Upregulation of NF-E2-related factor-2-dependent glutathione by carnosol provokes a cytoprotective response and enhances cell survival

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Aim: To explore whether glutathione (GSH) increased through Nrf-2 activation is involved in the cytoprotective effects of carnosol in HepG2 cells.

Methods: Human hepatoma cell line HepG2 were exposed to rosemary essential oil or carnosol. Cell viability was measured using an Alamar blue assay. The production of intracellular GSH was determined using monochlorobimane. The level of protein or mRNA was examined by Western blotting or RT-PCR, respectively.

Results: Rosemarry essential oil (0.005%-0.02%) and carnosol (5 and 10 mol/L) increased the intracellular GSH levels and GSH synthesis enzyme subunit GCLC/GCLM expression. Rosemary essential oil and carnosol increased nuclear accumulation of Nrf2 and enhanced Nrf2-antioxidant responsive element (ARE)-reporter activity. Transfection of the treated cells with an Nrf2 siRNA construct blocks GCLC/GCLM induction. Furthermore, pretreatment of the HepG2 cells with essential oil and carnosol exerted significant cytoprotective effects against H2O2 or alcohol. In TNFx-treated cells, the nuclear translocation and transcriptional activity of NF-κB was abolished for 12 h following carnosol pretreatment. Cotreatment with GSH also suppressed NF-κB nuclear translocation, whereas cotreatment with BSO, a GSH synthesis blocker, blocked the inhibitory effects of carnosol.

Conclusion: This study demonstrated that Nrf2 is involved in the cytoprotective effects by carnosol, which were at least partially mediated through increased GSH biosynthesis.

Keywords: carnosol; glutathione; Nrf2; NF-κB; human hepatoma cell line HepG2; cytoprotection

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Introduction

Non-nutritional constituents in many foods may have beneficial health effects such as anti-inflammatory and anti-carcinogenic properties. Rosmarinus officinalis (rosemary) originates from southern Europe and is a commonly used herbal flavoring agent[1]. Rosemary extracts exhibit potent antioxidant activities that reduce lipid peroxidation, the production of reactive oxygen species (ROS), and inflammation[2, 3]. Carnosol is a diterpene derived from rosemary where it is found in considerable quantities; approximately 0.2%–1%, in dried rosemary[4], and 10.3% in commercially available rosemary extracts[5]. Although carnosic acid is the major polyphenolic compound present in rosemary plants, carnosol, an oxidation product of carnosic acid, has stronger anti-inflammatory effects[6]. Several studies including ours had been reported previously that the cytoprotective effects of carnosol have a number of beneficial properties from a medicinal standpoint, including antioxidation, anti-inflammation and anti-cancer effects in various cell types[7–9]. However, details of the mechanisms underlying the hepatic cytoprotection of carnosol and its regulation remain to be elucidated.

To protect and survive against a variety of environmental or intracellular stresses, mammalian cells have developed robust cellular defensive systems, including mechanisms that alleviate oxidative stress. Among the factors involved in these defense responses are components of detoxifying systems, including phase II drug metabolizing enzymes such as glutathione S-transferase, NAD(P)H: quinone oxidoreductase, and UDP-glucuronosyltransferase[10], and anti-oxidant enzymes such as glutamine-cysteine ligase (GCL)[11, 12]. Previous studies have shown that these enzymes are coordinately regulated through transcription factor NF-E2-related factor-2 (Nrf2) activation in response to electrophiles[10]. Carnosol possesses

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Our findings demonstrate that both substances induce GCL expression and also an elevation of the intracellular GSH levels. In addition, the increased GSH levels were found to be associated with the inhibition of TNFα-induced NF-κB nuclear accumulation.

Materials and methods

Materials

Bacterially derived TNFα was purchased from Calbiochem (San Diego, CA). The p3xARE/Luc vector was constructed as described previously[24]. Antibodies raised against Nrf2 were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). p65 antibodies were obtained from Stressgen Biotechnologies (SB, San Diego, CA, USA). ECL reagents were purchased from Pierce (Rockford, IL, USA). Luciferase assay kits were purchased from Promega (Madison, WI, USA). Peroxidase-conjugated anti-rabbit and anti-mouse antibodies were obtained from Amersham (Arlington Heights, IL, USA) and nitrocel lulose was obtained from Schleicher & Schuell (Dassel, Germany). All other reagents, including carnosol and rosemary essential oil were purchased from Sigma (St Louis, MO, USA).

