as the standard. MDA content was expressed as nanomoles per milligram protein.

Determination of GST activity: GST activity assay kit (Cayman) was used to determine the activity of GST in the rat liver, according to the manufacturers' instructions. The absorption was monitored at 340 nm. GST activity was expressed as nmol/min/mg protein.

Determination of NQO1 activity: NQO1 activity assay kit (Abcam) was used to determine the activity of NQO1 in rat liver, according to the manufacturers' instructions. The absorbance was detected at 440 nm. NQO1 activity was expressed as U/mg protein.

Determination of Nrf2-DNA binding activity: To measure the DNA binding activity of Nrf2 in liver nuclear extracts Nrf2 Transcription Factor Assay Kit (Cayman) was used.

Histological Analysis

The liver tissues were fixed in 10% formalin solution for 24 h. Following embedded in paraffin, sections of the liver tissue were cut at 5 μ m and stained with hematoxylin-eosin (H-E) and Masson's trichrome methods. These samples were examined under light microscopy (Leica Micros Imaging Solutions Ltd., Cambridge, UK). The microscopic score of each tissue was calculated as the sum of the scores given to each criterion. Scores were given as absent (0), slight (1), moderate (2), and severe (3) for each criterion. The maximum score was 9.

Statistical Analysis

The IBM SPSS software version 22 was used for data analysis. Whether the data showed normal distribution was analyzed according to the Kolmogorov-Smirnov Test. For the data that had normal distribution, the One-Way ANOVA test was used for statistical analyses. The Posthoc LSD Test was used for the comparisons among the groups. For the data that didn't have a normal distribution, the Kruskal-Wallis test and the Post-hoc Mann-Whitney U test were used. Arithmetic mean \pm Standard Deviation (SD) was used for the biochemical data. Differences were considered significant when p values were less than 0.05 and 0,0001 for the biochemical and histological data, respectively.

RESULTS

Histological Findings

Histopathological analysis results were given in Table 1. The liver tissues of the control group did not present any histopathological alterations and significant damage was shown in the liver tissue of the DMBA-treated group (p <0.0001). Liver tissues of the DMBA+watercress (WS) group showed markedly reduced damage defined by less amount of mononuclear cell infiltration, hemorrhage, eosinophilic cells with a pyknotic nucleus (p < 0.0001). Histopathological images of the control, DMBA, and DMBA+WS groups are shown in Figures 3, 4, and 5, respectively. In the control group, the normal parenchymal structure was observed in liver tissue (Figure 3 (A,B)). There is no histological alteration of the central vein and surrounding hepatocytes in the portal areas. The liver tissue of the DMBA group (Figure 4 (A-E)) showed significantly histopathological damage. Liver necrosis (Figure 4 (A,E)), hepatocytes with a pyknotic nucleus and eosinophilic cytoplasm (Figure 4 (A,B)), hemorrhage (Figure 4 (A,B)), mononuclear cell infiltration (Figure 4 (C)), vascular congestion (Figure 4 (C,E)), and sinusoidal dilatation (Figure 4 (D)) were observed. The liver tissue of the DMBA+WS group presented a decrease in histopathological damage. In liver tissue of the DMBA+WS group, a small amount of mononuclear cell infiltration (black arrows), hemorrhage (Figure 5 (A)), eosinophilic cells with a pyknotic nucleus (black arrows) were observed (Figure 5 (B)).

Biochemical Findings

Alterations in serum ALT level related to liver function are represented in Figure 6. It was observed that serum ALT level increased as a result of the application of DMBA, and then decreased in the DMBA+WS group (p < 0.05). The DNA binding activity level of the Nrf2 transcription factor was decreased significantly (p < 0.05) in the DMBA group, and a significant increase (p < 0.05) was observed in the DMBA+WS group (Figure 7). Our results showed decreased activities of NQO1 and GST in the DMBA group. The administration of watercress significantly modulated the activities of NQO1 and GST (Figure 8). Furthermore, activities of SOD, GPx, and CAT were decreased in the DMBA group compared to the control and increased in the DMBA+WS group (Table 2). MDA levels were significantly elevated in the DMBA group and decreased by watercress administration (Table 2).

