Research article

Induction of Nrf2/ARE-mediated cytoprotective genes by red ginseng oil through ASK1–MKK4/7–JNK and p38 MAPK signaling pathways in HepG2 cells

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ABSTRACT

Background: The induction of cellular defensive genes such as phase II detoxifying and antioxidant enzymes is a highly effective strategy for protection against carcinogenesis as well as slowing cancer development. Transcription factor Nrf2 (nuclear factor E2-related factor 2) is responsible for activation of phase II enzymes induced by natural chemopreventive compounds.

Methods: Red ginseng oil (RGO) was extracted using a supercritical CO2 extraction system and chemical profile of RGO was investigated by GC/MS. Effects of RGO on regulation of the Nrf2/antioxidant response element (ARE) pathway were determined by ARE-luciferase assay, western blotting, and confocal microscopy.

Results: The predominant components of RGO were 9,12-octadecadienoic acid (31.48%), bicyclo[10.1.0]tridec-1-ene (22.54%), and 22,23-dihydrostigmasterol (16.90%). RGO treatment significantly increased nuclear translocation of Nrf2 as well as ARE reporter gene activity, leading to upregulation of heme oxygenase-1 and NAD(P)H:quinone oxidoreductase 1. Phosphorylation of the upstream kinases such as apoptosis signal-regulating kinase (ASK)1, mitogen-activated protein kinase (MAPK) kinase (MKK)4/7, c-Jun N-terminal kinase (JNK), and p38 MAPK were enhanced by treatment with RGO. In addition, RGO-mediated Nrf2 expression and nuclear translocation was attenuated by JNK inhibitor SP600125 and p38 MAPK inhibitor SB202190.

Conclusion: RGO could be used as a potential chemopreventive agent, possibly by induction of Nrf2/ARE-mediated phase II enzymes via ASK1–MKK4/7–JNK and p38 MAPK signaling pathways.

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1. Introduction

Carcinogenesis is a multistep process including initiation, promotion, and progression, in which the accumulation of genetic alterations leads to transformation of an initiated cell into a population of preneoplastic cells and eventually a tumor with invasive and metastatic capacities [1]. Recently, many dietary phytochemicals have exhibited beneficial effects on health including prevention, delay, and inhibition of cancer progression [2]. Cancer chemoprevention is defined as the use of natural, synthetic, or biological chemical agents to prevent the early preneoplastic stage of carcinogenesis [3]. Among the underlying mechanisms of chemopreventive agents is the induction of phase II detoxifying/antioxidant enzymes involved in carcinogen detoxification, and antioxidants, such as heme oxygenase (HO)-1, NAD(P)H:quinone oxidoreductase (NQO)1, aldo–keto reductase (AKR), and
glutathione S-transferase (GST). Accumulating evidence shows that the transcription factor Nrf2 (nuclear factor-E2-related factor 2) is involved in the regulation of phase II enzymes through the activation of antioxidant response element (ARE), which is located in the promoter region genes encoding these enzymes [4–6]. Under physiological conditions, Nrf2 is sequestered in the cytoplasm by binding to Keap1 (Kelch-like ECH-associated protein 1, cytosolic repressor). Upon stimulations, however, Nrf2 dissociates from Keap1 and translocates into the nucleus where it binds to ARE to transcriptionally induce defensive genes [7,8]. Studies of Nrf2-deficient mice have highlighted the importance of phase II detoxifying/antioxidant enzymes in the deactivation of chemical carcinogens, as these mice were more susceptible to carcinogenesis than normal mice were [9,10].

Many studies have demonstrated that activation of the Nrf2/ARE pathway is related to the upstream kinases, including mitogen-activated protein kinases (MAPKs), protein kinase C, phosphatidylinositol 3-kinase (PI3K), or transmembrane kinase [11]. MAPK families, including c-Jun N-terminal kinase (JNK), extracellular signal-regulating kinases (ERKs) and p38 MAPK are key signaling molecules that respond to mitogenic stimulation or environmental stress, resulting in expression of target proteins [12]. Specific MAPK inhibitors have been shown to block the induction Nrf2 as well as phase II detoxifying/antioxidant enzymes [13,14]. MAPKs are regulated by their upstream kinases: ERK is activated by MAPK kinase (MEK)1 and MEK2, JNK by MKK4 and MKK7, and p38 MAPK kinase by MKK3, MKK4, and MKK6, which are activated by upstream MAPK kinase kinases (MAPKKKs) [15,16].

