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# Chemical and biological mechanisms of phytochemical activation of Nrf2 and importance in disease prevention

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#### Abstract

Plants are an incredibly rich source of compounds that activate the Nrf2 transcription factor, leading to upregulation of a battery of cytoprotective genes. This perspective surveys established and proposed molecular mechanisms of Nrf2 activation by phytochemicals with a special emphasis on a common chemical property of Nrf2 activators: the ability as "soft" electrophiles to modify cellular thiols, either directly or as oxidized biotransformants. In addition, the role of reactive oxygen/nitrogen species as secondary messengers in Nrf2 activation is discussed. While the uniquely reactive C151 of Keap1, an Nrf2 repressor protein, is highlighted as a key target of cytoprotective phytochemicals, also reviewed are other stress-responsive proteins, including kinases, which play non-redundant roles in the activation of Nrf2 by plant-derived agents. Finally, the perspective presents two key factors accounting for the enhanced therapeutic windows of effective phytochemical activators of the Keap1–Nrf2 axis: enhanced selectivity toward sensor cysteines and reversibility of addition to thiolate molecules.

### 7.1 Introduction

Numerous phytochemicals have shown great promise for prevention and treatment of various human diseases. For example, ClinicalTrials.gov lists 25 different intervention trials investigating the effects of standardized preparations of broccoli sprouts. These trials include the amelioration of symptoms of diseases as diverse as autism<sup>1</sup>, cystic fibrosis, influenza, asthma, and chronic obstructive pulmonary disease. The trials also encompass studies on the prevention of breast, lung, and prostate cancers, carcinogenesis from aflatoxin exposure, and cardiovascular disease. The purpose of this Perspective is to *i*) review a key molecular mechanism of phytochemicals in the prevention and amelioration of diseases: the activation of the transcription factor NF-E2-related factor 2 (Nrf2), *ii*) outline common

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<sup>&</sup>lt;sup>1</sup>ClinicalTrials.gov identifiers, in order as listed above. Amelioration studies: <u>NCT01474993</u>, <u>NCT01315665</u>, <u>NCT01269723</u>, <u>NCT01183923</u>, <u>NCT00994604</u>. Prevention studies: <u>NCT00982319</u>, <u>NCT00255775</u>, <u>NCT00607932</u>, <u>NCT01437501</u>, <u>NCT00252018</u>.

chemical features of potent Nrf2 activators, and *iii*) provide perspectives for harnessing these features for more effective disease prevention or treatment.

#### 7.2 Upregulation of Cytoprotective Genes by Nrf2

Oxidative stress and associated inflammation contribute to progression of many chronic degenerative diseases in humans [1, 2]. Exposure to oxidant and electrophilic agents from air, water, food, and other environmental sources has also been implicated in a large (70–90%) component of cancer and cardiovascular disease risks [3]. The Nrf2 transcription factor has emerged as a key player in protecting cells against various intrinsic and extrinsic assaults. Nrf2 regulates more than 600 genes, including over one hundred that encode cytoprotective proteins [4], named for their ability to protect cells against oxidative stress, reactive electrophilic species, and other types of stress (reviewed in [5]). In brief, these proteins include antioxidant enzymes, NADPH regeneration enzymes, glutathione biosynthesis enzymes, heat shock proteins, enzymes that facilitate the elimination of xenobiotic toxicants such as detoxification enzymes and drug-efflux pumps, as well as subunits of the 26S proteasome. Collectively, these Nrf2-regulated genes share at least one

copy of the antioxidant response element (ARE,  $5' - \frac{A}{G}TGA \frac{C}{G} \frac{T}{A} \frac{C}{N} \frac{A}{N}GC \frac{A}{T} - 3'$ ) in their promoter region [4]. Nrf2 binds to the ARE as a heterodimer with one of several small Maf transcription factors, leading to upregulation of gene transcription.

Upregulation of this battery of cytoprotective proteins through Nrf2 activation is a critical component of an organism's ability to cope with intrinsic and extrinsic stress factors, including inflammation, reactive oxygen/nitrogen species (ROS/RNS), shear stressors such as endothelial stressors, and environmental toxins. Studies on  $Nrf2^{-/-}$  mice, which have low and largely non-inducible levels of many cytoprotective proteins [4], serve as striking and comprehensive examples of the importance of these proteins in maintaining health and preventing disease. Nrf2-deficient mice are prone to develop disorders that are caused by ROS and inflammation, including macular degeneration [6], neurodegeneration in a murine model of Parkinson's disease [7], cardiac disorders [8, 9], and chemically-induced tumorigenesis [10-12]. Furthermore,  $Nrf2^{-/-}$  mice are more susceptible to damage of the blood-brain barrier following brain injury [13], formation of carcinogen-DNA adducts in the lung exposed to diesel exhaust [14], acute pulmonary injury induced by butylated hydroxytoluene [15], and hepatic damage by acetaminophen [16]. In addition, they are deficient in their intrinsic capacity for skin wound healing [17]. Since loss of Nrf2 increases pathological cell and tissue damage in response to intrinsic and extrinsic factors, it is believed that upregulation of Nrf2 serves both cytoprotective and preventive roles in diverse pathophysiological situations.

In support of the importance of Nrf2 in preventing human diseases, inherited DNA polymorphisms that reduce the abundance of Nrf2 are associated with various pathologies, including chronic gastrisis, ulcerative colitis, skin pathologies such as skin vitiligo, and adult respiratory distress syndrome (reviewed in [18]). In fact, a number of clinical trials that assess the effects of broccoli sprouts include direct measurements of Nrf2 activation. For example, Nrf2 levels, along with markers of oxidative stress, will be assessed after administration of macerated broccoli sprouts in both healthy volunteers and those with

cystic fibrosis<sup>2</sup>, in nasal epithelial cells obtained by curettage, as well as in alveolar macrophages and bronchial epithelial cells of patients with chronic obstructive pulmonary disease<sup>3</sup>.

#### 7.3 Phytochemical Activation of Nrf2

#### 7.3.1 Overview

Plants have been an incredibly rich source for the identification of compounds that activate cytoprotective genes. Development of a simple microtiter-plate based assay [19] to assess induction of the cytoprotective enzyme NAD(P)H: quinone oxidoreductase 1 (QR1) in mouse Hepa1c1c7 cells has greatly facilitated the ability to screen for and identify many cytoprotective phytochemicals. For example, a collective effort of colleagues at the University of Illinois at Chicago and Purdue University has identified 66 compounds from 18 plant species that are active in the QR1 assay [20]. Representative phytochemicals that have been shown to activate cytoprotective genes are listed in Table 1.

Several plant families important for human diets are particularly rich in ARE inducers. For example, many organosulfur activators have been isolated from garlic and onion, edible members of Allium family. The Cruciferous family of vegetables, including broccoli, cabbage, Brussels sprouts, horseradish, mustard and watercress, produces a particularly large and functionally diverse number of potent ARE inducers (Table 1). These plants contain glucosinolates, the thioglucoside conjugates of the ARE-activating species. Altogether, over 120 glucosinolates have been identified from various plants [21]. These are enzymatically converted to the ARE-inducing forms either by myrosinase, a thioglucosidase that is localized in a separated cellular compartment and is released upon maceration or chewing, or by intestinal microflora after ingestion [22]. Three general classes of ARE inducers produced by myrosinase-catalyzed hydrolysis of glucosinolates are isothiocyanates, indoles and epithionitriles (Table 1). The enzymatic hydrolysis mechanisms involved in their release are reviewed elsewhere [23, 24]. Finally, phenolic compounds, another important class of Nrf2-activating agents, have been isolated from diverse plant families, including grapes (Vitaceae) and teas (Theaseae), which are rich sources of flavonoids and related catechins.

Importantly,  $Nrf2^{-/-}$  mice experiments highlight the key role of Nrf2 in mediating the cytoprotective effects of the phytochemicals discussed above. Thus, sulforaphane, one of the active components of broccoli sprout extracts has been reported to inhibit skin [12] and forestomach [10] carcinogenesis in wild-type (wt) mice, but its ability to do so is significantly attenuated in  $Nrf2^{-/-}$  animals. In addition, sulforaphane was able to protect the blood-brain barrier post-injury only in wt Nrf2 mice [13]. Similarly, resveratrol, a flavonoid-like molecule produced by many plants, protected against high-fat-diet-induced oxidative stress in aortas of wt but not  $Nrf2^{-/-}$  mice [9].

