

Nrf2 as a Master Redox Switch in Turning on the Cellular Signaling Involved in the Induction of Cytoprotective Genes by Some Chemopreventive Phytochemicals

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Abstract



A wide array of dietary phytochemicals have been reported to induce the expression of enzymes involved in both cellular antioxidant defenses and elimination/inactivation of electrophilic carcinogens. Induction of such cytoprotective enzymes by edible phytochemicals largely accounts for their cancer chemopreventive and chemoprotective activities. Nuclear factor-erythroid-2-related factor 2 (Nrf2) plays a crucial role in the coordinated induction of those genes encoding many stress-responsive and cytoprotective enzymes and related proteins. These include NAD(P)H:quinone oxidoreductase-1, heme oxygenase-1, glutamate cysteine ligase, glutathione S-transferase, glutathione peroxidase, thioredoxin, etc. In resting cells, Nrf2 is sequestered in the cytoplasm as an inactive complex with the repressor Kelch-like ECH-associated protein 1 (Keap1). The release of Nrf2 from its repressor is most likely to be achieved by alterations in the structure of Keap1. Keap1 contains several reactive cysteine residues that function as sensors of cellular redox changes. Oxidation or covalent modification of some of these critical cysteine thiols would stabi-

lize Nrf2, thereby facilitating nuclear accumulation of Nrf2. After translocation into nucleus, Nrf2 forms a heterodimer with other transcription factors, such as small Maf, which in turn binds to the 5'-upstream *cis*-acting regulatory sequence, termed antioxidant response elements (ARE) or electrophile response elements (EpRE), located in the promoter region of genes encoding various antioxidant and phase 2 detoxifying enzymes. Certain dietary chemopreventive agents target Keap1 by oxidizing or chemically modifying one or more of its specific cysteine thiols, thereby stabilizing Nrf2. In addition, phosphorylation of specific serine or threonine residues present in Nrf2 by upstream kinases may also facilitate the nuclear localization of Nrf2. Multiple mechanisms of Nrf2 activation by signals mediated by one or more of the upstream kinases, such as mitogen-activated protein kinases, phosphatidylinositol-3-kinase/Akt, protein kinase C, and casein kinase-2 have recently been proposed. This review highlights the cytoprotective gene expression induced by some representative dietary chemopreventive phytochemicals with the Nrf2-Keap1 system as a prime molecular target.

Chemoprevention, a Realistic Strategy to Reduce the Risk of Cancer



Cancer chemoprevention has attracted much attention as one of the most practical and realistic strategies in reducing the global burden of cancer [1]. The field of cancer chemoprevention research became prominent and focused at the beginning in the late 1960s when Dr. Lee Wattenberg conceptualized this strategy and suggested a mechanistic framework [2], [3], [4]. A broad spectrum of substances has been reported to retain chemopreventive potential, and it is noticeable that many of them were discovered

in the plant kingdom. Numerous phytochemicals derived from fruits, vegetables, grains, spices, and herbs are capable of intervening in multi-stage carcinogenesis via distinct mechanisms [5].

The journey of a population of normal cells towards malignancy apparently involves three distinct stages – initiation, promotion and progression. Tumor initiation, a rapid and irreversible process, begins with the genotoxic damage of cellular DNA upon exposure to endogenous or exogenous carcinogens. The initiation stage of chemically induced tumorigenesis involves the metabolic activation of carcinogens and subsequent

covalent modification of genomic DNA, leading to activation of oncogenes and/or inactivation of tumor suppressor genes. Tumor promotion is recognized as a reversible process characterized by clonal expansion of initiated cells to form a solid mass of proliferating preneoplastic cells. Progression, the final stage of neoplastic transformation, involves the growth of a tumor with invasive and metastatic potential [6].

The recent progress in molecular biology of cancer has identified key components of the intracellular signaling network, especially protein kinases and transcription factors, which function abnormally during the course of cellular transformation and malignancy. In response to carcinogenic insults, the micro-environment of the intracellular signaling network becomes disrupted, thereby favoring malignant transformation of cells. Therefore, one of the essential approaches currently being adopted in studying the mechanisms of chemopreventive phytochemicals is the assessment of their effects on the specific components of the signal transduction network that often becomes awry – either amplified or repressed – during carcinogenesis.

Potential of Cellular Antioxidant Defence by Edible Phytochemicals

Oxidative stress acts as a predisposing factor to multistage carcinogenesis. Reactive oxygen species (ROS), such as superoxide anion, hydrogen peroxide, hydroxyl radical, etc., are constantly generated in cells as unwanted by-products of aerobic metabolism. Although a low physiological level of ROS is scavenged efficiently by the cellular antioxidant defense system, an imbalance between the generation of ROS and the cellular antioxidant capacity turns into a state of “oxidative stress”. Oxidative stress contributes to tumorigenesis by either directly attacking DNA to cause genetic alterations or through epigenetic mechanisms that can be achieved by modulating cellular signal transduction pathways [7]. Certain environmental insults (e.g., microbial infection, ultraviolet radiation, and chemical toxicants) act as tumor initiators and/or promoters by inducing steady-state increases in the generation of ROS [7].

In response to oxidative stress, cells attempt to fortify the antioxidant arsenal as the first line of defense. The antioxidant enzymes are involved in not only deactivation of ROS, but also elimination or detoxification of electrophiles [8], [9]. Because of such dual roles, antioxidant enzymes are not unequivocally distinguishable from phase-2 detoxification enzymes, and *vice versa* [10]. Although antioxidant or detoxifying enzymes are rapidly induced in response to oxidative or electrophilic stress, such adaptive survival response is normally transient and prone to be overwhelmed by excess amount of ROS or electrophiles. In this context, we may need to fortify or adequately maintain the cellular cytoprotective mechanism through dietary as well as pharmaceutical manipulations. A wide variety of dietary polyphenols and other classes of phytochemicals have been reported to induce the expression of enzymes involved in both cellular antioxidant defenses and elimination/inactivation of electrophilic carcinogens [10]. Induction of such cytoprotective enzymes by edible phytochemicals is recognized as one of the highly effective strategies for preventing cancer in the human population [11].

Nrf2 as a Key Player in Intracellular Signal Transduction Leading to Cytoprotective Enzyme Induction

Nuclear factor-erythroid-2-related factor 2 (Nrf2), a member of the NF-E2 family of the basic leucine zipper transcription factors, is essential for the coordinated induction of those genes encoding many stress-responsive or cytoprotective enzymes and related proteins, such as NAD(P)H:quinone oxidoreductase-1 (NQO1), superoxide dismutase (SOD), glutathione S-transferase (GST), glutathione peroxidase (GPx), heme oxygenase-1 (HO-1), glutamate cysteine ligase (GCL), catalase, and thioredoxin [8], [9], [10], [11], [12].

The *Nrf2*-deficient mice failed to induce genes responsible for carcinogen detoxification and protection against oxidative stress [3], [14], [15]. Thus, there was a significantly higher burden of benzo[*a*]pyrene (B[a]P)-induced gastric neoplasia in *Nrf2*-deficient mice, which were less responsive to the chemopreventive agent oltipraz [16]. A naturally occurring chemopreventive agent, sulforaphane, reduced B[a]P-induced forestomach tumorigenesis in ICR mice, most likely via the induction of phase 2 detoxification/antioxidant enzymes, as this protective effect was abrogated in *Nrf2*-null mice [17]. Besides its role in regulating carcinogen detoxification and cellular antioxidative defense, Nrf2 also has anti-inflammatory functions [18], [19] and thus represents a novel therapeutic approach for the treatment or prevention of inflammatory disorders [20]. In a recent study, Khor et al. reported that the aggravation of dextran sulfate sodium-induced colitis in *Nrf2*^{-/-} mice was associated with decreased expression of HO-1, NQO-1, UDP-glucuronyltransferase 1A1 (UGT1A1), and GST μ -1 [21]. In addition, levels of pro-inflammatory mediators, such as cyclooxygenase-2, inducible nitric oxide synthase, interleukin (IL)-1 β , IL-6, and tumor necrosis factor alpha (TNF- α), were significantly elevated in the colonic tissues of *Nrf2*^{-/-} mice as compared to their wild-type counterparts [21]. Therefore, targeted activation of the Nrf2 signaling accounts for the chemopreventive effects elicited by some dietary phytochemicals capable of suppressing oxidative and/or inflammatory stress.