Table 1. The mean histopathological damage score of all groups. Lowercase letters a, b, c in the same column show the differences between groups. p<0.0001.				
Group	Histopathological damage score (Mean±SE)			
Control	0.39 ± 0.59^{a}			
DMBA	2.11±0.88 ^b			
DMBA+watercress	$1.30 \pm 0.80^{\circ}$			
Lowercase letters a, b, c in the same column show the differences between groups. $p<0.0001$				



Figure 3. (**A**,**B**) Liver specimen from the control group showing a normal appearance. In the control group, liver tissue was normal histological and hepatocytes arranged in the form of cords around the vena centralis were observed. A: H-E; X20, B: H-E; X 40.



Figure 4. (A-E) Liver specimen from the DMBA-treated group. In the DMBA-treated group, necrosis of the liver tissue (black star) (A, E), hepatocytes with eosinophilic stained pyknotic nuclei (thick black arrows) (A, B), haemorrhage (thin black arrows) (A, B), mononuclear cell infiltration (thin black arrows)) (C), vascular congestion (white star) (C, E) and sinusoidal dilatation (thin black arrows) (D) were observed. A, B, C, E: H-E; X20, D: H-E; X 40.



Figure 5. (A,B) Liver specimen from the DMBA+watercress group. In the DMBA+watercress group, mononuclear cell infiltration (black arrows) (A), haemorrhage (black arrows) (A), and hepatocytes with eosinophilic stained pyknotic nuclei (black arrows) (B) were observed. A: H-E; X 20, B: H-E; X 40.

Table 2. Effects of DMBA and DMBA + Watercress treatments on antioxidant parameters in rat liver tissues. Data are presented as mean±SD					
Group (n)	CAT (U/mg protein)	SOD (U/mg protein)	GPx (U/mg protein)	MDA (nmol/mg protein)	
Control (9)	7550.4±582.04	38.398±11.7	130.12±2.74	134.97±9.42	
DMBA (9)	6642.2±521.69ª	19.272±6.02ª	108.23±4.59ª	212.46±8.57ª	
DMBA+Watercress (9)	7425.4±262.54 ^b	67.691±14.9 ^a , ^b	118.66±3.92 ^a , ^b	152.98±3.07 ^a , ^b	
*p <0.05 compared to the control group; bp < 0.05 compared to the DMBA group.					







Figure 7. DNA binding activity level (%) of Nrf2 transcription factor in all groups. Data are presented as mean±SD. ^ap<0.05 versus control; ^bp<0.05 versus DMBA

DISCUSSION

The present study investigated both histopathological and biochemically whether watercress has a protective effect in DMBA-induced liver injury by Nrf2/ARE signalling pathway. DMBA is a potent carcinogen and causes liver damage via reactive metabolites and oxidative stress (26). In this study, we used histopathological findings and serum ALT level to evaluate the severity of DMBA-induced liver damage. We observed that there was a significant increase in ALT level in the DMBA group in comparison to the control group. Administration of watercress caused a significant decrease in ALT level compared to the DMBA group. Moreover, the histopathological analysis showed normal parenchymal structure in the control group, whereas DMBA showed significant histopathological damage



Figure 7. GST and NQO1 activities in all groups. Data are presented as mean \pm SD. ^ap<0.05 versus control; ^bp<0.05 versus DMBA

in the liver tissue. Reduction in histopathological damage was observed in the watercress treated group and a small amount of mononuclear cell infiltration, haemorrhage, eosinophilic stained pyknotic nucleus cells were observed. Taken together, histopathological data and serum ALT level of our study, we can suggest that watercress ameliorated DMBA-induced liver damage. In parallel with our results, Azarmehr et al. (27) reported that watercress has a hepatoprotective effect against acetaminophen-induced liver damage. Additionally, Doustimotlagh et al. (28) showed that Nasturtium officinale R. Br and quercetin combination cyclophosphamide-induced protected against hepatotoxicity in rats.