<sup>&</sup>lt;sup>2</sup>Clinicaltrials.gov identifier NCT01315665

<sup>&</sup>lt;sup>3</sup>Clinicaltrials.gov identifier <u>NCT01335971</u>

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#### 7.3.2 The Chemistry Required for Phytochemicals to Activate Nrf2

**7.3.2.1 Reactivity with Thiolates**—Despite the high level of overall structural diversity among ARE inducers, many cytoprotective phytochemicals from different plant sources share common thiol-reactive chemical motifs (collectively shown in Figure 7.1 as red moieties). For example, the presence of an  $\alpha$ , $\beta$ -unsaturated carbonyl group, a potential Michael acceptor, is a particularly common feature in ARE inducers, including withanolides (e.g. withaphysacarpin), chalcones (e.g., isoliquritigenin), butenolides (e.g.  $\beta$ -angelica lactone), oxidized terpenoids (e.g. zerumbone, E-gugglesterone, and citrals), and curcuminoids (e.g. curcumin). Other inducer classes are similarly electrophilic epithionitriles (e.g. 1-cyano-2,3-epithiopropane), isothiocyanates (e.g. sulforaphane), and organopolysulfides (e.g. allicin). Talalay and colleagues in 1988 first recognized that the diverse structures share the ability to react with thiolate groups [25]. They hypothesized that gene induction takes place by virtue of an intracellular sensor that contains one or more reactive cysteine residues, modification of which by inducing agents would lead to target gene activation. This seminal hypothesis is supported by numerous studies that followed, which directly linked the presence of particular functional groups to Nrf2-ARE induction, as shown with zerumbone [26], chalcones [27], flavonoids [28], and withanolides [20, 29]. Furthermore, a strong correlation has been identified between inducer potencies and chemical reactivities toward thiolates [30].

Finally, in 1999 a key repressor of Nrf2 was discovered, the Keap1 protein, which was found to possess an unusually large number of cysteines (25 and 27 in the mouse and human proteins, respectively) [31]. As described in detail in Section 7.3.2.2, a subset of these cysteines, C151 in particular, has been found to be important for Nrf2 activation by phytochemicals.

#### 7.3.2.2 Phytochemicals that Do Not Have the Ability to React with Thiolates

While the critical feature of an Nrf2 activator appears to be the ability to react with thiolates, a large number of phytochemicals, including phenols, monosulfides, furans, and indoles, would need to acquire this property through metabolic and/or chemical processing. Studies on biotransformations of many such molecules are surprisingly limited, despite the current interest in health benefits of phytochemicals. Herein, we summarize the available experimental evidence from literature implicating mechanisms of converting phytochemicals to thiol-reactive species. We also discuss two classes for which transformation to a thiolate-reactive species is considerably more difficult to ascertain, carotenoids and 1,3-polyphenols.

One relatively well-characterized chemical class of phytochemicals that require electrophilic conversion for inducer activity are electron-rich phenolic compounds featuring "additive" distribution of electron-donating groups (Figure 7.1, blue structural fragments). These are compounds that contain phenolic hydroxyl groups in a conjugated system with an even number of carbons separating them. Representative examples of this class include 1,2-diphenols (e.g., epigallocatechin gallate (EGCG), caffeic acid, and piceatannol) and vinylogous 1,6-diphenols, with an alkene spacer acting as the "electron conduit" (e.g., quercetin and kaempferol). The 1,2- 1,4-, and 1,6-diphenols, but not the mono- or 1,3-

dihydroxy variants, can be readily oxidized in vivo by a variety of mechanisms (Figure 7.2) to the corresponding quinoids and will be referred to herein as quinoid-forming phenols. In fact, such phenols have been associated with induction of carcinogen-detoxifying enzymes by one of the earliest observations in the field [32]. The resulting Michael acceptors can then readily react with cysteine thiolates (Figure 7.2). For example, the well-characterized flavonoid quercetin is expected to be a very weak electrophile intrinsically due to the electron-donating effect of the 3-hydroxyl group and both aromatic rings (see Figure 7.3.A for numbering scheme). However, in human blood plasma, quercetin is readily oxidized to a significantly more electrophilic quinone methide [33] (Figure 7.3.A). Importantly, the oxidized species is much more reactive toward thiolates, and its conjugation products have been detected with glutathione (GSH), N-acetylcysteine (NAC), cysteine [34, 35], and protein cysteine residues [36]. Similarly, EGCG contains several aromatic 1,2-dihydroxy units, and thus can form quinones that react with isolated and protein thiolates, as demonstrated in both biochemical experiments and cells [37, 38]. EGCG has also been found in mouse urine as the S-cysteinyl-EGCG conjugate after a high oral dose [39]. Furthermore, the oxidative conversion of phenolic compounds to Michael acceptors has been shown to correlate strongly with cytoprotective enzyme induction in studies evaluating tert-butylhydroquinone (tBHQ) [40] and EGCG analogs [38], as well as a broad series of phenols [41]. As further evidence that the oxidation of phenols is a prerequisite to ARE induction,  $Cu^{2+}$  or other oxidized transition metal cations in the media strongly stimulated the ARE induction potential of para- and ortho-hydroquinones [42]. These metal ions act as catalysts in the oxidation of phenols to Michael reaction acceptors (Figure 7.2) under aerobic conditions. Importantly, transition metal salts had no effect on inducer activity of the corresponding quinones or sulforaphane [43].

The organosulfur compounds from garlic and onions were shown by the Wattenberg group in 1988 to have interesting structural requirements for inducer activity [44]. Garlic organopolysulfides and derivative thiosulfinates, such as allicin (Figure 7.1), were able to induce the cytoprotective enzyme glutathione S-transferase, as might be expected from their ability to modify cellular thiols. However, the monosulfide diallylsulfide (DAS) is also an inducer, and the mechanism by which it activates Nrf2 is yet to be delineated. In addition, the diallyl forms of the di- and tri-sulfides were much more potent than the propyl version. In surmising why the diallyl sufides might have higher potency, it is interesting to note that cytochrome P450 2E1 (CYP2E1) is implicated in sequential conversion of the sulfide to the corresponding sulfoxide (DASO) and sulfone (DASO<sub>2</sub>) derivatives [45]. Further oxidation of the sulfone metabolite generated an electrophile that was shown to act as a suicide inhibitor of CYP2E1, as well as other unidentified cytochromes implicated in bioactivation of various cytotoxins [45]. While the exact mechanism of this inhibition remains to be established, the irreversible nature of the antagonism implies generation of a reactive intermediate capable of covalent modification of the enzyme involving, in all likelihood, an active site cysteine [46]. In addition, LC-MS/MS analysis of bile fluids from rats treated with DAS, DASO and DASO<sub>2</sub> identified several GSH conjugates, implicating epoxidation of the allylic group as an important metabolic activation step for all three compounds [47]. Unlike the epoxides of DAS and DASO, the DASO<sub>2</sub>-derived epoxide provides a unique entry into a sulfone-activated Michael acceptor containing a sulfonylprop-2-en-1-ol group

(see Figure 7.3.B for the proposed mechanism). This so-called "soft" electrophile, with highly distributed charge density is much more likely to react with a thiolate, a similarly polarizable "soft" nucleophile, rather than the "hard" epoxide (see Section 7.3.3 for a brief discussion of hard and soft reagents, and other effects affecting electrophile chemoselectivity). Therefore, oxidation of the sulfide, in combination with epoxidation of the allylic group, could be responsible for the inactivation of CYP2E1 and, possibly, induction of Nrf2 by modifying sensor cysteines.