In resting cells, Nrf2 resides in the cytoplasm by forming an inactive complex with the repressor Kelch-like ECH-associated protein 1 (Keap1), which is anchored to the actin cytoskeleton. This tight interaction presents Nrf2 for ubiquitination followed by proteasomal degradation. Keap1 associates with Cullin 3 (Cul3) and Rbx1 to form a functional E3 ubiquitin ligase complex that targets multiple lysine residues located in the N-terminal Neh2 domain of Nrf2 for ubiquitination. So, Keap1 functions as a substrate adaptor protein for a Cul3-dependent E3 ubiquitin ligase complex and thereby regulates steady-state levels of Nrf2 [22], [23]. Under oxidative or electrophilic stress, Nrf2 is stabilized and translocated to the nucleus, where it transactivates ARE-regulated genes [9], [11]. Widely accepted mechanisms underlying nuclear translocation of Nrf2 and subsequent transactivation of ARE-regulated genes include (a) stabilization of cytoplasmic Nrf2 through blockade of its ubiquitination and proteasomal degradation by Keap1-Cul3 complex [24], and (b) dissociation of Nrf2 from Keap1 via phosphorylation of serine [25] or threonine [26] residues of Nrf2 (● Fig. 1). After migration to the nucleus, Nrf2 undergoes heterodimeric combinations with other transcription factors, such as small Maf protein and binds to the 5'-upstream *cis*-acting regulatory sequence, termed antioxidant response elements (ARE) or electrophile response elements

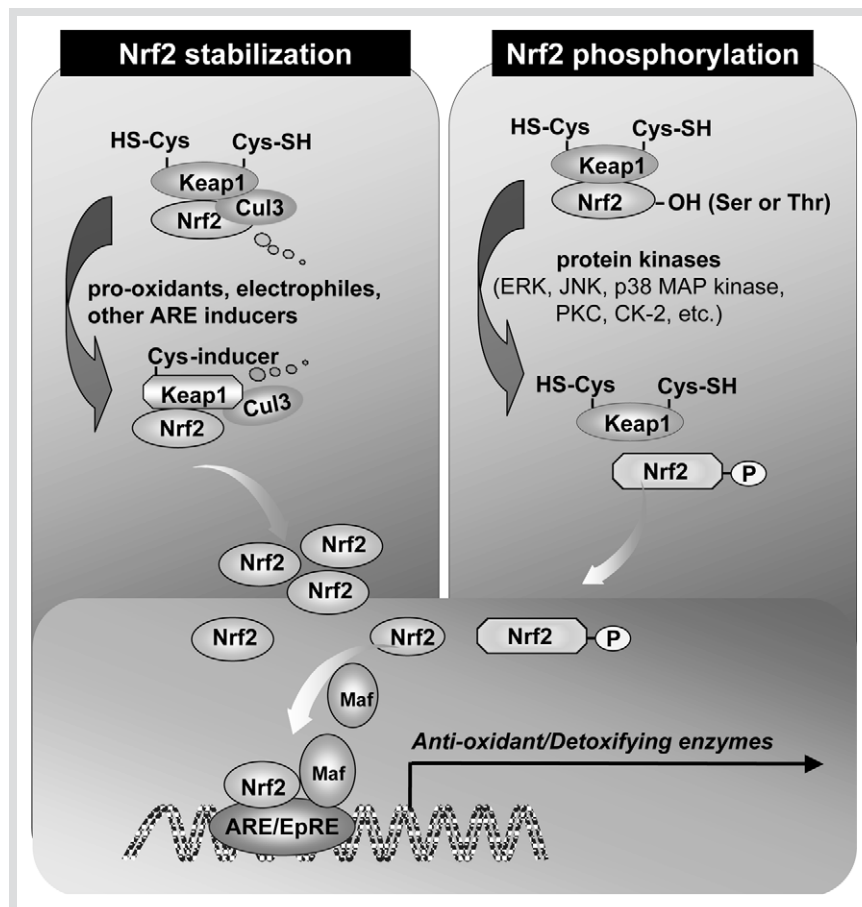


Fig. 1 Proposed mechanisms for Nrf2-ARE-mediated signaling. Nrf2 can be activated by at least two mechanisms that include: (i) stabilization of Nrf2 via Keap1 cysteine thiol modification and (ii) phosphorylation of Nrf2 by upstream kinases. Under physiological conditions, Nrf2 is sequestered in the cytoplasm as an inactive complex with Keap1. (*left*) Keap1 is a substrate adaptor protein for Cul3-dependent ubiquitin ligase complex. Under basal conditions, Keap1 targets Nrf2 for ubiquitin-dependent degradation and represses Nrf2-dependent gene expression. Some ARE inducers (e.g., pro-oxidants, electrophiles and phase-2 enzyme inducing chemopreventives) can modify critical cysteine residue(s) (e.g., Cys151) of Keap1, which results in a conformational change in Keap1 and the inability of Cul3 to ubiquitinate Nrf2. Covalently modified or oxidized Keap1 is then polyubiquitinated by Cul3 and undergoes proteasomal degradation, thereby allowing Nrf2 to get stabilized and accumulated in nucleus. (*right*) Alternatively, activation of protein kinases, such as PKC, JNK, ERK and CK2 induces Nrf2 phosphorylation, which may stimulate the dissociation of Nrf2 from its repressor Keap1 and subsequent translocation into nucleus. Phosphorylation of Nrf2 is also considered to facilitate its interaction with the transcriptional coactivator CBP/p300 and recruitment of components of transcription initiation machinery.

(EpRE), located in the promoter region of genes encoding various antioxidant/detoxifying enzymes [29].

Certain dietary chemopreventive agents target Keap1 by oxidizing or chemically modifying its specific cysteine thiols [5], [9], [10], [11], [30], [31], [32]. The oxidation of Keap1 cysteine thiols can be mediated by some edible polyphenols. In this context, it is noticeable that among flavonoids, the higher their intrinsic potential to generate oxidative stress and redox cycling, the stronger their potency as inducers of ARE-mediated gene expression [33]. Polyphenols can undergo oxidation either enzymatically or spontaneously to form quinoids, which can be reduced back to parental molecules, and during such redox process ROS can be generated. For instance, the green tea polyphenol (-)-epigallocatechin gallate (EGCG) was found to produce substantial amount of H_2O_2 under cell culture conditions [34], [35]. Therefore, it is paradoxical that the activation of Nrf2-ARE signaling by antioxidant polyphenols to induce cytoprotective enzymes is attributable, at least in part, to their prooxidant activity.

Keap1 is a cysteine-rich homodimeric, multi-domain zinc metalloprotein [36]. Some inducers of cytoprotective enzymes or their reactive metabolites/intermediates can directly interact with critical sensor cysteine residues of Keap1, which has been proposed to diminish the affinity of Nrf2 for Keap1, stimulating their dissociation. Many polyphenols undergo oxidation to produce reactive quinone derivatives that are electrophilic Michael reaction acceptors. It has been suggested that some quinone derivatives of polyphenol inducers can oxidize two highly reactive cysteine thiol groups of Keap1, resulting in disulfide bond formation [37]. This possibility was corroborated by the linear correlation between the antioxidant enzyme-inducing potencies of di-

phenols and the extent of their redox potential in terms of ability to release electrons [37]. Other types of phytochemicals with Michael acceptor functionalities directly modify a critical cysteine thiol of Keap1 without undergoing redox cycling. Of particular examples are sulforaphane and structurally related isothiocyanates [38].