Ginseng (Panax ginseng Meyer) is a representative herbal, which has been widely used in Korea, China, and Japan for about 2,000 years. Red ginseng is made by steaming and drying fresh ginseng, a process that chemically transforms components and acquires special physiological activities such as antioxidant [17], antidiabetic [18], antiobesity [19], and anticarcinogenic [20] effects. Although several pharmacological effects of red ginseng have been studied, they are mostly water-soluble fractions. Moreover, the biological activities and molecular mechanisms of lipid-soluble moieties of red ginseng have been poorly characterized. Lipophilic fractions from red ginseng extracted with hydrophobic organic solvents such as petroleum ether and hexane have been shown to have antiinflammatory and antitumor effects [21,22]. Our group has recently reported anti-inflammatory and hepatoprotective mechanisms of supercritical CO2-extracted red ginseng oil (RGO) in vitro and in vivo [23,24]. In addition, our previous study has demonstrated the safety of RGO in an acute toxicity study using male and female Sprague–Dawley rats [25].

In the present study, we examined the chemical profile of supercritical CO2-extracted RGO and its underlying mechanisms in the induction of cellular defense system in HepG2 cells.

2. Materials and methods

2.1. Chemicals

Primary antibodies against Nrf2, NQO1, β-actin, and Lamin B were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA); HO-1, phospho-(p)-ERK1/2, p-JNK, p-p38, p-Akt, p-MAPK/ERK kinase (MEK)1/2, p-MKK3/MKK6, p-B-Raf, mixed lineage protein kinase (MLK)3, stress-activated protein kinase/ERK kinase-1 (SEK1)/MKK4, p-MKK7, phospho-apoptosis signal-regulating kinase (p-ASK1), and phospho-transforming growth factor β-activated kinase (p-TAK1) antibodies were purchased from Cell Signaling Technology (Boston, MA, USA). Peroxidase-conjugated secondary anti-rabbit and anti-goat antibodies were purchased from Santa Cruz Biotechnology, and Alexa-Fluor-555-conjugated secondary antibody was from Cell Signaling Technology (Boston, MA, USA). Specific inhibitors including U0126 (MEK1/2 inhibitor), SP600125 (JNK inhibitor), SB202190 (p38 MAPK inhibitor), and LY294002 (PI3 kinase inhibitor) were obtained from Cell Signaling Technology. Triton X-100, polyethylene glycol and other chemicals were from Sigma-Aldrich (St. Louis, MO, USA).

2.2. Preparation of RGO

RGO was prepared as previously described [24]. Dried red ginseng powder was placed into the extraction vessel of a pilot-scale supercritical fluid extraction system (Ilshin Autoclave Co. Ltd., Daejeon, Korea). Extractions with supercritical CO2 were operated at 6,500 psi (relative to 450 bar) in combination with temperature at 65°C. Extracted constituents were collected in a vial that was prefilled with a trapping solvent and maintained at 4°C during the extraction step.

2.3. GC/MS

Analysis of RGO was performed using an HP-5MS capillary column (30 m × 0.25 mm, 0.25 μm; Agilent Technologies, Santa Clara, CA, USA) in a Gas chromatography–mass spectrometry (GC/MS) (5,975C; Agilent Technologies). Samples were injected into the column and run using split mode (split ratio = 10:1). The helium carrier gas was programmed to maintain a constant flow rate of 1 mL/min. Oven temperature was initially 80°C for 3 min and then finally raised to 300°C at 4°C/min. Compounds were tentatively identified by comparing mass spectra of the peaks with those in the mass spectrum library of NIST 11.

2.4. Cell culture

Human hepatoma HepG2 cell line was purchased from American Type Culture Collection (Manassas, VA, USA). HepG2-C8 cell line was established by the stable transfection of HepG2 cells with p-ARE-T1-luciferase reporter gene as previously described [26]. Both cell lines were maintained in F-12 medium with 10% fetal bovine serum, 100 U/mL penicillin, 100 μg/mL streptomycin, 1% essential amino acids, 1% GlutaMAX, and 0.1% insulin. The cells were maintained in a humidified incubator at 37°C in 5% CO2.