Cell culture

HepG2 cells (ATCC HB-8065) were grown in high glucose Dulbecco’s modified Eagle’s medium (DMEM, Invitrogen) supplemented with 10% fetal bovine serum (FBS, Invitrogen), 100 U/mL penicillin and 100 µg/mL streptomycin. The cells were seeded at a density of 1.5 million cells per mL in T-25 culture flasks with 5 mL of complete medium and cultured at 37 °C with 5% carbon dioxide. The culture medium was then replaced with serum free DMEM and the cells were incubated for 12 h prior to experimental treatment.

GSH Assay

GSH levels were determined as described previously[17]. Briefly, cells were cultured at 37 °C in the presence or absence of the treatment reagents indicated in the corresponding figures, washed with PBS and incubated with monochlorobimane (MCB, 40 µmol/L) in the dark for 20 min at room temperature. After two further washes with PBS, the cells were solubilized with 1% SDS and 5 mmol/L Tris HCl (pH 7.4). Fluorescence was measured by spectrofluorophotometry (Shimadzu, RF-5301PC), with excitation and emission wavelengths of 405 and 510 nm, respectively and samples were assayed in triplicate. The assay for detecting in vitro GSH levels was performed identically but without cell lysates. The levels of intracellular GSH were quantified using a GSH solution as a standard.

Cell viability assay

Cell viability was performed using an Alamar blue assay kit (Sero tec, Oxford, UK) in accordance with the manufacturer’s instructions. This assay is based on the detection of metabolic activity in living cells using a redox indicator that changes from an oxidized (blue) to a reduced (red) form. The intensity of the red color is proportional to the viability of the cells, and is calculated by the difference in the absorbance values at 570 and at 600 nm and expressed as a percentage of the control.

RNA isolation and RT-PCR

Total cellular RNA was extracted using the phenol-guanidinium isothiocyanate method[28]. Equal amounts (5 µg) of RNA from the different treatments were then reverse-transcribed for 50 min at 42°C using 50 units of Superscript II (Invitrogen, Carlsbad, CA). Amplifications of the cDNA were performed in 25 µL of PCR buffer (10 mmol/L Tris-HCl, 50 mmol/L KCl, 5 mmol/L MgCl2, and 0.1% Triton X-100, pH 9.0) containing 0.6 units of Taq DNA polymerase (Promega, Madison, WI) and 30 pmol of the specific primers GCLM forward, 5’-CAGCGAGGAGCTTCATGATTG-3’; reverse, 5’-TGATCACAGAATCCAGTGCC-3’; GCLC forward, 5’-GGTCTTGAATCTCTGCAAGAAG-3’; reverse, 5’-ATGGAGATGTGTTTATCTGTTCC-3’[19] and GAPDH forward 5’-TATCGTGGAGGACTCATGACC-3’; reverse 5’-TACATGGCAACTGTGAGGGG-3’. Reaction products were separated electrophoretically in a 2.5% agarose gel and stained with ethidium bromide.

Quantitative PCR

For transcript quantification purposes, real-time PCR was performed. Then 1 µL of the reverse-transcriptase product was used in a 25 µL volume reaction containing 200 µmol/L of each dNTP, 5 pmol of each primer, 1×PCR buffer II, 1.5 mmol/L MgCl2, 0.2×SYBR Green I (Molecular Probes), and 1.25 U of AmpliTaq Gold (Applied Biosystems). Amplification was performed using ABI 7500 Real-Time PCR System (Applied Biosystems) programmed as 94 °C for 12 min followed by 40 cycles of (94 °C for 30 s, 57 °C for 20 s, 72 °C for 30 s). Variability in the initial quantities of cDNA was normalized to the internal control, GAPDH. A negative control was included in each set of experiments. Melting curve analysis was performed to enhance specificity of the amplification reaction, and the 7500 software was used to compare the amplification in the experimental samples during the log-linear phase.

Plasmids, transfections and luciferase assays

A NF-kB/Luc fragment containing tandem repeats of double-stranded oligonucleotides spanning the NF-kB binding site of ICAM-1: 5’-TGAAATCCC-3’[20], sense: 5’-CCCGGGTGGAATTTCCCTGCAAATTTCCCGGG-3’; 3’-TTAGATCCGGAATTTCAGGAATTTCTTCCACCCCGG-3’ was introduced into the pGL3 promoter plasmid (Promega, Madison,
The HepG2 cells were then grown to 60%–80% confluence and transfected with a total of 1 μg of NF-κB/Luc or Nrf2/Luc using lipofectamine reagent (Invitrogen, Carlsbad, CA, USA) in accordance with the manufacturer’s instructions. For luciferase assays, the cell lysate was first mixed with luciferase substrate solution (Promega), and the resulting luciferase activity was measured in a luminometer. For each experiment, luciferase activity was determined in triplicate and normalized with respect to β-galactosidase activity levels.