The dissociation of Nrf2 from Keap1 as a consequence of Keap1 cysteine thiol modification by ARE inducers used to be considered as a plausible mechanism underlying Nrf2 activation by electrophiles and prooxidants, but this supposition has recently been challenged by some investigators [24]. According to the new paradigm, the direct interaction of the highly reactive cysteine residues of Keap1 with phase 2 enzyme inducers as well as electrophiles causes conformational changes of this repressor protein, which abrogate the capability of Keap1 to aid proteasomal degradation of Nrf2. Thus, modification of the cysteine 151 residue of human Keap1 by sulforaphane failed to directly release Nrf2 from Keap1, but rather induced conformational changes of Keap1 followed by its proteasomal degradation through polyubiquitination, thereby stabilizing Nrf2 [24]. Besides the C151 residue, other cysteine residues of Keap1 (e.g., C273 and C288) have also been reported to be essential for the Keap1-dependent repression of Nrf2 [39]. Detailed mechanisms underlying the regulation of Nrf2 by Keap1-Cul3 complex in intact cells as well as in those under stress have been reviewed elsewhere [11], [40], [41] and are beyond the scope of this article. In addition to activation of Nrf2 through oxidation or chemical modification of Keap1 cysteine thiols, phosphorylation of specific serine [27] or threonine [26] residues of Nrf2 may facilitate the release of Nrf2 from the Keap1 repression and its subsequent nu-

clear translocation and interaction with the co-activator CBP/p300 (● Fig. 1). Alternatively, phosphorylation of the tyrosine 141 residue located in the bric-a-brac, tramtrack, broad complex (BTB) domain makes Keap1 stable, while dephosphorylation at this residue promotes rapid degradation of Keap1 and hence stabilization of Nrf2 [42]. Another novel mechanism of Nrf2 stabilization has recently been proposed by Clements et al. [43]. According to this study, DJ-1/PARK7 (a cancer and Parkinson's disease-associated protein) inhibits Nrf2-Keap1 binding and is indispensable for Nrf2 stabilization and subsequent NQO1 induction [43]. Activation of several upstream kinases, such as mitogen-activated protein (MAP) kinases [25], phosphatidylinositol 3-kinase (PI3K)/Akt [44], protein kinase C (PKC) [27], [45] and casein kinase-2 (CK-2) [46] have been considered to facilitate nuclear translocation and transcriptional activation of Nrf2. On the other hand, glycogen synthase kinase-3 β (GSK3 β) negatively regulates Nrf2 signaling via phosphorylation of Nrf2 at tyrosine [47] or serine [48] residues. Several ARE inducers have been shown to phosphorylate Akt and enhance nuclear translocation of Nrf2 [49], [50], [51], [52], [53], which may result from the inactivation of GSK3 β by activated Akt. Nrf2 activation by dietary chemopreventives through modulation of one or more of the upstream kinases have recently been reviewed [8], [9], [12], [54]. ● Table 1 lists several representative dietary phytochemicals (structures shown in ● Fig. 2) that have the ability to activate Nrf2-ARE signaling. Additional information on chemopreventive phytochemicals targeting the Nrf2-Keap1 system is available in several reviews [5], [7], [8], [9], [10], [11] as well as in the following section.

Chemopreventive Phytochemicals with Nrf2-ARE Activating Ability

▼ Sulforaphane

Sulforaphane [1-isothiocyanato-(4*R,S*)-(methylsulfinyl)butane], a representative isothiocyanate present in broccoli and other cruciferous vegetables, exerts its cancer chemopreventive and cytoprotective effects mainly by the induction of genes encoding phase 2 detoxifying and antioxidant enzymes [55]. The gastrointestinal GPx is a selenoprotein that was suggested to act as a barrier against hydroperoxide absorption but has also been implicated in the control of inflammation and malignant growth [56]. The anticarcinogenic role of GI-GPx is evident from enhanced gastrointestinal tumor formation in *gpx2/gpx1* double knockout mice [57]. The GI-GPx promoter harbours an ARE binding site that was activated by sulforaphane or *tert*-butylhydroquinone (tBHQ) as well as through overexpression of Nrf2 in human hepatoma (HepG2) or human colon cancer (Caco-2) cells [56]. Thus, the antioxidant and anti-inflammatory as well as chemopreventive effects of sulforaphane may be attributable to upregulation of GI-GPx as well as induction of conventional phase 2 detoxifying enzymes.

Sulforaphane, by activating Nrf2 signaling, abrogated toxicity as well as the accumulation of arsenite in cultured murine hepatocytes [58]. Using a human bladder urothelial cell line stably transfected with Nrf2-siRNA, Wang and colleagues [59] demonstrated that compromised Nrf2 expression sensitized the cells to As(III)-induced toxicity. On the other hand, the activation of the Nrf2 pathway by sulforaphane or the known Nrf2-inducer tBHQ rendered these cells more resistant to As(III). Sulforaphane upregulated the expression of detoxifying enzymes including NQO1,

GST and GCL in the small intestine of *nrf2*-wild-type mice, while the *nrf2*-null mice displayed lower levels of these enzymes [60]. Livers obtained from the *nrf2* wild-type and the *nrf2* knockout mice after treatment with vehicle or sulforaphane for 3 and 12 h exhibited differential gene expression profiles determined by the Affymetrix 39K oligonucleotide microarray [61]. In this study, the Nrf2-dependent, sulforaphane-inducible genes were identified. These include phase 1/2 xenobiotic metabolizing enzymes and phase 3 transporters. Unexpected clusters of genes include those for heat shock proteins, ubiquitin/26S proteasome subunits, and lipid metabolism. Topical application of sulforaphane (100 nmol) for 14 consecutive days inhibited skin carcinogenesis induced by 7,12-dimethylbenz[*a*]anthracene plus 12-*O*-tetradecanoylphorbol 13-acetate in C57BL/6 mice, whilst no such chemopreventive effects were elicited in the *nrf2*-deficient mice [62].

Sulforaphane induces Nrf2-driven phase 2 enzyme expression by modulating the activation of MAP kinases [54], [63], [64], [65], [66], [67]. Sulforaphane and its sulfide analogue, erucin elevated the mRNA expression of NQO1, UGT1A1, and multidrug transporter protein (MRP) 2 in Caco-2 cells by activating PI3K/Akt- or MAP kinase kinase (MEK)/extracellular signal-regulated protein kinase (ERK)-mediated signaling [65]. Involvement of ERK1/2 and JNK in sulforaphane-induced ARE-transcription activities was also observed in murine keratinocytes [66]. Unlike ERK and JNK, p38 MAP kinase appeared to negatively regulate Nrf2 activation [67]. The p38 MAP kinase phosphorylated purified Nrf2 protein at a site presumably different from those phosphorylated by other kinases [67]. Overexpression of p38 MAP kinase isoforms caused an increase in the interaction between Nrf2 and Keap1 and attenuated constitutive and inducible Nrf2 translocation into nucleus [67]. However, the role of p38 MAP kinase in Nrf2 activation is still controversial. Sulforaphane induced HO-1 expression in HepG2 cells by downregulating p38 MAP kinase, thereby activating the Nrf2-ARE signaling [67]. The 26S proteasome is responsible for degradation of abnormal proteins and may play a role in cell survival upon oxidative stress. Sulforaphane enhanced the expression of the catalytic subunits of the proteasome as well as proteasomal peptidase activities in murine neuroblastoma Neuro2A cells [68]. Sulforaphane treatment protected these cells from hydrogen peroxide-mediated cytotoxicity in a manner dependent on proteasomal function. In addition to the modulation of upstream kinases, the mechanism of Nrf2 activation by sulforaphane involves a direct modification of cysteine residue(s) on Keap1, facilitating dissociation of Nrf2 from Keap1 [38]. As sulforaphane is an electrophile, it can react with protein thiols to form thionoacyl adducts. Although sulforaphane activates Nrf2-ARE signaling through specific modifications of the Keap1 protein, this isothiocyanate displays a distinctly different pattern of Keap1 modification compared with conventional ARE inducers that modify Keap1 by alkylation. Thus, sulforaphane modifies Keap1 predominantly in the Kelch domain, rather than in the central linker domain, which is targeted by previously characterized ARE inducers [38]. Moreover, sulforaphane treatment *in vivo* did not lead to the accumulation of ubiquitinated Keap1, which was consistent with the previous finding that sulforaphane inhibits Keap1-dependent ubiquitination of Nrf2 without inducing Keap1 ubiquitination [23]. A single cysteine-to-serine substitution (C151S) at residue 151 within the BTB domain of Keap1 conferred significant resistance to inhibition by sulforaphane [22]. Treatment of cells with sulforaphane decreased the level of Cul3 that was co-purified