2.5. Cell viability assay

HepG2 cells were plated at 105 cells/well in 24-well plates and treated with vehicle (polyethylene glycol: dimethyl sulfoxide, 1:1, v/v) or various concentrations of RGO for 24 h. After incubation, each well was washed twice with phosphate-buffered saline (PBS) followed by incubation with 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) for 4 h at 37°C. Dark blue formazan crystals were dissolved with dimethyl sulfoxide and absorbance was measured with a PowerWave XS microplate reader (BioTek Instruments, Inc., Winooski, VT, USA) at 570 nm. The percentage of viable cells was estimated by comparison with vehicle-treated control cells.

2.6. ARE–luciferase activity assay

HepG2-C8 cells were plated at 106 cells/well in 12-well plates and incubated for 24 h. After starvation overnight, cells were treated with RGO for 12 h, washed twice with PBS, and lysed with the reporter lysis buffer supplied for luciferase assay system (Promega, Madison, WI, USA). After centrifugation at 3,000 g for 10 min, a 10-μL aliquot of the supernatant was assayed for luciferase activity with a GloMax luminometer (Promega). The
Luciferase activity was normalized against protein concentration, determined using a BCA protein assay (Pierce Biotechnology, Rockford, IL, USA), and expressed as fold of induction over luciferase activity of vehicle-treated control cells.

2.7. Preparation of whole-cell, nuclear and cytosolic extracts

Whole-cell extract was prepared using cell lysis buffer containing 20 mM Tris (pH 7.5), 135 mM sodium chloride, 2 mM EDTA, 2 mM dithiothreitol, 25 mM β-glycerophosphate, 2 mM sodium pyrophosphate, 10% glycerol, 1% Triton X-100, 1 mM sodium orthovanadate, 10 mM sodium fluoride, 10 μg/mL aprotinin, 10 μg/mL leupeptin, and 1 mM phenylmethylene sulfonfonyl fluoride. Nuclear–cytoplasmic fractionation was conducted using NE-PER Nuclear and Cyttoplasmic Extraction Reagents Kit (Pierce Biotechnology), and protein concentrations were determined by BCA protein assay.

2.8. Western blotting

Proteins (whole-cell extracts: 30 μg/lane, nuclear extracts: 10 μg/lane, and cytosolic extracts: 30 μg/lane) were separated on 10% sodium dodecyl sulfate polyacrylamide gels. The separated proteins were electrophoretically transferred onto polyvinylidene fluoride membranes (Millipore, Billerica, MA, USA) with a semidyen transfer system (Bio-Rad, Hercules, CA, USA). The membranes were incubated with blocking solution (5% nonfat milk) for 1 h, and then incubated overnight with proper primary antibodies. After hybridization with primary antibodies, the membranes were washed three times with PBST solution and incubated with horseradish peroxidase-conjugated secondary antibodies for 1 h. The membranes were again washed three times with PBST, and detected using Western Blotting Luminol reagents (Santa Cruz Biotechnology).

2.9. Confocal microscopy

Cells were cultured in coverglass-bottom dishes. After starvation, the cells were pretreated with specific inhibitors for 1 h and then with RGO for 2 h. The cells were fixed with 4% paraformaldehyde in PBS for 10 min and permeabilized with 0.3% Triton X-100 for 10 min. After incubation with blocking buffer (3% bovine serum albumin in PBS) for 1 h, cells were incubated with Nrf2 antibody (1:100) in 0.3% bovine serum albumin for 2 h. The cells were stained for 2 h with Alexa-Flour-555-conjugated secondary antibody (1:200) in the dark, and finally incubated with 1 μg/mL of 4',6'- diamidino-2-phenylindole for 20 min in the dark. Images were obtained using a LSM 510 laser confocal microscope (Zeiss, Jena, Germany).

2.10. Statistical analysis

The data were expressed as mean ± standard deviation values by Microsoft Excel 2010 software (Microsoft Corporation, Redmond, WA, USA). The values were compared with the control using analysis of variance followed by unpaired Student t tests. A value of p < 0.05 was considered significant.