Furan-containing compounds, such as the diterpenes cafestol and kahweol, are known to act as the principal Nrf2 activators in coffee despite lacking a thiol-reactive functional group [48, 49]. Although the exact mechanism of induction is yet to be established, recent mass spectrometric study of bile fluids from mice injected with cafestol was interpreted by the authors to suggest that epoxidation is the key step in converting coffee furans into thiolreactive species [50]. However, the study did not distinguish between direct addition to the epoxide and addition at a remote double bond conjugated to the epoxide or Michael addition to a  $\gamma$ -ketoenal, arising from ring-opening reaction of the oxidized furan (depicted in Figure 7.3.C). The distinction is important because furan epoxides and corresponding dicarbonyl derivatives have long been associated with severe cytotoxicity of furan-containing compounds [51], due to both high reactivity and tendency to react with oxygen and nitrogen nucleophiles, in addition to thiolates. This, however, is inconsistent with the generally cytoprotective properties of the coffee furans. The particular structural environment of electrophiles derived from these furans may account for the observed shift in the balance of toxic and protective effects (Figure 7.3.C). Thus, kahweol epoxide presents a doubly conjugated epoxide (epoxydiene) requiring a soft thiolate nucleophile to attack the soft electrophilic center next to a highly sterically congested quaternary carbon. The corresponding ketoenal is also deactivated toward non-specific additions by both crossconjugation of the dienone functionality and  $\beta_{\beta}$ -dialkyl substitution of the unsaturated aldehyde [52]. Alternatively, participation of less reactive metabolites, such as conjugated lactones, for example, which have been detected as metabolic derivatives of furancontaining compounds [53], may account for the low toxicity of these phytochemicals. Further studies will be required for understanding the unique properties of these furans.

Indole-3-carbinol (I3C), a glucosinolate breakdown product, and its digestive product 3,3'diindolylmethane (DIM) are ARE inducers [54], and have many other cancer chemopreventive effects [55]. However, while DIM has been described as an effective inducer of the ARE with little associated long-term toxicity [56], I3C has a weak level of Nrf2 activation and is associated with a complex interplay of pro- and anticarcinogen effects (for a review see [56]). Although no specific mechanism accounting for the ARE induction ability of DIM has been established, reaction schemes for producing electrophilic species capable of covalent modification of thiols could be inferred from known metabolic and chemical transformations of I3C, DIM, and 3-methylindole (3MI), a byproduct of tryptophan metabolism by intestinal microflora [57]. In the case of the latter compound, a highly electrophilic thiolate-reactive derivative 3-methyleneindolenine, an  $\alpha,\beta$ -unsaturated imine, is produced via cytochrome P450-mediated dehydrogenation [58]. This species is postulated to be responsible for the cytotoxicity of this substituted indole [58, 59]. The reactivity of the  $\alpha,\beta$ -unsaturated imine toward nucleophiles is promoted by a concomitant

rearomatization of the indole nucleus, which makes it a rather non-discriminating and thus toxic agent, capable of modifying a variety of functionalities found in a cellular environment. 3-Methyleneindolenine also serves as an intermediate in the dehydrationconjugate addition-retro-aldol cascade in the acid-catalyzed conversion of I3C to DIM [60] (Figure 7.3.D). Therefore the low level of Nrf2 activation and pro- and anticarcinogen effects associated with I3C could be explained by the formation of both the toxic electrophile and DIM, respectively, in the course of the same chemical process. On the basis of established biotransformations of DIM [61], it is tempting to propose that an oxidation event similar to that seen for 3MI can lead to the extended conjugation-stabilized Michael acceptor, 3-(3-indolylmethylene)-indolenine (Figure 7.3.D). We must note that no thiol conjugates of DIM metabolites have been isolated, perhaps, due to reversibility of such additions to the conjugation-stabilized electrophile (see Section 7.3.2.4 for further discussion of reversibility importance). Significantly, a sulfate-conjugated hydration product of the proposed species has been isolated as one of the major biotransformants from cultured cancer cells [61]. The Michael acceptor produced by both I3C and 3MI may be the principal agent responsible for associated toxicities. Unlike 3MI, however, biotransformation of I3C can also lead to DIM, a more stable indole derivative [62] that is unlikely to produce a nondiscriminating electrophile. This example underscores that the level of reactivity of phytochemical-derived electrophiles could be the key determinant in a sensitive balance of cytotoxic and cytoprotective effects.

Carotenoids, lycopene in particular, have been shown to activate the ARE via Nrf2 [63]. The carotenes are unsaturated hydrocarbons and thus contain no thiol-reactive species. However, the authors point out that the ethanolic extract of the lycopene preparation, containing unidentified hydrophilic lycopene derivatives, activated the ARE with a similar potency as lycopene. Therefore, it seems likely that oxidation of the polyene, leading to formation of an electrophilic species (e.g. citral [64]), is a prerequisite for Nrf2 activation.

In addition, there are a handful of phenols (non-quinoid forming) that have no electrophilic moieties, or facile non-enzymatic autoxidative paths to obtaining these moieties, which have been shown to activate cytoprotective enzymes, such as resveratrol [65] and galangin [28]. The Nrf2-dependence of ARE activation by resveratrol has been particularly wellestablished in studies in  $Nrf2^{-/-}$  mice [9] and normal human small airway epithelial cells (SAEC) [66]. Resveratrol and other related phenols can be hydroxylated to quinoid-forming species by a member of the cytochrome P450 (CYP) superfamily of monooxygenases, such as pro-carcinogen activating CYP1A1, CYP1A2, and CYP1B1 [67, 68]. These enzymes are transcriptionally controlled by the aryl hydrocarbon receptor (AhR). So-called bifunctional ARE inducers, such as β-naphthoflavone (β-NF), activate Nrf2 by first binding to and activating AhR, which in turn leads to upregulation of CYP enzymes [69]. However, resveratrol is a known inhibitor of AhR [70, 71], and therefore in uninduced normal cells resveratrol may not be hydroxylated by CYPs prior to Nrf2 activation. Resveratrol activation of QR1 through the estrogen receptor  $\beta$  (ER $\beta$ ) in breast cancer cells has been explored as another mechanistic possibility. Various studies support a model in which binding of phytoestrogens, including resveratrol, to ER $\beta$  causes ER $\beta$  to bind and activate the QR1 ARE [72-75]. However, this mechanism appears to be restricted to cancer cells overexpressing

ERβ [75]. A third mechanism has been suggested based on the observation that resveratrol binds to and inhibits quinone reductase 2 (QR2) with low nanomolar affinity [76]. QR2's function is not clearly elucidated, but it is known to catalyze the reduction of quinones, among several other classes of electron-deficient compounds. The authors suggested that resveratrol may activate Nrf2 by inhibiting QR2, resulting in the accumulation of endogenous quinones that can then induce electrophilic stress by modifying cellular thiols [76]. A similar indirect induction mechanism could apply to other phytochemical inhibitors of QR2 that inhibit this reductase at physiologically relevant concentrations, including quercetin, kampferol, apigenin, or genistein [76, 77]. This hypothesis warrants further investigation. Finally, a fourth mechanism accounting for activation of Nrf2 by non-quinoid-forming phenols by means of kinase activation is explored for genistein in Section 7.3.4.3.

**7.3.2.3 Role of ROS in Nrf2 Activation by Phenolic Compounds**—One important area of consideration is to what extent ROS are involved in the activation of Nrf2 by phytochemical inducers. Phenolic ARE inducers can play multiple roles in redox status of a cell (Figure 7.2). They are able to act as both direct antioxidants, scavenging free radicals, and, if converted to thiolate-reactive electrophiles, as indirect antioxidants by upregulating antioxidant genes [78]. Depending on the experimental conditions, they can also act as prooxidants [79]. As shown in Figure 7.2, the non-enzymatic autoxidation of a polyphenol can lead to ROS formation. This generation of ROS may play a role in Nrf2 activation, as well as have potential off-target effects such as toxicity. The ability of ROS to upregulate Nrf2 is well-established, for example by treatment of cells with  $H_2O_2$  [80]. One mechanism by which ROS may activate Nrf2 is depicted in Figure 7.2. In this scenario, generation of superoxide and other ROS can lead to oxidation of GSH to GSSG [81]. GSSG then could modify sensor thiolates such as Keap1 cysteines [82, 83], thereby activating Nrf2 (see also section 7.3.4.1). In addition, sensor thiolates can be directly oxidized by ROS (reviewed in [84] and [85]).

There are indications that ROS mediate signaling for some ARE inducers. For example, the generation of ROS in cells [86, 87] and cell media [88] by EGCG has been well established. Importantly, ROS-scavengers NAC, GSH, superoxide dismutase (SOD), and catalase all inhibited the induction of heme oxygenase 1 (HO-1) by EGCG, as shown in bovine aortic endothelial cells (BAECs) [89]. Thus, EGCG-induced ROS, rather than an EGCG-derived electrophile, mediated EGCG-induced HO-1 expression under these conditions. A role for ROS as a secondary messenger in Nrf2 activation, specifically H<sub>2</sub>O<sub>2</sub>, has also been shown for a different ARE inducer class, dithiolethiones, in Hepa 1c1c7 cells [80]. In addition, a role for ROS in Nrf2 activation by 21 flavonoids (including fisetin, kaempferol, and quercetin) was explored [90]. A high level of correlation was observed between the flavonoids' ability to activate the ARE and their computed energy levels of the highest occupied molecular orbitals ( $E_{HOMO}$ ), representing the tendency of the flavonoid to donate electrons in redox processes, for example. Thus, more oxidizable flavonoids possessing less negative  $E_{HOMO}$  values were generally more potent inducers of ARE-mediated gene expression. Therefore ROS may be important secondary messengers for flavonoids.