Table 1 Cellular antioxidant defence through activation of the Nrf2 signaling by chemopreventive phytochemicals

Compound (Dose)	Molecular mechanisms	Experimental systems	Ref.
Sulforaphane			
(2–200 μ M)	\uparrow Stabilization and nuclear accumulation of Nrf2 via cysteine thioacetylation of Keap-1	Human Keap-1-transfected HEK293 cells and in vitro study	[38]
(20 μ M)	\downarrow Phosphorylation of p38 MAP kinase, \uparrow Nrf2-ARE activity, \uparrow HO-1 expression	HepG2 cells	[67]
(9 μ mol/mouse per day for 7 days)	\uparrow mRNA expression of NQO1, GCL, GST and glutathione reductase, \uparrow activities of NQO1, GST, glucose-6-phosphate dehydrogenase, UDP-glucose dehydrogenase, maleic enzyme, and carboxylesterase	<i>nrf2</i> -wild type mice	[60]
(3 μ mol/g for 14 days)	\uparrow Expression and activities of NQO1 and GST, \uparrow expression of GCS	<i>nrf2</i> -wild type mice	[119]
Curcumin			
(0.05% in diet)	\uparrow Activity, mRNA and protein expression of GST and NQO1 in liver and lungs, \uparrow expression, nuclear translocation and ARE binding of Nrf2 in liver and lungs, \uparrow B[a]P detoxification	Inbred Swiss albino mice treated with B[a]P	[73]
(20 μ M)	\uparrow GSTP-1 mRNA expression, \uparrow Nrf2-ARE-regulated <i>GSTP-1</i> reporter gene activity	HepG2 cells	[76]
(15 μ M)	\uparrow ARE-mediated expression of HO-1 and GCLM, \uparrow PKC δ , \uparrow p38 MAP kinase, \uparrow Nrf2	Human monocytes	[77]
(10–30 μ M)	\uparrow Phosphorylation of p38 MAP kinase, \uparrow dissociation of Nrf2-Keap1, \uparrow Nrf2 binding to <i>ho-1</i> -ARE, and \uparrow expression and activity of HO-1	NRK-52E cells and LLC-PK ₁ cells	[74]
(10–15 μ M)	\uparrow Nuclear translocation and DNA binding of Nrf2-ARE, \uparrow GCLC and GCLM mRNA and protein expression, \uparrow GSH	HBE1 cells	[75]
(10 μ M)	\uparrow Activation of PKC and p38 MAP kinase, \uparrow Nrf2 activation, and \uparrow expression of HO-1	HepG2 cells	[78]
(20 μ M)	\uparrow HO-1 expression, \uparrow Nrf2 nuclear translocation, \uparrow ARE luciferase activity	rat vascular smooth muscle cells	[80]
(200 mg/kg)	\uparrow DNA binding of Nrf2-ARE in liver, \uparrow HO-1 expression and activity in liver, \downarrow dimethylnitrosoamine-induced hepatotoxicity	Curcumin given by gavage to male Albino rats	[79]
EGCG			
(25 μ M)	\uparrow MAP kinases, \uparrow ARE luciferase activity	HepG2 cells	[85]
(20 μ M)	\uparrow HO-1 expression, \uparrow Nrf2 nuclear translocation, \uparrow Akt, \uparrow p38 MAP kinase	B lymphoblasts	[49]
(50 μ M)	\uparrow HO-1 expression, \uparrow Nrf2 nuclear level, \uparrow ARE luciferase activity, \uparrow ERK, \uparrow Akt	Bovine aortic endothelial cells	[50]
(100 μ M)	\uparrow Nuclear localization of Nrf2, \uparrow Nrf2-ARE DNA binding, \uparrow Akt, \uparrow ERK, \uparrow mRNA and protein expression of GCL, HO-1 and MnSOD	MCF-10A cells	[51]
(20 mg/kg; gavage)	\downarrow Growth of implanted colon tumor in nude mice, \uparrow protein expression of Nrf2, and mRNA expression of UGT1A, UGT1A8, UGT1A10 and Nrf2 in implanted colon tumors	Colon tumors orthotopically implanted in cecum of nude mice	[86]
10 or 20 mg/kg	\uparrow Expression of UGT1A10 and Nrf2 protein and mRNA, \downarrow formation of aberrant crypt foci	Balb/c nude mice treated with or without 2-amino-3-methylimidazo[4,5-f]quinoline	[121]
Diallyl sulfide			
(1 mM)	\uparrow HO-1 mRNA and protein expression, \uparrow Nrf2 nuclear translocation, \uparrow Nrf2 DNA binding activity, \uparrow ERK and p38, protected cells from hydrogen peroxide-induced toxicity	HepG2 cells	[89]
Diallyl trisulfide			
(100 μ M)	\uparrow HO-1 and NQO-1 gene expression, \uparrow Nrf2 protein level and ARE activity via activation of calcium-dependent signaling	HepG2 cells	[90]
S-Allylcysteine			
100 mg/kg	\uparrow Activities of SOD, catalase, GPx, GST and NQO1, \uparrow nuclear translocation of Nrf2, \downarrow chromium-induced cell death	K ₂ Cr ₂ O ₇ -treated rat liver	[120]
Resveratrol			
(15 μ M)	\uparrow Expression and activity of HO-1 via activation of Nrf2	PC12 cells	[99]
(10 μ M)	\uparrow GCL activity, \uparrow GSH synthesis, \downarrow CSE-induced posttranscriptional (aldehyde/carbonyl adduct or nitration) modification of Nrf2 and Keap1, \uparrow Nrf2 nuclear translocation	Human small airway epithelial cells and A549 cells	[100]