3. Results

3.1. Chemical composition of RGO

Table 1

| Retention time | Compound | Molecular formula | Peak area (%) |
|---------------|----------|-------------------|---------------|
| 26.98         | Palmitic acid | C16H32O2 | 2.89          |
| 29.23         | Linoleic acid | C18H32O2 | 31.48         |
| 44.36         | Ergosta-4,6,22-trien-3 alpha-ol | C29H48O | 4.53          |
| 45.00         | α-tocopherol | C29H48O2 | 1.01          |
| 46.54         | Stigmasterol | C29H48O | 3.14          |
| 47.27         | β-sitosterol | C29H48O | 16.90         |
| 47.64         | γ-sitosterol | C29H48O | 4.31          |
| 35.79         | Likely bicyclo[10.1.0]tridec-1-ene | C18H22 | 22.54         |
| 43.22         | Stigmastan-3,5-diene | C29H46 | 0.69          |
| 46.87         | Triteraccontane | C33H48 | 0.93          |
| 49.05         | Stigmast-4-en-3-one | C28H46O | 1.01          |

The chemical composition of the supercritical CO2-extracted RGO was analyzed for the first time using GC/MS (Table 1). The identified compounds were composed of three major groups including acids (34.37%), alcohols (29.89%), and hydrocarbons (24.07%). RGO contained fatty acids including linoleic acid (18:2, 31.48%), and palmitic acid (16:0, 2.89%). Ginseng seed oil has been reported to have oleic acid (C18:1) as the most predominant fatty acid (80%) followed by linoleic acid (16%) and palmitic acid (2%) [27]. Unlike ginseng seed oil, oleic acid was not detected and linoleic acid was the most predominant fatty acid in RGO. A previous study analyzing fatty acid composition of a lipophilic fraction from red ginseng extracted with petroleum ether has shown that linoleic accounts for about 75% of fatty acids in the fraction followed by palmitic acid (10%) [22], which corresponds to our results. For alcohols, β-sitosterol was the most abundant (16.90%) in RGO while ergosta-4,6,22-trien-3-ol (4.53%), γ-sitosterol (4.31%), stigmasterol (3.14%), stigmasterol, and α-tocopherol (1.01%) were also detected. Stigmasterol and β-sitosterol have been previously found in ginseng seed oil [27]. A hydrocarbon bicyclo[10.1.0]tridec-1-ene was detected with a significant amount (22.54%) in RGO. This compound has been found in the fruit of an Iranian berry (Bereberis integrigina) [28] and the essential oil from buds of Tussilago farfara L. [29]. Three compounds including linoleic acid, bicyclo[10.1.0]tridec-1-ene, and β-sitosterol occupied nearly 71% of RGO.

3.2. Effect of RGO on cell viability

Hepatocytes play a crucial role in the processes of metabolism, detoxification, modification, and excretion of exogenous and endogenous substances, implying that it is a target for the toxic action of xenobiotics or their reactive metabolites [30]. Enhancement of cellular defense system importantly contributes to protection of cells/tissues from damage caused by toxicants or oxidative stress. Consequently, HepG2 human hepatoma cell line is considered as a good model for studying the inductive effect of natural products on phase II detoxifying/antioxidant enzymes, which directly involve in cytoprotective action [31]. In the present study, the viability of HepG2 cells was determined to evaluate the cytotoxic effect of RGO using MTT assay. RGO treatment for 24 h resulted in a decrease of cell viability in a dose-dependent manner (Fig. 1). The treatment of HepG2 cells with 50 μg/mL RGO was unlikely to cause significant cell death, and therefore, RGO concentrations < 50 μg/mL were selected for subsequent experiments of activation of Nrf2/ARE-mediated cytoprotective genes.

3.3. Induction of phase II detoxifying/antioxidant enzymes by RGO

Phase II enzymes play important roles in the protection of cells/tissues from endogenous and/or exogenous carcinogens [32]. Accordingly, we first examined the inductive effect of RGO on the phase II enzymes expression after 24 h treatment. The protein
expression levels of HO-1 and NQO1 significantly and dose-dependently increased in RGO-treated HepG2 cells (Fig. 2). RGO treatment at the concentration of 50 \( \mu \text{g/mL} \) elevated protein levels of HO-1 and NQO1 by 160% and 200%, respectively, compared to untreated controls. Similarly, expression of HO-1 and NQO1 was strongly induced in the treatment of sulforaphane (SFN), a known phase II genes activator.