Interestingly, we note that there are three outliers in the flavonoid correlation analysis [90] with much greater abilities to activate the ARE than predicted by their  $E_{\text{HOMO}}$  values (and

hence their tendency to donate electrons in the formation of ROS). This boost in induction potency could be associated with a particular set of structural features. Thus, only these three flavonoids (quercetin, morin, myricetin) out of 21 tested can be predicted to form extended conjugation-stabilized quinone methides involving the C ring, as products of a two-electron oxidation sequence (Figure 7.3.A). We hypothesize that while generation of ROS by flavonoids may contribute to activation of Nrf2, the ability to form a conjugationstabilized electrophile, which can react with thiolates such as quinoid, produces a much greater extent of ARE activation due to its ability to react directly with sensor thiolates, such as Keap1 cysteines. Importantly, the quinone methide derived from quercetin has a remarkably high stabilization effect of the extended conjugation, as shown by its ability to form reversible thiol adducts [34]. Finally, in support of this hypothesis, out of a series of five structurally similar flavonoids, only the quinoid-forming ones (kaempferol, quercetin and luteolin) were able to react with a thiol (GSH) to form *mono-* and *bis-*GSH conjugates, without forming radicals and ROS [91, 92]. In contrast, the other two flavonoids (apigenin and naringenin), which are not able to form conjugation-stabilized electrophiles, generated radicals and ROS and could not form conjugates with GSH.

In considering whether generation of ROS is a secondary messenger in Nrf2 activation by flavonoids and other phytochemicals, it is important to note that the two-electron oxidation of phenols needs not necessarily lead to ROS formation. If a polyphenol can be recognized as a substrate of an oxidizing metalloenzyme, it can be converted to the Michael acceptor without production of ROS (Figure 7.2). To our knowledge, this theory remains to be tested. Dosage is likely critical as to whether useful or harmful levels of ROS are produced by treatment with ARE inducers [93]. Much work remains to determine the role(s) that ROS play in Nrf2 activation by phytochemicals, including whether ROS generated by therapeutic and physiologically relevant concentrations of phytochemicals have deleterious off-target effects, or are relatively harmless and perhaps relevant for participating in numerous signaling mechanisms, which are beyond the scope of this Perspective (reviewed in [94], [95], and [85]). The dosage amount is likely very important, as illustrated through the example of the synthetic oleanane triterpenoid CDDO-Im. While at low concentrations (100 nM) CDDO-Im is a potent Nrf2 inducer with undetectable adverse effects, above 300 nM ROS-mediated toxicity is observed [96]).

#### 7.3.3 Basis of Phytochemicals as Therapeutic, Rather than Toxic, Agents

The cytoprotective roles of phytochemicals discussed thus far could be considered surprising, given that most if not all were produced for biodefense against insects, bacterial parasites and other animals, including humans, consuming plant parts. The questions arises then as to how these molecules have a cumulatively cytoprotective effect. One critical factor is dosage, as at high concentrations most of these molecules have been shown to display at least some level of toxicity. At low concentrations then, the residual "toxicity" maintains cells in a state of adaptive stress, providing them tools for counteracting a variety of adverse conditions [97]. The ability of a phytochemical to induce such a state selectively plays a critical role in the cytotoxicity/cytoprotection balance. This balance appears to be rather sensitive to subtle changes in a chemical structure. Two considerations that appear to be significant are selectivity for stress-sensing cysteines and reversibility of thiol modification.

There are various factors that contribute to preference of an electrophile for stress-sensing cysteines. First, cytoprotective compounds (or their bioactivated derivatives) contain "soft" electrophilic centers, which generally favor an attack by a corresponding soft nucleophile, best represented in a cellular environment by a thiolate anion. In phytochemicals, soft electrophilicity is associated with a carbon center conjugated through a network of  $\pi$ -bonds to a reactive functional group, such as a carbonyl (Figure 7.3.A), sulfone (Figure 7.3.B), epoxide (Figure 7.3.C), or imine (Figure 7.3.D), or with an electrophilic center activated directly by large polarizable atoms (e.g. epithionitriles, isothiocyanates, etc.). The molecular orbital effects at the root of these preferences are important but are beyond the scope of this perspective (See [98] for relevant discussion). Soft electrophilicity, however, is not sufficient for shifting the therapeutic window toward cytoprotection, as highly reactive albeit soft agents will modify unintended thiols (i.e., hemoglobin, human serum albumin, glutathione, etc) [3] and even non-thiol targets (i.e., nucleic acids).

Second, stress-sensing cysteines are likely maintained by their immediate environment in a highly reactive state. This could include pKa reduction [99] and the presence of Brønsted acids to orient and activate the incoming electrophiles through hydrogen bonding [52] (See Figure 7.4.A). It is highly likely that one or several Keap1 cysteines are in fact presented in such an environment (see Section 7.3.4.1).

Third, the structural context of presenting the electrophilic functionalities is likely to play an essential role in an electrophile's ability to react selectively. Thus, another contributing factor noted in enhancing reactivity with sulfur nucleophiles is the presence of a neighboring group, such as a hydroxyl functionality in a vicinity of a reactive  $\beta$ -carbon of cinnamates, chalcones, curcuminoids, bis(2-hydroxybenzylidene)-derivatives [30], and some withanolides [20]. A neighboring hydroxyl can guide an incoming thiolate anion via a transient charge-dipole interaction (hydrogen bond) resulting in a highly selective process that enhances the effectiveness of such agents (Figure 7.4.B).

In addition, a sterically hindered electrophile may exercise a superior degree of selectivity toward uniquely reactive thiolates (e.g., Keap1 cysteines). For example, both the high effectiveness and low toxicity profile of synthetic oleanane triterpenoids has been ascribed to a special combination of high intrinsic electrophilicity of the Michael acceptor activated by both keto and cyano groups (Figure 7.4.C) and sterics, specifically the presence of a Michael acceptor functionality directly adjacent to a highly congested quaternary carbon center [100] (Figure 7.4.D). Congestion in the vicinity of the reactive  $\beta$ -carbon reduces the reactivity of the Michael acceptor by crowding the transition state, where trigonal planar geometry is converted into a more sterically demanding tetrahedral state. This should allow only the most reactive thiolates to overcome the steric barrier and form covalent adducts. It must be noted, however, that mere reduction of reactivity without adjustment of other factors is expected to result in a situation where the enhanced selectivity and reduced toxicity come at the expense of potency. Thus, of two stereoisomers of guggulsterone, a terpenoid that lacks any known toxicity, only the E-form displays moderate ARE induction [101], due to, presumably, a poor accessibility by nucleophiles of the trisubstituted  $\alpha,\beta$ unsaturated ketone furnished with a quaternary center as the  $\alpha$ -carbon substituent. The relatively low toxicity of coffee furans, as compared to other furans, can also be explained

by steric factors, if the operative electrophile is a conjugated epoxydiene formed directly from kahweol (Figure 7.3.C) or after oxidative processing of cafestol.

The reversibility of addition, the fourth factor on our list, is emerging as a crucial characteristic of highly effective inducers with low associated toxicity [102]. Clearly, the entropically-favored reversal of electrophilic additions is expected to *i*) alleviate many potentially adverse outcomes associated with unintended alkylation events, *ii*) activate multiple stress sensors at a greatly reduced concentration, and *iii*) exhibit catalytic turnover (recycling) in maintaining the state of adaptive stress. ARE inducers must compete with prominent blood proteins—hemoglobin and serum albumin—and glutathione, among many other potential nucleophiles. Glutathione, the most available cellular thiol responsible for detoxification, can be conjugated with electrophiles in an uncatalyzed manner, or the conjugation can be catalyzed by glutathione S-transferases (GST) (reviewed in [103]). Once formed, if sufficiently stable, these conjugates are excreted from the cell by trans-membrane multidrug resistance-associated proteins (MRPs), leading to dilution of the ARE inducer and the signaling event (Figure 7.2). Thus, highly reversible ARE inducers may escape elimination from the cell by de-conjugating from GSH prior to being transported by an MRP, maintaining their cellular concentration more effectively than ARE inducers that form much more stable conjugates. For "reversible" ARE inducers, conjugation with glutathione has been hypothesized to be an intermediate step [104], leading via continuous deconjugations to activation of less represented but highly reactive "sensory" thiolates, responsible for antioxidant effects (Figure 7.2).