Compound (Dose)	Molecular mechanisms	Experimental systems	Ref.
(25–50 μ M)	\uparrow Expression of NQO1 protein and mRNA, \uparrow NQO-1 activity, \uparrow nqo-1-ARE activity, \uparrow phosphorylation and nuclear translocation of Nrf2	K562 cells	[97]
Lycopene (2–4 μ M)	\uparrow GCS-ARE and NQO1-ARE activity, \uparrow mRNA and/or protein expression of GCS and NQO-1, \uparrow glutathione, \downarrow ROS	MCF-7 and HepG2 cells	[105]
Capsaicin (200 μ M)	\uparrow ROS, PI3K/Akt expression, \uparrow HO-1 protein and mRNA expression, \downarrow HO-1 activity, \downarrow NQO1 expression and activity, \uparrow activation of Nrf2	HepG2 cells	[52]
Piperine (50 μ M)	\uparrow Phosphorylation of MAP kinases, \uparrow JNK-mediated HO-1 expression, \uparrow Nrf2 nuclear translocation	Mouse auditory House Ear Institute organ of Corti-1 (HEI-OC1) cells	[122]
Carnosol (10 μ M)	\uparrow HO-1 mRNA and protein expression, \uparrow binding of Nrf2 to ho-1-ARE, \uparrow activation of ERK, p38 MAP kinase and JNK	PC12 cells	[110]
Cinnamaldehyde (100 μ M)	\uparrow Expression of HO-1, \uparrow nuclear translocation of Nrf2, \uparrow ho-1-ARE reporter activity	Human endothelial cells	[111]
Zerumbone (1–25 μ M)	\uparrow Expression of GSTP-1, GCL, GPx and HO-1, \uparrow nuclear translocation of Nrf2	RL34 cells	[113]
Cafestol (0.025% diet)	\uparrow Expression and activities of NQO-1 and GST in cytosolic fraction of small intestinal tissue	<i>nrf2</i> wild type mice	[119]
Kahweol	\uparrow Expression of HO-1 mRNA and protein, \uparrow HO-1 activity, \uparrow phosphorylation of Akt and p38 MAP kinase, \uparrow nuclear translocation of Nrf2	SH-SY5Y cells	[123]
Chalcone (25 μ M)	\uparrow HO-1 expression, \uparrow nuclear Nrf2 level, \uparrow ARE luciferase activity, \uparrow thioredoxin reductase promoter activity	Endothelial cells	[114]
Xanthohumol (4 μ M)	\uparrow NQO1 expression, \uparrow activation of Nrf2 via covalent modification of Keap-1 at cysteine 27 residue	Murine Hepa1c1c7 cells	[118]
Eupatilin (150 μ M)	\uparrow Phosphorylation of ERK, \uparrow nuclear translocation of Nrf2, \uparrow HO-1 expression	Feline ileal smooth muscle cells	[124]
Isoorientin (5 μ g/mL)	\uparrow Phosphorylation of PI3K/Akt, \uparrow nuclear accumulation of Nrf2, \uparrow expression of HO-1, NQO-1 and peroxiredoxin	HepG2 cells	[125]
Quercetin 50 μ M	\uparrow HO-1 expression, \uparrow nuclear level of Nrf2, \downarrow ROS generation, \downarrow H ₂ O ₂ - or <i>t</i> -butyl hydroperoxide-induced cell death	Adult human retinal pigment epithelial cells	[126]
10–40 μ M	\uparrow NQO-1 mRNA expression, \uparrow NQO1-ARE DNA binding, \uparrow nuclear translocation of Nrf2, \downarrow Nrf2 ubiquitination, \uparrow <i>nrf2</i> -mediated ARE activity	HepG2 cells	[127]
100 μ M	\uparrow GSH level, \downarrow lipid peroxidation \uparrow HO-1 activity, \uparrow nuclear Nrf2 expression	Ethanol-stimulated human hepatocytes isolated from liver cancer patients	[128]
25 μ M	\uparrow HO-1-ARE and Gl-GPx-ARE activity, \uparrow Gl-GPx promoter activity, \uparrow nuclear Nrf2 expression	HepG2 cells	[129]
3-O-Caffeoyl-1-methylquinic acid (20–200 μ M)	\uparrow mRNA expression of HO-1, GCL, GR, and GST, \uparrow HO-1 protein expression, \uparrow phosphorylation of JNK and ERK, \uparrow nuclear translocation of Nrf2	Human umbilical vein endothelial cells	[130]
Brazilin (10 mM)	\uparrow Phosphorylation of Akt and ERK, \uparrow expression of HO-1 mRNA and protein, \uparrow nuclear translocation of Nrf2	HEI-OC1 cells	[131]
Chlorophyllin (50 μ M)	\uparrow Phosphorylation of PI3K/Akt, \uparrow nuclear accumulation of Nrf2, \uparrow expression of HO-1 and NQO1	Human umbilical vein endothelial cells	[132]

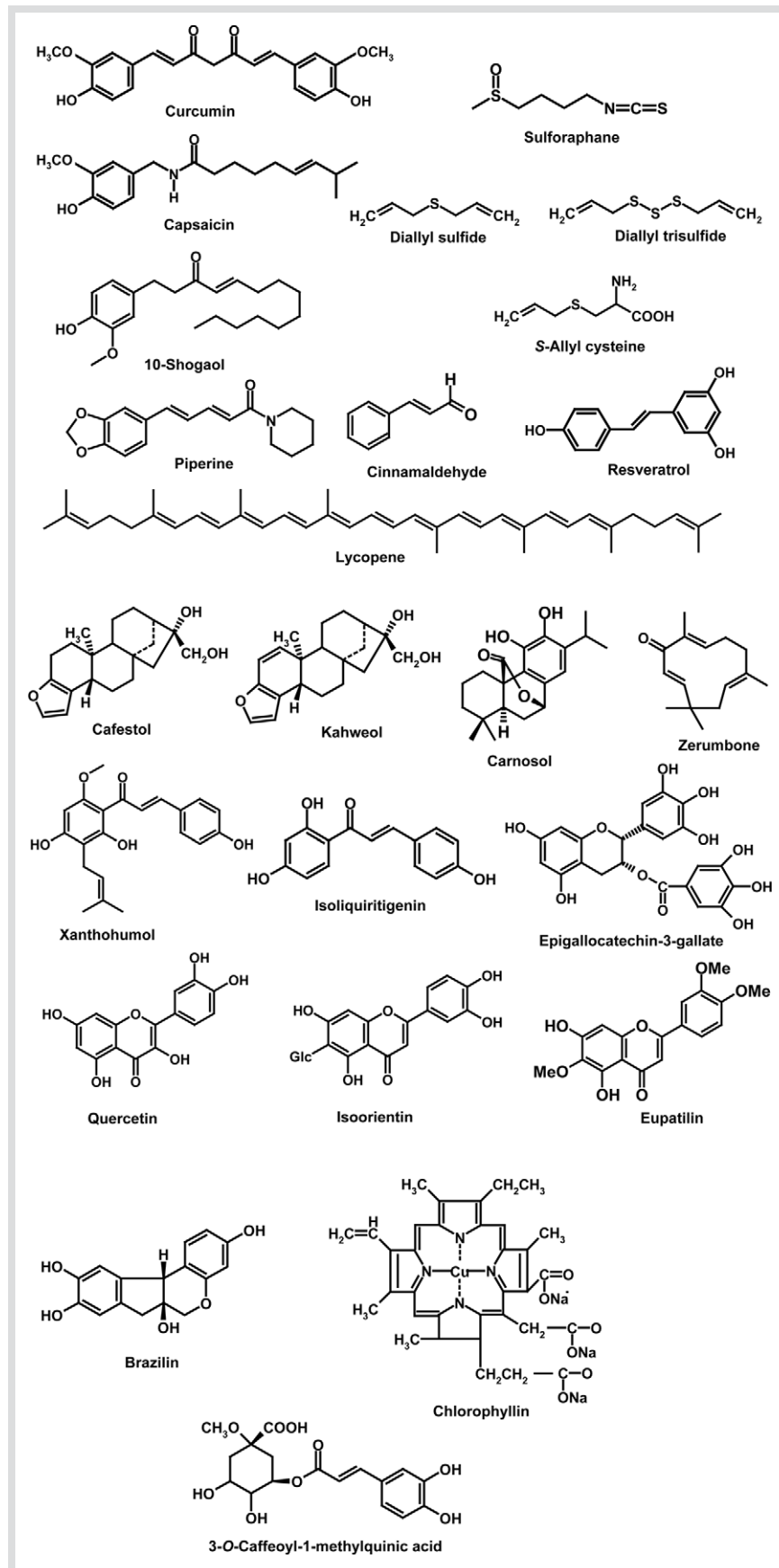


Fig. 2 Selected chemopreventive phytochemicals that can activate Nrf2-ARE signaling.

with the wild-type Keap1-CBD protein following affinity purification. Importantly, the amount of Cul3 that was co-purified with the Keap1-C151S protein was greater than the amount of

Cul3 that was co-purified with the wild-type Keap1 protein following treatment with sulforaphane [22]. Although a series of studies reported that sulforaphane activated Nrf2 signaling

largely by cysteine thiol modification and subsequent dissociation of Nrf2 from Keap1, a recent study by Egger and colleagues [24] demonstrated that thiol modification of Keap1 cysteine 151 by sulforaphane failed to cause direct dissociation of Nrf2 from Keap1. The study, rather, provided a new insight into the mechanism of Nrf2 activation by sulforaphane. According to this study, the covalent modification of Keap1 cysteine 151 by sulforaphane leads to structural changes of Keap1, which then undergoes polyubiquitination and proteasomal degradation, thereby allowing Nrf2 to escape from Cul3-dependent proteasomal degradation [24]. A sulforaphane analogue, 6-methylsulphonylhexyl isothiocyanate isolated from Japanese wasabi was also found to induce cytoprotective gene expression via the Nrf2-ARE signaling [69]. Likewise, phenethyl isothiocyanate induced HO-1 expression and ARE activity in human prostate cancer (PC3) cells via ERK and JNK-mediated phosphorylation of Nrf2 and subsequent Nrf2 nuclear translocation [25].