Overproduction of ROS triggers serious damage to various cells associated with increased MDA level that is an important marker of lipid peroxidation and oxidative damage. In our study, the highest tissue MDA level was observed in the DMBA group. MDA level was significantly lower in the watercress group when compared with the DMBA group (29). Activities of SOD, CAT, GPx were significantly decreased in DMBAtreated rats when compared to control group and administration of watercress enhanced the activities of SOD, CAT, GPx. The alteration in antioxidant status increases the risk of oxidative damage to nucleic acids, proteins, lipids, and small intracellular molecules (30). Many experimental studies reported that enhancement of oxidative stress is contributed to DMBA-induced cancer (31). In this study, administration of watercress significantly reduced DMBA-induced elevation lipid peroxidation and also decreased the activities of liver enzymatic antioxidants. In line with our results, Sadeghi et al. (32) reported that hydroalcoholic extract of watercress ameliorated oxidative stress and liver injury in bile duct ligation-induced cholestatic rats by preventing the hepatic protein oxidation and enhancing the activity of the antioxidant enzymes.

It has been reported that up-regulation of Keap1 and down-regulation of Nrf2 expression increased cellular ROS (33). Nrf2-mediated antioxidant signalling pathways are critical for the prevention of oxidative stress-induced cell injury (34). It has been showed that various dietary phytochemicals target Nrf2/Keap1 signalling and induce the expression of antioxidant and phase II enzymes to inhibit the development of DMBAinduced carcinogenesis model (35). Our results revealed that the binding percentage of the Nrf2 transcription factor had a statistically significant decrease in the DMBA group comparison to the control group. In the watercress group, the percentage of Nrf2/ARE binding significantly was activated compared to the DMBA group.These results can be associated with sulfur-containing glucosinolate derivatives, such as PEITC. These natural isothiocyanates reduce Nrf2 degradation which results in the translocation to the nucleus of Nrf2 (36). Previous studies reported that dietary isothiocyanates are an important cancer chemoprevention compound for the Nrf2/ARE system, which induces the activation and expression of phase II enzymes (37). Saravan et al. (38) were reported that thymoquinone protected liver tissue from DMBA toxicity by regulating phase I and phase II detoxification enzymes. In present study, there was a statistically significant decrease in phase II enzymes (NQO1 and GST) activities in the DMBA group compared to the control group. Administration of watercress significantly increased the activities of these enzymes compared to the DMBA group. According to our study, it can be suggested that there is a good correlation between the protective effect of watercress and supporting the DMBA detoxification by the Nrf2 signalling pathway. Induction of phase II detoxifying enzymes by Nrf2 assists to eliminate toxic reactive intermediates generated via xenobiotic metabolism (39).

There are two major limitations for our study. First, we were only be able to measure ALT to evaluate the level of liver damage. Changes in ALT level were supported with histopathological analysis, but we could not measure other liver function parameters due to financial conditions. Another limitation is that we did not design and conduct a study on the molecular mechanism about which way watercress affects the Nrf2/ARE pathway. This topic has been planned for our future research.

Overall, this experimental study suggests that watercress ameliorated oxidative stress and liver injury through the regulation of antioxidant and phase II enzymes by the activation Nrf2/ARE signalling pathway in DMBAinduced liver damage in rats.

CONCLUSION

The study revealed that the watercress plant is effective through the activation of Nrf2/ARE signalling resulting in increased antioxidant and phase II detoxification enzyme activities which are important for the chemoprevention of liver damage caused by DMBA. Our data support that activation of Nrf2 is a promising strategy for chemoprevention. In this context, we think that watercress functions as an Nrf2 activator and effect of its PEITC content on the molecular mechanism should be investigated with further studies

ETHICAL DECLARATIONS

Ethics Committee Approval: The study was carried out with the permission of the Local Ethics Committee of the Animal Experiments of İnönü University (Date: 20/02/2015, Decision No: 2015/A-25).

Referee Evaluation Process: Externally peer-reviewed.

Conflict of Interest Statement: The authors have no conflicts of interest to declare.

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Author Contributions: All of the authors declare that they have all participated in the design, execution, and analysis of the paper, and that they have approved the final version.

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