For example, the reversible addition of sulforaphane to thiols has long been suspected to be responsible for pleiotropic nature of its cellular activity [78, 105]. In addition, the dramatically enhanced Nrf2-induction potential of CDDO and its variants over traditional phytochemical inducers may be due in large part to the facile reversibility of their thiol conjugates [100]. While a systematic analysis of chemical features responsible for the reversibility of thiol conjugations would benefit the field, certain generalizations about these features can be made herein. Thus, unusual stabilization of the electrophile through crossconjugation (Figure 7.4.E) or extension of conjugation (Figure 7.4.F) could be invoked in a few cases documenting the favorability of addition reversal [34, 52, 100]. Both effects should lead to the enhanced delocalization of the partial positive charge induced by the associated carbonyl, reducing the electrophilicity of  $\beta$ - ( $\delta$ -, etc.) carbon centers. For example, in celastrol, a quinone methide triterpene that has sub-micromolar potency for ARE activation, extended conjugation can account for the observed reversibility of its reaction with GSH [106]. Similarly, reversibility due to enhanced stabilization of the electrophile could explain the reduced toxicity and unique effectiveness of quercetin, when compared to structurally related flavonoids and flavonols in particular. Thus, the glutathionyl adducts of the quercetin-derived quinone methide have been shown to be reversible on the order of minutes to hours [34]. In addition, we suggest that reversibility of addition may play an important role in shifting the effectiveness-to-toxicity balance in yetto-be established cases. For example, the divergent toxicity profiles of I3C and DIM (see section 7.3.2.1) could be explained by the reversibility of the DIM-derived thiol conjugate due to conjugation extension by a second aromatic group.

Alternatively, the destabilization of an adduct through steric congestion at positions adjacent to the nascent C–S bond may also facilitate elimination, as carbon hybridization changes from trigonal planar to tetrahedral. Thus, we hypothesize that the unusually low toxicity of CDDO [100] is likely the result of addition reversibility due to steric congestion in the corresponding adduct (Figure 7.4.D). In contrast, ARE inducers that do not form relatively reversible adducts, such as the EGCG *o*-quinone, for example, may be eliminated from the cell rather quickly through sequestration by GSH and subsequent elimination by an MRP. For these compounds, generation of superoxide and ROS as signaling intermediates may represent a more effective pathway for Nrf2 activation (Figure 7.2).

#### 7.3.4 Mechanisms of Nrf2 Activation

7.3.4.1 Modification of Keap1 Cysteines: Identification of Key Sensors for ARE **Inducers**—A key mechanism by which phytochemicals activate Nrf2 is revealed by their common ability to react with cysteine residues. As mentioned in Section 7.3.2.1, the Nrf2repressor protein Keap1 is particularly cysteine-rich. Keap1 directly binds to Nrf2 [31] and represses the transcription factor in at least two ways. First, Keap1 contains a Crm1dependent nuclear export signal sequence that appears to be important for maintaining Nrf2 primarily in the cytoplasm under basal conditions [107-109]. In addition, and likely most important for phytochemical signaling, Keap1 serves as a bridge between Nrf2 and the Cul3-E3-ubiquitin ligase complex, leading to ubiquitination of Nrf2 and subsequent degradation by the 26S proteasome (reviewed in [110]). There are two separate sites in Nrf2 that bind to Keap1, termed the ETGE and DLG motifs [111]. Keap1 forms a homodimer through its N-terminal BTB domain, and it is the C-terminal Kelch domain of Keap1 that binds to the Nrf2 ETGE and DLG motifs. The seven lysines of Nrf2 that are targets for Cul3-mediated ubiquitination are located between the ETGE and DLG motifs [112]. Binding of the two Kelch domains of a homodimer of Keap1 to the ETGE and DLG sites of an Nrf2 protein molecule is believed to be critical for presentation of those lysines to an E2 ligase for ubiquitination [113]. Ubiquitination-directed degradation maintains the Nrf2 protein at a low level under normal conditions. Upon exposure to stress or compounds that modify cysteines, Nrf2 escapes Keap1 repression, and thereby is no longer ubiquitinated and degraded. This in turn leads to Nrf2 accumulation, resulting in activation of ARE-regulated genes.

Two primary methods have been used to illustrate which of the Keap1 cysteines (27 in human Keap1) are implicated in sensing ARE inducers: overexpression of Keap1 containing cysteine mutated to serine or alanine in mammalian cell lines or zebrafish, and mapping of cysteine modification in the purified protein by peptidic digestion and mass spectrometry. These complementary methods have revealed that a "cysteine code" may exist, in which particular types of ARE inducers react most readily with particular Keap1 cysteines. Cysteines 226 and 613 were shown to sense heavy metals [114], while C273 and C288 have been implicated in sensing alkenals, some cyclopentenone prostaglandins and nitro-fatty acids [114-116]. Keap1 C151 has been shown to be required for sensing many electrophilic ARE inducers [115, 117-121]. For example, a potent imidazole derivative of CDDO (CDDO-Im), tBHQ, a quinone-forming phenol, and sulforaphane are all highly dependent on C151 to upregulate the ARE [107, 115, 120]. Finally, the importance of Keap1 C151 in

sensing ARE inducers is illustrated by the Keap1 C151S transgenic mouse [122]. This mouse is not only less responsive to tBHQ for upregulation of cytoprotective enzymes but also the basal levels of these enzymes are repressed [122]. Thus Keap1 C151 emerges as the key sensor for both endogenous agents as well as phytochemicals. The role of this cysteine as a sensor has been proposed to have evolved as a means of sensing nitric oxide [114]. Interestingly, mutation of C226 and C623 to serine has no effect on ARE activation by sulforaphane or tBHQ [114]. Therefore C151 is currently the only known Keap1 sensor cysteine for a phytochemical, sulforaphane, and a synthetic quinoid-forming polyphenol, tBHQ.

The role of Keap1 C151 as a principal sensor for phytochemicals is further supported by peptide mapping studies. Keap1 C151 is the only cysteine consistently and highly modified by all phytochemicals tested thus far: isoliquiritigenin, 10-shogaol, xanthohumol [123] and sulforaphane [124] (Figure 7.5). There have been discrepant results for modification of Keap1 cysteines, in particular C151 by sulforaphane. Initially, peptide-mapping studies of human Keap1 cysteines modified by sulforaphane found C151 to be one of the least readily modified cysteines (Figure 7.5)[125]. Based on the high dependence of sulforaphane ARE activation on C151, as well as the known labile, reversible nature of dithiocarbamates [126], i.e. the products of sulforaphane reaction with thiols, further experiments were conducted using a streamlined method to limit the reverse reaction after labeling Keap1 cysteines with sulforaphane [124]. Under these conditions, C151 emerged as one of the four most readily modified cysteines of Keap1. The adduction of Keap1 C151 by sulforaphane was also observed indirectly for Keap1 overexpressed in Cos1 cells using a biotin-switch technique, where C151 appeared to be modified to a greater extent than other cysteines [114]. The entire pattern of modification of Keap1 by sulforaphane generally shows large variability depending on the method used (Figure 7.5 and [124]), in particular as to whether C151 is detected as modified, illustrating the importance of considering the reversible nature of electrophile-cysteine adducts. Interestingly, several cysteines were detected as readily modified regardless of the method used, including C77, C226, C368, and C489 (Figure 7.5), indicating that certain dithiocarbamate-cysteine adducts are much more stable than others. The C151 dithiocarbamate adduct, on the other hand, is both one of the most reactive and reversible. Modeling of the BTB domain of Keap1 containing C151 [114, 120] depicts various residues nearby, including K131, R135, K150 and H154, that could participate both in lowering the pKa of C151 and providing acid/base catalysis for promoting both the forward (addition) and reverse (elimination) reactions. As a confirmation, a single mutation of K150 to threonine significantly reduces the response to C151-dependent ARE inducers [115].