Curcuminoids

Curcumin is a diferuloylmethane derived from the rhizomes of turmeric (*Curcuma longa* Linn, Zingiberaceae). Its chemopreventive effects have been extensively investigated and well defined [70], [71]. Curcumin targets the Nrf2-ARE signaling pathway to induce phase 2 detoxifying enzymes. Dietary curcumin enhanced not only the Nrf2 levels but also its nuclear translocation and the ARE binding in liver and lung of mice. Wild-type C57BL/6J and C57BL/6J/*nrf2*(-/-) mice were given a single oral dose of curcumin (1,000 mg/kg) and liver and small intestine were collected for DNA microarray analysis. Among those well-defined genes, 822 (664 induced and 158 suppressed) and 222 (154 induced and 68 suppressed) were curcumin-regulated Nrf2-dependent genes identified in the liver and small intestine, respectively [72]. As expected, many phase 2 detoxification/antioxidant enzyme genes were among the identified genes. Induction of carcinogen detoxifying enzymes, such as GST isoforms and NQO1 by dietary curcumin in mice paralleled the curcumin-mediated activation of Nrf2, leading to increased detoxification of benzo[a]pyrene [73].

Balogun et al. [74] reported that curcumin disrupted the Nrf2-Keap1 complex, leading to increased Nrf2 binding to ARE and subsequent increases in the expression and the activity of HO-1 in porcine renal epithelial proximal tubule (LLC-PK₁) cells and/or rat kidney epithelial (NRK-52E) cells via activation of p38 MAP kinase. Curcumin also enhanced the expression of GCL at mRNA and protein levels in immortalized human bronchial epithelial (HBE1) cells by promoting nuclear translocation and DNA binding of Nrf2 [75]. The treatment of HepG2 cells with curcumin resulted in the elevation of GSTP-1 mRNA and Nrf2-ARE-regulated GSTP1 reporter gene activity [76]. Curcumin activated ARE-mediated expression of HO-1 and GCLM in human monocytes via activation of PKC delta, an enzyme upstream of p38 MAP kinase and Nrf2 [77]. Curcumin induced HO-1 expression by activating Nrf2 via ROS generation, activation of PKC and p38 MAP kinase, and the inhibition of phosphatase activity in human hepatoma (HUH-7) cells [78]. ROS production appeared to be mitochondrial in origin, as curcumin treatment resulted in depolarization of the mitochondrial membrane potential. Both HO-1 induction and Nrf2 activation were ROS-dependent. In another study, curcumin induced HO-1 expression and nuclear translocation of Nrf2 in cultured MDA-MB468 breast cancer cells [49]. The compound also induced nuclear localization of Nrf2 and HO-1 expression effectively in wild-type mouse em-

bryonic fibroblasts, but not in those from *nrf2*-deficient mice [49].

Oral administration of curcumin at 200 mg/kg body weight for four consecutive days resulted in enhanced nuclear accumulation and the ARE-binding of Nrf2 and HO-1 upregulation in rat liver [79]. The induction of HO-1 by curcumin accounted for its protection against dimethylnitrosamine-induced hepatotoxicity in rats. Structurally, curcumin has two α,β -unsaturated carbonyl groups and can hence act as a Michael reaction acceptor, causing thiol modification of Keap1, followed by Nrf2 nuclear translocation. Consistent with this notion, tetrahydrocurcumin, which lacks an electrophilic α,β -unsaturated carbonyl functional moiety, failed to induce Nrf2-ARE binding as well as HO-1 induction when given orally to rats [79]. Pae et al. reported that curcumin inhibited serum- and TNF α -induced growth of rat vascular smooth muscle cells (rVSMC) and human aortic smooth muscle cells, respectively by activating Nrf2-ARE-mediated induction of HO-1 and increasing the expression of p21, while its hydrogenated analogue tetrahydrocurcumin did not elicit such growth inhibitory effects [80]. Moreover, co-treatment of rVSMC cells with curcumin and the HO-1 inhibitor tin protoporphyrin partially abolished curcumin-induced p21^{WAF1/CIP1} expression and growth inhibition, suggesting that the antiproliferative effect of curcumin is mediated via upregulation of HO-1 [80]. Interestingly, curcumin accelerated the nuclear translocation of Nrf2 via activation of PI3K and p38 MAP kinase in vascular smooth muscle cells, which was associated with increased expression and the promoter activity of aldose reductase [81]. Demethoxy- and bis-demethoxycurcumin induced HO-1 promoter activity more effectively than did curcumin in a mouse pancreatic β -cell line [53]. The induction was dependent on the presence of ARE sites containing enhancer regions (E1 and E2) in the HO-1 promoter as well as nuclear translocation of Nrf2. Real-time quantitative RT-PCR analysis showed significant elevation in the mRNA levels of two other phase 2 enzymes, GCLM and NQO1 as well [53]. Treatment of HepG2 cells harbouring GI-GPx-promoter constructs with curcumin resulted in a significant induction of the reporter gene activity [56]. Analysis of the ability of a series of curcuminoids to induce quinone reductase (QR) activity in Hepa1c7 cells revealed that the introduction of aromatic *o*-hydroxy groups into the curcuminoid skeleton raises the inducer potency more than 30-fold [82]. According to this study, demethoxycurcumin and bis-demethoxycurcumin exhibited almost similar capability to induce QR activity as compared to that observed with curcumin [82]. However, demethoxycurcumin and bis-demethoxycurcumin induced HO-1, GCL and NQO-1 mRNA expression and HO-1 promoter activity, and activated Nrf2 more effectively than curcumin in mouse pancreatic-beta (MIN6) cells [53].

Epigallocatechin gallate (EGCG)

EGCG, the major active catechin component of green tea, has been known to possess anti-oxidant, anti-inflammatory and chemopreventive properties [83], [84]. EGCG was found to be the most potent Nrf2 activator among the green tea polyphenols, as evidenced by its pronounced ability to induce ARE-luciferase reporter gene transactivation [85]. EGCG has been reported to activate Nrf2 and induce expression of HO-1 in endothelial cells [50] and B-lymphoblasts [49]. While EGCG-induced HO-1 expression was attributed to activation of Akt and ERK1/2 in endothelial cells [50], p38 MAP kinase as well as Akt is involved in HO-1 induction and Nrf2 nuclear translocation in B lymphoblasts treated with EGCG [49]. We also observed that EGCG induced ac-

tivation of ERK1/2 and Akt through phosphorylation in cultured human mammary epithelial MCF-10A cells. The pharmacological inhibitors of MEK and PI3K, which are upstream kinases responsible for phosphorylation of ERK1/2 and Akt, respectively, attenuated the nuclear localization of Nrf2 induced by EGCG [51]. Similarly, a nontoxic dose of EGCG increased the ARE-luciferase activity and the expression of ARE-regulated genes in HepG2 cells by activating MAP kinases [85].