Transient transfection with siRNA targeting Nrf2
An siRNA targeting human Nrf2 5’-UCCCGUUUGUAGAU-GACAA-3’ and a control siRNA 5’-GCAAGCUGACCCU-GAAGUUCAU-3’ (non-sense) were purchased from Ambion (Austin, TX, USA). HepG2 cells were seeded onto 60-mm dishes, incubated for 24 h, and then transiently transfected with 100 nmol/L siRNA per dish at 90% confluence with Lipofectamine 2000. After 24 h of recovery in 10% serum medium, the cells were cultured in serum-free medium without serum for another 12 h prior to treatment.

Preparation of cytosolic and nuclear lysates
To separate the cytosolic and nuclear fractions from HepG2 cells, the cells were collected by scraping in cold PBS. The cell pellet was then lysed in 10 mmol/L HEPES, 1.5 mmol/L MgCl₂, 10 mmol/L KCl, 0.5 mmol/L DTT, 0.5 mmol/L PMSF and 0.3% nonidet P-40. After 5 min of centrifugation (3000 rounds per minute at 4 °C) the supernatant was collected and designated as the cytosolic fraction. Nuclear proteins were then extracted from this preparation using a buffer containing 25% glycerol, 20 mmol/L HEPES, 0.6 mol/L KCl, 1.5 mmol/L MgCl₂ and 0.2 mmol/L EDTA. Protein concentrations were determined using a protein assay DC system (Bio-Rad, Richmond, CA, USA).

Western blotting
Whole lysates of HepG2 were prepared as previously described[22]. A total of 1×10⁶ cells were lysed on ice in lysis buffer (1% NP-40, 0.5% sodium deoxycholate, 0.1% SDS and a protease inhibitor mixture) and whole-cell extracts were boiled for 5 min prior to separation on 10% SDS-PAGE, in which the protein samples were evenly loaded. The proteins were then transferred to a nitrocellulose filter in Tris-glycine buffer at 100 V for 1.5 h. The membranes were then blocked with PBS containing 5% nonfat milk and incubated with antibodies for two hours at 4 °C, with gentle shaking. The results were visualized by chemiluminescence using ECL (Pierce, Rockford, IL, USA), according to the manufacturer’s instructions.

Figure 1. The GSH levels are increased in HepG2 cells treated with rosemary essential oil. (A) The intracellular GSH levels of HepG2 cells incubated with 0.01% rosemary essential oil for 1, 6 and 12 h. Data values are expressed as a percentage of the untreated control, which was set at 100%. Results are the mean±SEM (n=3). *P<0.05 vs untreated cells. (B) The intracellular GSH levels of HepG2 cells incubated with the indicated doses of rosemary essential oil for 12 h. (C) Cell viability of HepG2 cells incubated with 0.005% and 0.01% rosemary essential oil for 24 h. Data are expressed as the mean±SEM of three independent experiments. No significant differences were found by ANOVA.
Upregulation of both GSH and GSH synthesis enzyme levels in HepG2 cells by carnosol and essential oil of rosemary

The rate-limiting enzyme in the de novo synthesis of GSH is glutamate-cysteine ligase (GCL), also known as γ-glutamylcysteine synthetase. GCL consists of a catalytic heavy subunit (GCLC) and a modulatory light subunit (GCLM). We assayed both GCLC and GCLM expression in essential oil and carnosol-treated HepG2 cells.

Treatment with 0.01% essential oil increased both the GCLC and GCLM expression levels over the course of the incubation time (Figure 3A). As shown in Figure 3B, an increase in both the GCLC and GCLM mRNA levels was detected following essential oil treatments at 0.01% and 0.02% for 12 h. In a similar experiment, carnosol exposure increased both the GCLC and GCLM gene expression levels after three hours treatment (Figure 3C). In addition, carnosol at 5 µmol/L and 10 µmol/L concentrations was found to increase GCLC and GCLM gene expression over 12 h treatments (Figure 3D). To further determine the effect of 0.01% essential oil and carnosol, cells were incubated for different times. Quantitative PCR reactions were performed to amplify GCLC and GCLM, the production of GCLC and GCLM were increased significantly by 0.01% essential oil or carnosol as early as 3 h and reached a maximum by 6 h (Figure 3E and 3F).