3.4. Induction of Nrf2/ARE machinery by RGO

To investigate whether the induction of phase II detoxifying/antioxidant enzymes by RGO was associated with the activation of Nrf2/ARE pathway, we examined the ARE luciferase activity and Nrf2 activation in RGO-treated HepG2-C8 and HepG2 cells, respectively. As shown in Fig. 3A, RGO significantly enhanced ARE luciferase activity in a dose-dependent manner and RGO at 50 \( \mu \text{g/mL} \) exerted greater induction than SFN; a potent natural inducer of Nrf2/ARE pathway [33]. ARE is a crucial regulatory element found in the promoter region of many cellular defensive genes encoding phase II detoxifying and/or antioxidant enzymes including NQO1 and HO-1 [5], suggesting that the induction of phase II enzymes by RGO could be mediated by the transcriptional activation of ARE. Based on the potent inductive ability of RGO on the ARE transcriptional activity, the effect of RGO on Nrf2 protein expression and its nuclear translocation was analyzed. Similar to luciferase activity, RGO upregulated protein expression of Nrf2 in a dose-dependent manner (Fig. 3B). In addition, the level of Nrf2 protein in the nucleus was gradually elevated with the dose of RGO, whereas protein level of Nrf2 in the cytoplasm decreased (Fig. 3C). These findings indicate that the nuclear translocation of Nrf2 by RGO might facilitate the transcriptional activation of ARE-dependent genes.

3.5. Regulation of HO-1 and NQO1 expression by RGO through ASK-MKK4/7-JNK and p38 MAPK signaling pathways in HepG2 cells

To analyze the underlying mechanisms responsible for RGO-induced phase II enzymes, HepG2 cells were treated with RGO for 1 h and induction of upstream kinase Akt and MAPKs including ERK, JNK and p38 MAPK was determined. As shown in Fig. 4, RGO dose-dependently stimulated phosphorylation of Akt and all three MAPKs. To further address the role of individual Akt and MAPK pathways in RGO-induced Nrf2 expression, specific kinase inhibitors (U0126 for MEK1/2, SB202190 for p38 MAPK, SP600125 for JNK, and LY294002 for Akt) were pretreated for 1 h prior to RGO treatment. The result showed that RGO-induced Nrf2 expression was markedly blocked by JNK inhibitor (SP600125) and
significantly inhibited by p38 inhibitor (SB202190), whereas U0126 and LY294002 were unlikely to diminish Nrf2 expression level (Fig. 5A). Furthermore, the RGO-mediated nuclear translocation of Nrf2 was significantly inhibited by SP600125 and SB202190, but not by U0126 or LY294002 (Fig. 5B). These results suggest that RGO-mediated Nrf2 induction is regulated through JNK and p38 MAPK signal transduction pathways. Activation of MAPKs requires tyrosine and threonine phosphorylation by activated MAPKKs (MEK1/2, SEK1/MKK4, MKK7, and MKK3/6). Among the MAPKs, the activation of JNK is catalyzed by two upstream kinases, MKK4 and MKK7 [16,34]. The activation of MKK4 and MKK7 is mediated by MAPKKKs including MLKs, ASKs, and dual leucine zipper kinases [35,36]. To further investigate the upstream kinases of JNK and p38 MAPK, we examined the regulatory role of RGO in the phosphorylation of MAPKKs (MKK3/6, MEK1/2, MKK4, and MKK7) and MAPKKKs (TAK1, B-Raf, ASK1, and MLK3). RGO-treated HepG2 cells resulted in the increased phosphorylation levels of MKK4/7 and ASK1, whereas RGO was unlikely to exhibit a significant effect on phosphorylation of MLK3, TAK1, and MKK3/6, implying that RGO directly regulated p38 MAPK phosphorylation without affecting upstream kinases including MKK3/6 and MLK3/TAK1 (Fig. 6). Our findings reveal that the RGO-induced expression of phase II enzymes could be regulated by Nrf2/ARE machinery through activation of ASK1–MKK4/7–JNK and p38 MAPK signaling transduction pathways.