EGCG inhibited the growth and liver/pulmonary metastasis of colon tumor implanted orthotopically in the cecum of nude mice, and this anticancer effect was proposed to be partly mediated by activating the Nrf2-UGT1A signal pathway [86]. C57BL/6J and C57BL/6J/*nrf2* (−/−) mice were given an oral dose of EGCG at 200 mg/kg or treated with vehicle. Both liver and small intestine were collected 3 h and 12 h after treatment, and global gene expression profiles were compared [87]. Genes that were either induced or suppressed more than two-fold by EGCG treatment compared with vehicle treatment in the same genotype group were filtered using the GeneSpring software. Among these well-defined genes, 671 EGCG-regulated Nrf2-dependent genes and 256 EGCG-regulated Nrf2-independent genes were identified in liver, whereas 228 EGCG-regulated Nrf2-dependent genes and 98 EGCG-regulated Nrf2-independent genes were identified in the small intestine [87].

Allyl sulfides

Garlic oil contains several organosulfur compounds, such as diallyl sulfide (DAS), diallyl disulfide (DADS), and diallyl trisulfide (DATS), capable of inducing carcinogen-detoxifying enzymes. DAS induced NQO1 by 5-fold in wild-type mice, whereas induction was completely absent in *nrf2* (−/−) mice, indicating that DAS also activates Nrf2 [88]. DAS induced a dose- and time-dependent increase of HO-1 protein and mRNA levels without provoking apparent toxicity in HepG2 cells [89]. Furthermore, protein expression, nuclear translocation, and DNA-binding activity of Nrf2 were induced by DAS treatment. Both ERK and p38 MAP kinase pathways appeared to play an important role in DAS-induced Nrf2 nuclear translocation and HO-1 gene expression [89]. DAS caused a transient increase of ROS. The thiol antioxidant *N*-acetyl-L-cysteine (NAC) blocked not only DAS-induced ROS production but also ERK activation as well as nuclear translocation of Nrf2, and also HO-1 expression. Likewise, the prototypic antioxidant enzyme catalase attenuated DAS-induced ERK activation, Nrf2 nuclear translocation and HO-1 protein induction, whereas Trolox was ineffective. DAS-treatment rendered the HepG2 cells resistant to oxidative stress caused by hydrogen peroxide or arachidonic acid, and this was attributable to its induction of HO-1 as pharmacologic inhibition of HO-1 activity blunted the cytoprotective effects of DAS [89]. It is noteworthy that the prooxidant activity of DAS contributes to Nrf2-driven antioxidant enzyme induction, which conferred the protection against oxidative cell death induced by external stimuli.

Chen et al. examined Nrf2-driven ARE activity and antioxidant gene expression by garlic organosulfur compounds in HepG2 cells [90]. Among the three allyl sulfides derived from garlic, DATS was most potent in terms of inducing ARE activation and expression of detoxifying enzymes, such as HO-1 and NQO1. Cotreatment with thiol antioxidants NAC and GSH abrogated the ARE activity and the Nrf2 accumulation induced by DATS. Three major MAP kinases, i.e., ERK, JNK, and p38, were activated by DATS treatment. However, the inhibition of these MAP kinases

did not affect DATS-induced ARE activity. Likewise, the PKC pathway was not directly involved in DATS-induced ARE activity, but instead the calcium-dependent signaling pathway appeared to play a role in the DATS-induced [90].

Resveratrol

Resveratrol (*trans*-3,5,4'-trihydroxystilbene) is another extensively investigated dietary chemopreventive phytochemical that is present in grapes and other plant species. Resveratrol exerts anti-oxidant, anti-inflammatory and chemopreventive activities by modulating various events in cellular signaling [91]. Resveratrol prevented chemically induced tumorigenesis in many experimental models [92], [93], [94], [95]. As a mechanism of carcinogen detoxification and cellular antioxidant defense, resveratrol induced NQO1 activity in Hepa1c1c7 cells [96]. In another study, human K562 cells treated with resveratrol exhibited increased NQO1 expression that peaked at 24–48 h [97]. A 2.5-fold rise in NQO1 protein levels was associated with a comparable elevation in the mRNA copy number and a 3- to 5-fold increase in NQO1 enzymatic activity. The stimulation of NQO1 gene expression by resveratrol involved the ARE signaling, accompanied by an increase in the state of phosphorylation of Nrf2 and its re-distribution to the nucleus [97]. The compound was found to induce HO-1 expression and activity in human aortic smooth muscle [98] and rat pheochromocytoma (PC12) cells [99] via activation of NF- κ B and Nrf2, respectively.

Treatment of human primary small airway epithelial and human alveolar epithelial (A549) cells with cigarette smoke extract (CSE) dose-dependently decreased glutathione (GSH) levels and GCL activity, effects that were associated with enhanced production of ROS [100]. Resveratrol restored CSE-depleted GSH levels by upregulation of GCL via activation of Nrf2 and also quenched CSE-induced release of ROS. Interestingly, CSE hampered nuclear translocation of Nrf2 in A549 and small airway epithelial cells by sequestering Nrf2 in the cytosol through post-translational modifications of this transcription factor, such as aldehyde/carbonyl adduct formation and nitration. On the other hand, resveratrol attenuated CSE-mediated Nrf2 modifications, thereby inducing nuclear translocation of Nrf2 and GCL gene transcription [100].

Pungent vanilloids

The chemopreventive potential of capsaicin (*trans*-8-methyl-*N*-vanillyl-6-nonenamide), a major pungent principle of hot chilli pepper (*Capsicum annum* L., Solanaceae), has been reviewed earlier [101], [102]. Capsaicin induced expression of HO-1 in HepG2 cells by activating PI3K/Akt-mediated activation of Nrf2 signaling in a ROS-dependent manner [52]. This study hypothesized that a quinone metabolite or other reactive forms of capsaicin would covalently modify NQO1, thereby suppressing the expression and activity of this antioxidant enzyme. The resulting overproduction of ROS is speculated to stimulate PI3K/Akt-mediated activation of Nrf2 [52]. (10)-Shogaol, a pungent ingredient of ginger (*Zingiber officinale* Roscoe, Zingiberaceae) has been reported to interact with the cysteine 151 residue of human Keap1 to form an alkylated adduct as detected by the FT-ICR mass spectrometer [103]. The alkylation of Keap1 by this electrophilic natural product may contribute to its antioxidant, anti-inflammatory and chemopreventive properties.

Lycopene

Lycopene, a natural antioxidant present predominantly in tomato products, has been reported to exert chemopreventive activi-

ty, especially against prostate and mammary carcinogenesis. The antioxidant properties of lycopene are thought to be primarily responsible for its chemopreventive effects [104]. Ben-Dor et al. have demonstrated that treatment of MCF-7 cells, transiently transfected with GCS-hARE-*tk-luc* reporter plasmid, with lycopene increased the GCS-ARE activity [105]. GCS-ARE and NQO1-ARE reporter activities were induced in HepG2 cells as well [105]. Lycopene elevated the mRNA and/or protein levels of GCS and NQO1, enhanced the cellular GSH level and reduced ROS generation in MCF-7 and HepG2 cells. Incubation of HepG2 cells with lycopene increased the nuclear translocation of Nrf2. In addition, the induction of NQO1 and GCS by lycopene was diminished in HepG2 cells ectopically expressing a dominant negative mutant Nrf2, suggesting that lycopene induced NQO1 and GCS via Nrf2 activation [105].

Miscellaneous

Epidemiological studies have revealed an inverse relationship between coffee consumption and the risk of certain types of cancer [106], [107], [108]. Dietary administration of coffee (3 or 6%) for 5 days showed a several-fold increased expression of mRNA transcripts of NQO1 and GSTA-1 in the liver and small intestine, and that of UGTA-6 and GCLC in the small intestine of *nrf2*^{+/+} mice as compared to *nrf2*^{-/-} animals [109]. The coffee-derived diterpenes, cafestol and kahweol, when treated to embryonic fibroblasts isolated from *nrf2*^{+/+} mice, increased NQO1 mRNA expression to a greater extent than that achieved with embryonic fibroblasts from *nrf2*^{-/-} mice. This study also demonstrated that the induction of NQO1 mRNA expression was dependent on the presence of an ARE element at the gene promoter segment and that the coffee-specific diterpenes induced NQO1 reporter gene activity via the activation of Nrf2 [109].