Effects of carnosol upon Nrf2 activation

In our present study, we have found that 0.01% rosemary essential oil and carnosol (5 μmol/L) increase the level of Nrf2 in the nucleus of HepG2 cells. We thus evaluated the specificity of carnosol for Nrf2 binding site, antioxidant response element (ARE) sequences, in our current study by transfecting HepG2 cells with luciferase reporter constructs harboring this element. Cells treated with 0.01% essential oil, and 5 µmol/L or 10 µmol/L carnosol, indeed displayed increased ARE-luciferase activity (Figure 4B). To further assess the mediating role of Nrf2 in the inhibitory effects of carnosol, a more targeted inhibition of Nrf2 using siRNA was undertaken. Cells were transfected with Nrf2 siRNA to reduce the Nrf2 protein level (Figure 4C) and this abolished the induction of both GCLC and GCLM expression following 12 h of carnosol pre-treatment (Figure 4D). These findings suggested that carnosol promotes GCLC and GCLM expression via the activation of the Nrf2 pathway.

The protective effects of carnosol and rosemary essential oil against oxidative stress and ethanol

We investigated the effects of rosemary essential oil and carnosol on H2O2-induced cytotoxicity and as shown in Figure 5A, it was found that HepG2 cells treated with 3 mmol/L H2O2 showed significantly reduced cell viability. In contrast, this cytotoxicity was significantly reduced in cells pretreated with 0.01% essential oil or 5 µmol/L carnosol.

It is well-known that excess alcohol in the liver induces hepatotoxicity. To evaluate whether essential oil and carnosol had protective effects in this regard, we examined their effects upon cell viability in the presence of excess alcohol. The addition of 50 mmol/L ethanol to the culture medium tended to reduce cell viability (Figure 5B). In the presence of essential oil or carnosol however, no loss of cell viability was evident upon exposure to excess alcohol (Figure 5B). These results suggest that the protection against damage from oxidative stress or excess alcohol is dependent on an increased GSH level.

Carnosol inhibits TNFα-induced NF-κB activation via upregulated GSH

To evaluate its effects on NF-κB activation in HepG2 cells,
cells were treated with GSH for 30 min before TNFα stimulation. As shown in Figure 6A, GSH indeed inhibits TNFα-induced NF-κB nuclear accumulation. We next investigated the possible role of GSH in mediating the inhibitory effects of carnosol in HepG2 cells. Pretreatment of cells with buthionine sulfoximine (BSO), a specific inhibitor of γ-glutamyl cysteine synthetase at concentrations of 100 µmol/L abolished the suppressive effects of carnosol upon NF-κB activation (Figure 6B). Furthermore, we tested whether carnosol inhibits TNFα-induced P65 activation at the transcriptional level. Following pretreatment for 12 h and a luciferase assay (Figure 6C), we found that TNFα-induced NF-κB activation was indeed inhibited. The results of these experiments indicate that the inhibitory effects of carnosol are the result of increased cellular GSH levels.

**Discussion**

GSH is a well-studied tri-peptide and has numerous roles in protecting cells from oxidants and maintaining the cellular thiol redox status\[23, 24\]. It was found in our previous study that cinnamaldehyde increases the cellular GSH levels in HepG2 cells after 9 h of exposure\[25\]. Carnosol is an electrophilic phytochemical present in the rosemary herb and our present study provides new evidence that both rosemary essential oil and carnosol enhance the GSH levels in HepG2 cells by upregulating the expression of GCLC and GCLM at non-cytotoxic concentrations. Our present experiments also show that the translocation of Nrf2 into the nucleus following treatment with rosemary essential oil and carnosol is associated with increases in its ARE transcriptional activity. We further demonstrate that Nrf-2 siRNA abolishes the induction of GCLC and GCLM by carnosol. Hence, the presence of activated Nrf-2 is required for the protective effects of carnosol. These protective effects of both rosemary essential oil and carnosol were manifested by the maintenance of cell viability under conditions of oxidative stress and following ethanol treatment. In addition, the inhibitory effects of carnosol upon TNFα-induced NF-κB activation are mediated through an increase in the intracellular GSH level.