While the modification pattern of a quinoid-forming polyphenol phytochemicals has not yet been assessed, the oxidized form of tBHQ was found to react with Keap1 C151, shown directly for purified Keap1 [127] (Figure 7.5) and indirectly for Keap1 overexpressed in Cos1 cells [114]. Interestingly, C151 is not found to be readily modified by the oxidized form of GSH, GSSG [82] (Figure 7.5). Phytochemicals for which ROS production is a key means of activating Nrf2 (Section 7.3.2.3) likely activate Nrf2 by first generating GSSG (Figure 7.2). Thus, their Nrf2 activation would presumably be much less dependent on Keap1 C151.

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While C151 is the cysteine most readily modified by phytochemicals based on current knowledge, other cysteines are sites of phytochemical addition in purified Keap1 protein (Figure 7.5). For example C319 is readily modified by all four phytochemicals shown in Figure 7.5 as well as oxidized glutathione. To our knowledge this cysteine has not yet been evaluated as a sensor for phytochemicals, and it has been shown not to be required to sense the soft cations, such as  $As^{3+}$  and  $Se^{4+}$ , but there is an indication that it may play a role in sensing the  $Zn^{2+}$  ion [114].

Notably, none of the phytochemicals evaluated in Figure 7.5 modified C273 or C288 appreciably, although glutathione modified C288 to some extent. These cysteines are considered to be potential sensors, as mutation of these cysteines to serine renders Keap1 unable to repress Nrf2 [107, 122]. As these cysteines have been implicated in sensing more reactive Michael acceptors—alkenals, prostaglandins and nitro-fatty acids [114-116]—they may serve to detect primarily these endogenous signaling agents, rather than more stable phytochemical electrophiles. Keap1 C226 and C613, which sense heavy metal salts and other soft cations [114], are readily modified by phytochemicals, with isoliquiritigenin and sulforaphane targeting the former and xanthohumol, 10-shogaol, and to some extent, sulforaphane targeting the latter. However, as described above, mutation of C226 or C613 to serine had no effect on the ability of Keap1 to sense sulforaphane [114]. It is likely that the C151 residue present in these mutant proteins could be modified by sulforaphane, leading to a loss of Keap1 repression of Nrf2. Further work is required to ascertain whether modification of Keap1 cysteines other than C151 contributes to Nrf2 activation by phytochemicals.

#### 7.3.4.2 Mechanism of Nrf2 activation upon modification of Keap1 Cysteines-

The means by which Nrf2 escapes Keap1-Cul3-directed ubiquitination and degradation is still under active investigation. Disruption of the Keap1-Nrf2 interaction upon modification of Keap1 cysteines, allowing Nrf2 to escape Keap1, was originally proposed on the basis of experiments using recombinant proteins [126]. However, further investigations of a similar nature showed that the overall affinity of the Keap1-Nrf2 complex is maintained after modification of Keap1 at sensor cysteines, including C151 and C288, by ARE inducers including sulforaphane and isoliquiritigenin [128]. While a reduced affinity of Keap1 for Nrf2 after treatment of cells with tBHQ has also been observed by pull-down assays [129], other groups have reported that Keap1 and Nrf2 remain associated after treatment of cells with either tBHQ [107, 130], sulforaphane [107], a synthetic triterpenoid derivative, dh404 [131], or quercetin [132]. Pull-down assays are a useful but imprecise means of determining relative affinities of protein-protein interactions, making it difficult to ascertain whether the observed perturbation of the Keap1-Nrf2 complex occurs in response to inducers in a cellular context. It is noteworthy that preventing Nrf2 degradation is sufficient to promote Nrf2 nuclear accumulation and ARE activation [133]. It has been hypothesized that the affinity of the weaker Nrf2 DLG site is decreased after modification of Keap1 cysteines including C151, C273 and/or C288, leading to decreased ubiquitination, without affecting substantially the overall stability of the complex [122].

Significantly, the stability of the Keap1–Cul3 interaction, also essential for ubiquitination, is reduced upon treatment with ARE inducers as observed by pull-down assays [112, 134].

These studies show that the reduced affinity is dependent on Keap1 C151, as mutation of C151 to serine largely abolished the effect. Modification of the C151 residue has clearly been shown to disrupt Keap1-mediated ubiquitination of Nrf2, as mutation of C151 to tryptophan had the same effect as electrophile treatment of cells [120]. The Keap1 C151W mutant was largely unable to mediate Nrf2 ubiquitination, leading to stabilization of Nrf2 with concurrent activation of ARE-regulated genes, and had reduced affinity for Cul3, as assessed by the pull-down assay. Remarkably, Nrf2 binding to Keap1 in this assay was unaffected by the tryptophan mutation. While other Keap1 C151W in cells, as well as in a zebrafish model [115], show that modification of Keap1 C151 alone is sufficient for signaling for Nrf2 activation.

It has been proposed that the negative charge of the C151 thiolate anion is essential for Nrf2 turnover, and that modification of C151 by an electrophile neutralizes the negative charge, leading to the signaling event [114]. However, a series of 13 mutations at position 151 showed that the size of the residues was the key determinant, rather than hydrophobicity or charge, with the largest residue, tryptophan, showing the largest effect [120]. In particular, asparagine, with a relatively small partial molar volume (i.e. size) and no capacity to carry a negative charge, rendered the corresponding Keap1 mutant effective in suppressing Nrf2. Therefore, the size of the residue at position appears to be the major determining factor for Keap1's ability to repress Nrf2. Since all phytochemicals that were found to react with the C151 thiolate have partial molar volumes larger than that of tryptophan, they would all be large enough to trigger the signaling mechanism.

Alternatively, formation of disulfide bonds between Keap1 cysteines has been proposed to mediate loss of Nrf2 repression. Disulfide-linked Keap1 dimers have been observed in extracts of cells treated with the chalcone derivative *bis*(2-hydroxybenzylidene)acetone, which is a phenolic Michael acceptor [135]. In addition, treatment of cells with oxidizing agents (e.g., H<sub>2</sub>O<sub>2</sub>, nitric oxide, hypochlorous acid, or S-nitrosocysteine) leads to a Keap1 dimer linked through C151 [136]. Non-enzymatic oxidation of phenolic compounds will generate phenoxyl radicals of variable stability (Figure 7.2), and it has been proposed that these reactive intermediates could mediate the formation of thiyl radicals, promoting the formation of a disulfide bond or bonds between Keap1 cysteines [78]. Furthermore, both ROS and GSSG could also participate in Keap1 disulfide bond formation (Figure 7.2). In addition, a Keap1 dimer that was stable under reducing conditions was induced by treatment of cells with a quinoid-forming polyphenol, tBHQ [107]. Formation of this non-disulfidelinked dimer was dependent on C151. Radical-mediated hydrogen abstraction from surface tryptophans or tyrosines by subsequent cross-linking may account for the formation of the non-reducible dimer [137]. Further studies are required to determine on a structural/ molecular level how direct modification of C151 and other Keap1 residues by ARE inducers or GSSG, or Keap1 dimer formation, lead to the loss of its ability to repress Nrf2.

#### 7.3.4.3 Role of Protein Kinases in Nrf2 Activation by Phytochemicals-In

addition to the direct Keap1-repression-inactivation mechanism discussed above, a number of kinases have been implicated in Nrf2 activation. The kinase pathways that currently appear to be the most important for Nrf2 activation by phytochemicals are the ones mediated

by phosphatidylinositol 3-kinase (PI3K)/Akt and mitogen-activated protein kinases (MAPKs). The PI3K/Akt pathway mediates carnosol-induced expression of the cytoprotective enzyme HO-1 in rat pheochromocytoma PC12 cells [138]. In addition, EGCG, guggulsterone and piceatannol activate expression of cytoprotective enzymes including HO-1 in the human mammary epithelial (MCF10A) cells in a PI3K/Akt–dependent manner [101, 139, 140]. EGCG activation of Nrf2 in BAECs is also PI3K/Akt dependent [89]. The Nrf2 activation by CDDO-Im is dependent upon PI3K/Akt activity in ARPE-19 retinal epithelial cells [141].