Carnosol, an *ortho*-phenolic diterpene present in rosemary (*Rosmarinus officinalis*, Lamiaceae), induced HO-1 expression at both protein and mRNA levels by increasing the binding of Nrf2 to ARE and induced the Nrf2-dependent activation of *HO-1* promoter in PC12 cells via upregulation of ERK, p38 MAP kinase and JNK pathways [110]. Cinnamaldehyde present in the dried stem bark of *Cinnamomum cassia* Presl. (Lauraceae), induced HO-1 protein expression, increased Nrf2 nuclear translocation and the ARE-luciferase reporter activity in human endothelial cells [111].

Zerumbone, a chemopreventive sesquiterpene derived from tropical ginger (*Zingiber zerumbet* Smith) [112] enhanced the cellular GSH level and induced a battery of antioxidant enzymes, such as GSTP-1, GCL, GPx and HO-1 in normal rat liver epithelial (RL34) cells. Treatment of RL34 cells with zerumbone (25 μ M) showed increased nuclear accumulation of Nrf2, while its reduced analogues such as α -humulene or 8-hydroxy- α -humulene failed to activate Nrf2 and induce aforementioned antioxidant enzymes, suggesting that the α,β -carbonyl moiety at the 8 position is crucial for Nrf2 activation and antioxidant enzyme induction by zerumbone [113].

Chalcone, an α,β -unsaturated flavonoid, upregulated the nuclear levels of Nrf2 and increased the ARE-luciferase activity and also the thioredoxin reductase promoter activity in bovine aortic endothelial cells [114]. It induced expression of thioredoxin reductase as well as HO-1 in the same cells. Furthermore, chalcone suppressed the activation of NF- κ B and STAT3 in both IL-6 and LPS-stimulated endothelial cells. The presence of an α,β -unsaturated carbonyl moiety is critical for the anti-inflammatory activity exerted by chalcone [114]. Some synthetic chalcone derivatives,

such as 2',4',6'-tris(methoxymethoxy)chalcone and 3',4',5',3,4,5-hexamethoxychalcone diminished NF- κ B activation, whereas they induced HO-1 expression [115], [116]. The structure-activity relationship assay of a series of unsubstituted and hydroxy-substituted chalcones showed that ring hydroxylation at position C2 or C2' or C4' enhanced the phase 2 enzyme inducing ability by 2- to 3-fold as compared to non-substituted chalcones or 4-hydroxy-substituted chalcone [117]. The simultaneous presence of a hydroxy moiety at C2 and C2' further increased the potency [117]. Xanthohumol, a sesquiterpene derived from hops (*Humulus lupulus* L.), exhibited chemopreventive activity. One of the mechanisms of chemoprevention with xanthohumol was the induction of antioxidant enzymes. Pretreatment of hepa1c1c7 cells with xanthohumol diminished menadione-induced DNA damage via up-regulation of NQO1 [118]. Xanthohumol induced NQO1 in an ARE-dependent manner, partly via the activation of Nrf2 through alkylation of Keap1 at the cysteine 27 residue [118]. In addition, liquid chromatographic-tandem mass spectrometric (LC-MS/MS) analysis revealed that xanthohumol alkylated the cysteine 151 residue, located at the BTB domain of Keap1, thereby contributing to the Nrf2-dependent ARE activation [103]. Likewise, isoliquiritigenin derived from licorice has also been shown to alkylate the cysteine 151 residue of Keap-1, thereby inducing the ARE activity [103].

Conclusion

▼ Considering the burden of health-care cost as a major global concern, the dietary chemoprevention provides an inexpensive, readily applicable, and easily accessible approach to cancer control and management. In general, dietary phytochemicals exert their chemopreventive effects through multiple mechanisms. The saturation of cellular defense mechanisms due to overwhelming external stress or their disruption as a consequence of deregulated intracellular signaling pathways responsible for inducing expression of a battery of carcinogen detoxifying/antioxidant enzymes would make cells/tissues more vulnerable to oxidative, nitrosative and inflammatory insults implicated in multi-stage carcinogenesis. Therefore, fortification of cellular defense mechanism or restoration of stress-response signaling by intaking dietary phytonutrients provides an important strategy for chemoprevention.

The bZIP transcription factor Nrf2 controls the cellular signaling involved in the transcriptional activation of genes encoding a panel of phase 2 detoxifying/antioxidant enzymes and stress-responsive proteins. Nrf2 plays an essential role in maintaining cellular homeostasis and hence represents a critical target for chemoprevention of oxidative stress- or inflammation-associated carcinogenesis. Under normal physiological conditions, Nrf2 forms an inactive complex with the negative regulator, Keap1 which controls the subcellular localization and steady-state levels of Nrf2. Cysteine residues present in Keap1 function as redox sensors, and oxidation or chemical modification of some of the highly reactive cysteine residues facilitates the dissociation of Nrf2 from Keap1 and subsequent nuclear translocation. Therefore Nrf2 constitutes a unique "redox switch" that can be turned on in response to redox imbalance caused by oxidative and electrophilic stresses. However, such adaptive response to external stress is normally transient and prone to be readily saturated. Therefore, it may be necessary to maintain proper levels of cellular defense capacity by inducing *de novo* synthesis of cy-

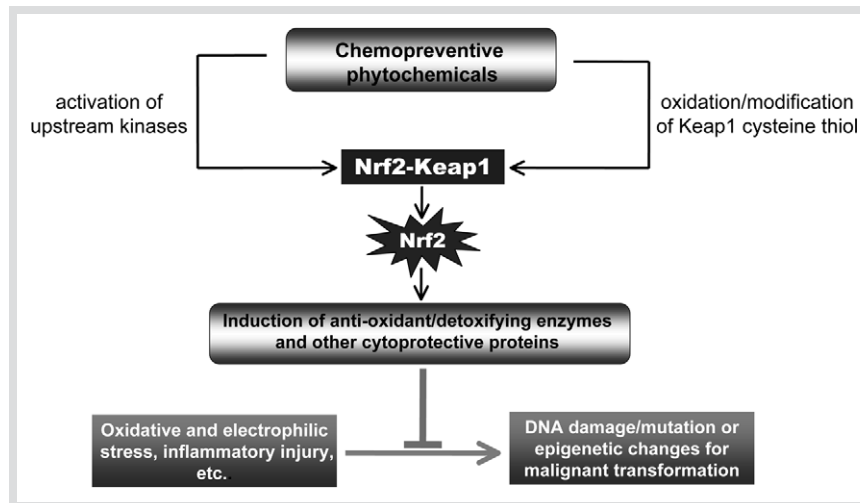


Fig. 3 Activation of Nrf2 signaling by chemopreventive phytochemicals. Chemopreventive phytochemicals can activate Nrf2 signaling by inducing phosphorylation of Nrf2 via activation of upstream protein kinases and/or through direct interaction with Keap1 cysteine thiols. The subsequent induction of cytoprotective enzymes can abrogate oxidative stress and inflammatory tissue injury, thereby blocking DNA damage or suppressing proliferation of initiated cells and malignant transformation.

toprotective proteins in a more sustained manner. In this context, it is of particular interest to note that many inducers of phase 2 and antioxidant proteins are present in our daily diets, especially those derived from edible plants, and intake of these dietary phytochemicals ensures long-term cytoprotection. While oxidation and/or direct modification of critical Keap1 cysteine thiols have been proposed as a major plausible mechanism underlying activation of Nrf2, phosphorylation of Nrf2 at specific serine and/or tyrosine residues also represents another important event in cytoprotective gene induction, which is modulated by many dietary chemopreventives. The activation of Nrf2-ARE signaling by dietary inducers will fortify cellular defense against oxidative, electrophilic and inflammatory stresses which are broadly implicated in multi-stage carcinogenesis as schematically represented in **Fig. 3**.

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