Oxidative stress occurs when the redox equilibrium, which is the ability of cells to protect against damage caused by production of free radicals, is disrupted. Oxidative stress has been implicated as a major cause of cellular injuries in a variety of human diseases. Previous studies have found that poly-
phenolic compounds have intrinsic antioxidant characteristics and protect cells against oxidative cell damage via a Michael acceptor function. These compounds include curcumin, phenylethyl isothiocyanate, epigallocatechin gallate, and other green tea polyphenols\[7\]. Recently, much attention has focused on the upregulation of phase II detoxifying and antioxidant enzymes via the activation of the Nrf2 transcription factor\[15, 26\]. Carnosol, a diterpene derived from the rosemary herb, is a representative member of a family of plant-derived phenols\[13\]. Our present experiments show that carnosol induces Nrf2-mediated GCLC and GCLM expression, which in turn controls GSH synthesis\[27\]. Furthermore, our present data demonstrate that carnosol induces Nrf2 translocation and activates ARE-luciferase promoter activity, indicating that it directly induces Nrf2 via its ARE.

Figure 5. The protective effects of rosemary essential oil and carnosol against oxidative stress and alcohol. Cells were initially preincubated for 12 h in the presence of the indicated doses of rosemary essential oil or carnosol. The medium was then removed and the cells were exposed to 3 mmol/L H_{2}O_{2} for 12 h. Cell viability was measured spectrophotometrically using an Alamar blue assay. Data are expressed as the means±SEM of at least three independent experiments. \(^{\text{a}}P<0.05\) vs control; \(^{\text{b}}P<0.05\) vs H_{2}O_{2} alone. (B) Cells were initially preincubated for 12 h in the presence of the indicated doses of essential oil or carnosol. The medium was then removed and the cells were exposed to 50 mmol/L ethanol for 12 h. Data are expressed as the means±SEM of at least three independent experiments. \(^{\text{a}}P<0.05\) vs control; \(^{\text{b}}P<0.05\) vs ethanol alone.

Figure 4. Effects of carnosol upon Nrf2 activation. (A) Nuclear extracts from HepG2 cells were prepared after treatment with 0.01% rosemary essential oil and 5 µmol/L carnosol for the indicated time periods. Immunoblots of nuclear lysates were then probed with Nrf2 specific antibodies. The nuclear lamin band intensities indicate equal loading of each well. (B) Cells were transfected with the ARE-luciferase construct (ARE) and then stimulated with 0.01% rosemary essential oil, 5 or 10 µmol/L carnosol. The cells were then lysed and analyzed for luciferase activity. Induction is indicated by an increase in the normalized luciferase activity in the treated HepG2 cells, relative to the control. Results are the means±SEM from at least three separate experiments. \(^{\text{a}}P<0.05\) vs untreated HepG2 cells. (C) HepG2 cells were transfected with control or Nrf2 siRNA for 36 h and the intracellular protein levels of Nrf2 were determined by Western blotting. (D) HepG2 cells were transfected with control or Nrf2 siRNA for 36 h and then exposed to 5 µmol/L carnosol for 12 h. The mRNA levels of GCLM and GCLC genes were determined by RT-PCR.
increased GSH in TNFα-treated cells. TNFα stimulation has been shown to activate NF-κB signaling pathways\textsuperscript{[32]} . We found in our current analyses that carnosol increases the GSH levels after six hours of treatment, and that BSO abolishes p65 translocation after a 12-h incubation with carnosol. In addition, treatment with GSH alone also inhibits p65 translocation. Thus, an increased GSH level is essential for carnosol-induced anti-inflammatory effects. Previous studies have demonstrated that changes in the thiol redox state of the cell might affect the posttranslational modification of p65, including the phosphorylation of critical residues shown to contribute to the nuclear import of this molecule\textsuperscript{[33, 34]} . Moreover, a recent study has revealed that a number of redox-sensitive transcription factors are modified by GSH and thereby inhibit their function\textsuperscript{[35]} . The p65 modification by GSH is thus implicated as an inhibitory mechanism by which carnosol regulates NF-κB activation through the increase in GSH levels.

Our present data thus reveal that carnosol induces cytoprotective mechanisms that may contribute to its putative beneficial effects in suppressing the liver cell response to oxidative stress, alcohol or cytokines during inflammation. Our present results thus expand our understanding of the role of phytochemicals in cytoprotection and potentially assist in the identification of new therapeutic strategies for diseases caused by oxidative damage and other environmental stresses.

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**Author contribution**

Chien-chung CHEN analyzed data and contributed analytic tools. Hui-ling CHEN performed research. Chia-wen HSIEH designed research. Yi-ling YANG contributed analytic tools. Being-sun WUNG designed research, analyzed data and wrote the paper.

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