4. Discussion

Consumption of natural products is one of the crucial strategies to prevent the risk of many diseases such as cardiovascular diseases, diabetes, and cancer. This could be achieved through inductive effects of chemopreventive phytochemicals on cellular defense systems, including phase II detoxifying and antioxidant enzymes to eliminate chemical carcinogens and oxidative stress. In the present study, we analyzed the chemical constituents and investigated the chemopreventive properties of RGO. The results demonstrated that RGO induces Nrf2 nuclear translocation and phase II enzymes, including HO-1 and NQO1, through activation of ASK1–MKK4/7–JNK and p38 MAPK signaling pathways in HepG2 cells. GC/MS analysis revealed that RGO contains many bioactive compounds, in which linoleic acid, β-sitosterol, and bicyclo(10.1.0) tridec-1-ene are major components accounting for about 70% of RGO. Recently, many findings have demonstrated the bioactivities...
of linoleic acid and β-sitosterol; however, no study has yet reported any biological characteristics of bicyclo(10.1.0)tridec-1-ene. Linoleic acid has been demonstrated to be an inducer of Nrf2 in mouse primary hepatocytes [37]. Our previous study showed that sitosterol might induce cellular protective systems through activation of the Nrf2/ARE pathway in HepG2 cells [38]. In addition, Zhang et al [39] have reported that phytosterins such as β-sitosterol, campesterol, and stigmasterol protected against 4-
nitrophenol-induced oxidative stress via enhancement of Nrf2-mediated detoxifying/antioxidant enzymes in rat testes. β-Sitosterol enhances antioxidant enzymes that contribute to the beneficial effects of olive oil consumption [40]. American ginseng oil may be useful as a functional ingredient since it contains various kinds of phytosterols, including squalene, campesterol, stigmasterol, sitosterol, oxidosqualene, and avenasterol, which are considered to promote human health by reducing cholesterol in humans [41].

It has been well documented that phase II enzymes could have significant effects on the prevention of tumor initiation through preventing activation of procarcinogens to reactive intermediates and increasing the neutralization of oxidative stress [32,42]. HO-1, catalyzing the conversion of heme into biliverdin, free iron, and CO [43], plays an important role in maintaining cellular redox homeostasis against oxidative stress [44]. NQO1, a detoxifying enzyme, catalyzes the two-electron reductive metabolism and detoxification of quinines and their derivatives, resulting in protection of cells from oxidative stress and cancer development [45,46]. In the field of cancer prevention, phytochemicals have been attracting a lot of interest due to their abilities to induce phase II detoxifying/antioxidant enzymes. A series of phytochemicals such as Epigallocatechin gallate (EGCG), resveratrol, sulforaphane, and quercitrin have been demonstrated as potent inducers of phase II and antioxidant enzymes, including HO-1 and NQO1 [47–50]. Nrf2 is a key transcription factor in the transcriptional induction of phase II detoxifying/antioxidant enzymes, via ARE. An in vivo study demonstrated that expression of phase II enzymes such as NQO1 and GSTA1 is markedly abrogated in nrf2-deficient mice as compared with wild-type animals [51]. Exposure of cells to Nrf2 inducers results in enhanced expression of many protective genes [49,50,52]. Therefore, activation of Nrf2 contributes to regulation of a powerful cluster of protective genes. One possible mechanism of Nrf2 activation is post-transcriptional modification of Nrf2 by protein kinases, which increase its stability and subsequent transcription activity. Protein kinases play a crucial role in converting various extracellular signals into intracellular responses through signal phosphorylation cascades. Phosphorylation of Nrf2 at serine and threonine residues by protein kinases such as PI3K/Akt, ERK, JNK, and p38 MAPK result in liberation of Nrf2 from Keap1 (Nrf2 inhibitor), enhancing Nrf2 nuclear accumulation, and finally triggering transcripational activation of phase II enzymes [78]. Phosphorylation is one of the key steps for activation of the Nrf2 pathway; however, the effect of individual protein kinases on Nrf2/ARE signaling systems may depend on cell/tissue types as well as inducers. Phenethyl isothiocyanate activates expression of Nrf2/ARE-mediated phase II enzymes through a JNK1-dependent pathway in HeLa cells [53], whereas p38 MAPK is involved in induction of Nrf2-mediated cellular defense systems by quercitin in HepG2 cells [54]. Tea catechin EGCG-induced HO-1 expression and Nrf2 nuclear accumulation have been shown through Akt and p38 MAPK signal transduction pathways [55].

In conclusion, the lipid-soluble constituents were determined in RGO with three major compounds including linoleic acid, bicyclo[10.1.0]tridec-1-ene, and β-sitosterol. Moreover, RGO potently induced expression and nuclear translocation of the transcription factor Nrf2, leading to transcriptional activation of ARE-mediated phase II detoxifying/antioxidant enzymes, possibly via ASK1–MKK4/7–JNK and p38 MAPK signaling transduction pathways in HepG2 cells. Our results suggest that RGO might be a new potential source of natural chemopreventive and cellular defensive agents.

**Conflicts of interest**

All authors have no conflicts of interest to declare.

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