The mechanism by which Nrf2 is regulated by PI3K/Akt has been investigated in some detail. The GSK-3 $\beta$  kinase, which is inactivated upon phosphorylation by Akt1, phosphorylates and negatively regulates Nrf2 via two distinct mechanisms [142, 143]. The phosphorylation of Nrf2 by GSK-3 $\beta$  leads to both cytoplasmic localization of Nrf2 [142], and Keap1-independent ubiquitination by the SCF/ $\beta$ -TrCP/Cul1 complex, leading to Nrf2 degradation [143]. A key cysteine residue, C124 in PTEN, a negative regulator of the PI3K/Akt axis [144], is directly modified by biotinylated CDDO in ARPE-19 cells [141], and the same cysteine is required for guggulsterone-induced Nrf2 accumulation in MCF10A cells [101]. Remaining to be demonstrated is whether the modification of PTEN C124 by electrophilic agents also inactivates GSK-3 $\beta$ , and this in turn leads to Nrf2 accumulation and Nrf2 nuclear localization.

While the inhibition of GSK-3 $\beta$  is implicated in the initial response to Nrf2 activators, the re-activation of GSK-3 $\beta$  has been proposed to occur several hours later, leading to eventual downregulation of the response by increasing Nrf2 nuclear export [145]. In this proposed mechanism, H<sub>2</sub>O<sub>2</sub> treatment of cells activates GSK-3 $\beta$ , leading to subsequent activation of Fyn/Src kinases, which then phosphorylate Nrf2 Y568. Phosphorylation of Y568 enhances the interaction of Nrf2 with the Crm1 nuclear export protein. This mechanism has not yet been evaluated for phytochemical ARE inducers. However, these investigations were prompted by the observation that genistein, an ARE inducer, is a tyrosine kinase inhibitor. Whether inhibition of Nrf2 Y568 phosphorylation is in part or whole responsible for the ARE induction by genistein or other natural phenols is unknown.

The role of MAPKs in activation of Nrf2 is somewhat controversial. A large number of studies with various phytochemicals, including sulforaphane, phenethyl isothiocyanate (PEITC), curcumin, quercetin and EGCG, have implicated MAPKs in Nrf2 activation (reviewed in [146]). In order to assess the importance of this kinase family in Nrf2 signaling, MAPKs were overexpressed in HEK293T cells, and potential sites of MAPK-dependent phosphorylation on Nrf2 were identified by mass spectrometry. As a result, five phosphorylated residues were identified [146]. An Nrf2 mutant with all five sites mutated to alanine showed a moderate decrease in the transcriptional activity of Nrf2, concomitant with a slight reduction in its nuclear accumulation. However the stability of Nrf2 protein, which is primarily controlled by Keap1, was not affected. The general conclusion drawn from this work is that direct phosphorylation of Nrf2 by MAPKs has limited contribution to modulating Nrf2 activity [110, 146]. However, the Nrf2 protein with five mutated residues was only evaluated for blocked phosphorylation by overexpression of JNK2, rather than each MAPK relevant for Nrf2 activation [146]. Therefore, an additional site on Nrf2 beyond

those five identified by mass spectrometry may in fact be a relevant target. Regardless of whether direct Nrf2 phosphorylation is the means by which MAPKs activate Nrf2, the sheer number of studies that implicate this pathway in Nrf2 activation by phytochemicals calls for further mechanistic investigation.

Activation of Nrf2 is also mediated by other kinases such as PERK [147, 148] and casein kinase 2 [149, 150]. However, to our knowledge, these pathways have not been evaluated for phytochemical activation of Nrf2. In addition, protein kinase C (PKC) isoforms are involved in the upregulation of cytoprotective genes by several phytochemicals. PKC-dependent ARE gene upregulation was observed for curcumin in HUH7 hepatoma cells and human monocytes [151, 152], piceatannol in BAECs [153], and epigallocatechin in human monocytic cells [154]. Nrf2 nuclear accumulation and ARE activation are also PKC-dependent for tBHQ in HepG2 and rat H4IIEC3 hepatoma cell lines [129, 155, 156], as well as nontumorigenic human keratinocytes (HaCaT cells) [157]. However, ARE activation by quercetin or kaempferol was not dependent on PKC in Hepa1c1c7 hepatoma cells [90], nor was diallyl trisulfide ARE activation dependent on PKC in HepG2 cells [158]. Further investigations are required to ascertain under what circumstances a phytochemical might activate Nrf2 through PKC isoforms.

#### 7.4 Summary and Future Directions

Nrf2 is emerging as a master control of cytoprotective mechanisms important for defense against environmental challenges and stress factors. Phytochemical activation of Nrf2 promises to be an important mechanism in the prevention and amelioration of a wide variety of human diseases. In addition, Nrf2 is under investigation as a drug target [159], and there is much to be learned from investigations of phytochemical ARE inducers with regards to what makes an effective agent with minimal toxicity. A large effort has been put towards understanding the mechanisms by which Nrf2 is activated, although phytochemicals are still relatively understudied compared to other compounds [160].

In general, a common trait of agents that activate Nrf2 effectively is the ability to react with thiols either directly or upon bioactivation (spontaneous or enzyme-catalyzed). Alternative mechanisms of Nrf2 de-repression involving induction of oxidative stress, for example, have also been discussed. The electrophilic and/or oxidative nature of Nrf2 activators means that a balance between toxic and protective effects is necessary for safe administration of these agents. This balance is described in conventional drug development as a "therapeutic window" or effective dosage range. Within this range, beneficial effects are exhibited at optimal strength without being compromised by adverse processes. Our primary recommendation for future studies in phytochemical activation of Nrf2 is that, as with conventional drugs, efforts are put towards understanding how this dosage range is widened. This will enable identification or development of more effective agents for disease prevention and strategies for administering those agents.

Widening this dosage range does not appear to be a trivial exercise. The challenge lies in the fact that a strategy for increasing potency that merely relies on higher reactivity of these electrophiles is likely to enhance the extent of deleterious events. These may range from

effect dilution through GSH conjugation followed by efflux to severe cytotoxicity associated with covalent modification of unintended targets. At the same time, reduction of toxicity through attenuation of electrophilicity would be expected to compromise potency. However, nature provides some important examples of how an effective therapeutic window can be achieved, which involves both partners in the modification reaction: sensor cysteines and naturally occurring cytoprotective agents.

The chemistry of the sensor cysteines plays an important role in ensuring advantageous biological effects of phytochemicals. The unique reactivity of soft thiolates of the sensor cysteines implicated in ARE induction, including Keap1 C151, ensures a high degree of "chemoselectivity" through the intrinsic preference for  $\pi$ -conjugated and thereby polarizable (soft) electrophilic centers, such as those found in effective phytochemical activators of Nrf2 (Figures 1 and Section 7.3.3). Chemoselectivity is further ensured by the high reactivity of the sensor cysteines implicated in ARE induction, including Keap1 C151. What remains to be explored is whether yet further preference toward particular sensor cysteines could be provided by the remaining structural features of the covalent modifiers, such as overall size or presence of particular functionalities that could interact favorably with the protein environment surrounding the thiol.

Certain phytochemical ARE inducers, either relatively potent ones or those with low or no observable toxicity, also provide clues as to features that widen the therapeutic window. Thus, preference for sensor thiolates can be enhanced by neighboring hydroxyl groups guiding a thiolate toward an attack on an electrophile. Alternatively, since sensor thiolates are so reactive, poor Michael acceptors could conjugate to them preferentially and, thereby, act as viable ARE inducers, while evading less reactive "off-target" nucleophiles. Examples of phytochemicals that are expected to be relatively unreactive Michael acceptors include those deactivated by alkene polysubstitution, bulky neighbors, and/or extended conjugation (Section 7.3.3). Remarkably, these deactivating factors are also expected to facilitate the reverse elimination process either by enhancing the stability of the electrophile, or by destabilizing the adduct. The reversibility of conjugates is emerging as an important feature for high potency and low toxicity. As discussed in Section 7.3.3, continuous de-conjugation may simultaneously contribute to both toxicity reduction due to instability of off-target conjugates and potency enhancement through efflux prevention. In addition, there is much to learn as to how modification of Keap1 cysteines leads to Nrf2 de-repression, for example, whether the size of modifiers influences the extent of the response, as suggested by a mutagenesis study [120].

Another factor important in the therapeutic window for phytochemicals is the ability to induce Nrf2 by a variety of pathways. Key sensor cysteines of phytochemical ARE inducers are being identified—Keap1 C151 and C124 of PTEN kinase—and other cysteines will likely emerge as important in sensing ARE inducers. Depending on the nature of the chemical and biological events associated with a given phytochemical in a given condition, different signaling "nodes" can be activated, all leading to Nrf2-ARE regulation. For example, the ultimate messenger molecule producing the Nrf2-dependent signal may not only be the phytochemical itself and/or its metabolite, but a concomitantly formed ROS/RNS or GSSG, as the GSH:GSSG ratio drops. These secondary signaling messengers

target not only Keap1 and PTEN, but also numerous other kinases, including those involved in Nrf2 regulation. For example, various flavonoids including quercetin and EGCG can noncovalently associate with and modulate a number of protein kinases with high affinity (reviewed in [161]). There appears to be an intricate network of regulatory proteins that modulate Nrf2 activity, and multiple sensing steps can lead to the ability of Nrf2 upregulation to be tuned as is appropriate. Importantly, the presence of such a network implies that, if multiple nodes are activated by a single phytochemical, the convergence of the responses may lead to a more pronounced effect. This rationale also provides the basis for the unexpected effectiveness exhibited by mixtures of Nrf2 activators. Indeed, impressive synergism has been observed in a few studies conducted thus far, including for glucosinolate break-down products [54] and a commercially-available phytochemical supplement mixture [162]. Nrf2 activation, while clearly important, is certainly not the only mediator of the cytoprotective and disease-preventive attributes of phytochemicals (reviewed in [163]). Thus, phytochemicals that modulate a number of targets in disease prevention within their "therapeutic window" will likely be most effective in humans.

In summary, despite the multitude of cellular targets affected by phytochemicals, upregulation of Nrf2 is emerging as a key factor in their cytoprotective properties. And while it is unlikely that a single phytochemical or even a plant source will emerge as a magic bullet for disease prevention or amelioration, we expect that unraveling the chemical and biological aspects of their action may lead to unprecedented opportunities. These prospects could range from dietary/supplement recommendations to phytochemical "cocktails" specially formulated for synergistic effects and to nature-inspired synthetic molecules harnessing the most effective features of the plant-derived prototypes.

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#### Fig. 7.1.

Structures of phytochemicals arranged by the chemistry that leads to Nrf2 activation. All compounds shown activate Nrf2. Thiol-reactive chemical motifs are shown in red, and "additive" distribution of electron-donating groups, which can be oxidized to quinoids, are shown in blue.



Note: Keap1 thiols in scheme could be replaced with those of other sensor proteins: PTEN C124, etc.

## $R^{\bullet} \rightarrow RH = DIRECT ANTIOXIDANT EFFECT (Free radical trapping)$ $O_2 \rightarrow O_2^{--} = PROOXIDANT EFFECT (ROS generation)$ Nrf2 Activation = INDIRECT ANTIOXIDANT EFFECT (Upregulation of cytoprotective genes)

#### Fig. 7.2.

The three roles a quinoid-forming polyphenol (represented by the R-catechol) can play: prooxidant, direct antioxidant, and indirect antioxidant. As a *direct antioxidant*, in the presence of a high concentration of free radical species, a polyphenol can trap the radical, forming a relatively stable radical species. As a *prooxidant*, in the presence of catalytic amounts of a transition metal, a polyphenol can promote the formation of superoxide and other ROS, enroute to formation of a Michael acceptor. An alternate path to oxidation of the polyphenol is catalyzed by a metalloenzyme and occurs without the production of ROS. Once the quinoid group is formed, the Michael acceptor group can react with a thiolate molecule. There is evidence that a quinone reacts with a key Keap1 sensor cysteine, C151, leading to Nrf2 activation, as described in the text. Upon activation, Nrf2 upregulates a battery of antioxidant enzymes and other cytoprotective enzymes, known as the *indirect antioxidant* effect. Reaction of the quinoid with GSH and subsequent elimination from the cell will lead to dilution of sensor cysteines, or formation of Nrf2 activation by radicals or ROS not depicted are oxidation of sensor cysteines, or formation of disulfides among sensor cysteines.

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#### Fig. 7.3.

Biotransformations implicated in conversions of quercetin (**A**), DAS (**B**), coffee triterpenoids (kahweol shown here) (**C**), and I3C and DIM (**D**) into thiol-reactive conjugated electrophiles: quinoids (quercetin), sulfone (DAS), epoxide or  $\gamma$ -ketoenal (kahweol) and indolenines (I3C & DIM). The established or proposed reactive groups are highlighted in red.



#### Fig 7.4.

Structural features of Michael acceptors affecting the rates of forward and reverse reactions with thiols. Keap1 C151 is shown as an example thiol. (A) Brønsted acid catalysis by a neighboring residue (Keap1 K150). (B) Neighboring group (proximity) catalysis through hydrogen bonding with a hydroxyl adjacent to the  $\beta$ -carbon. (C) Alkene polyactivation, favoring addition. (D) Steric congestion, preventing additions by less reactive nucleophiles via transition state crowding. Cross-conjugation (E) and extended conjugation (F) stabilize an electrophile by diluting the partial positive charges at the electrophilic centers and thereby reducing its reactivity and promoting reversibility.

Cysteine		XAN	ISO	SHO	SUL[124] SUL[125] tBQ			GSSG
-	C13	0	0	0	0	0	0	0
N-term	C14	0	0	0	0	0	0	0
	C23	0	$\bigcirc$	0	0	0		0
	C38	$\bigcirc$	0	0		0	0	0
BTB	C77	0		0			0	
	C151					0		0
	C171	0	0	0	0	0	0	0
Intervening region	C196	0	$\bigcirc$	0	0	0	0	0
	C226	0	•	0			•	0
	C241	0	0		0	0	0	0
	C249	0	0		0		0	0
	C257	O	0		0	0	0	0
	C273	0	0	0	0	0	0	0
	C288	0	0	0	0	0	0	
	C297	0	0	0	0	0	0	
	C319					0	0	
Kelch	C368	0	0					
	C395	0		0	0	0	0	0
	C406	0	0	0		0	0	0
	C434		0			0	0	
	C489		0	0			0	0
	C513	0		0	0		0	0
	C518	0		0	0		0	0
m	C583				0		0	0
	(613		0			0	$\mathbf{O}$	
C-té	C624		0		0		$\mathbf{O}$	
-	C024		$\mathbf{O}$		$\mathbf{O}$		$\mathbf{O}$	$\bigcirc$

#### Fig. 7.5.

Keap1 cysteines readily modified by phytochemicals, glutathione, and a quinone-forming polyphenol. Abbreviated name, chemical name and reference: XAN, xanthohumol [123]; ISO, isoliquiritigenin [123]; SHO, 10-shogaol [123]; SUL, sulforaphane [124, 125]; tBQ, *tert*-butylquinone [127]; and GSSG, oxidized glutathione [82]. Cysteines are ranked in order of most readily modified for all but tBQ, as determined by increasing concentrations of the electrophile, with the darkest circles being the most readily modified, and the empty circle indicating weakly modified or not modified cysteines. For tBQ, the cysteines are not ranked.

#### Table 1

Structures of Phytochemicals that Upregulate Cytoprotective Enyzmes through Nrf2

	Structural/Functional	Representatives	Plant Sources		
	Classes				
	Chalcone, MAª, phenol Chalcone, MA, phenol	Isoliquiritigenin Xanthohumol	Licorice, shallot, tonka bean Hops		
Polyphenols	Cinnamate, 1,2-diphenol Cinnamate, 1,2-diphenol Diarylheptanioid MA 1.2-diphenol	Caffeic acid Ferulic acid Curcumin	Lignin-containing plants Apples, cabbages, plums Tumeric		
	Diarylheptanioid, MA, 1,2-diphenol Masked 1,2-diphenol	Yakuchinone B Capsaicin	Alpinia oxyphylla Miquel Spicy peppers		
	Terpenoid 1,2-diphenol Flavanol, 1,2,3-triphenol	Carnosol EGCG	Rosemary Green tea		
	1,2-Diphenol Flavonoid mixture	Ellagic acid Silymarin (mixture)	Berries Milk thistle		
	Masked 1,2-diphenol, MA	Silibin (single compound) 10-Shogaol Apigenin	Ginger Chamomile thyme		
	Flavone, phenol Isoflavone, phenol	Luteolin Genistein	Celery, thyme Soy		
	Isoflavone, phenol Flavonol, 1,2,6-triphenol	Biochanin A Quercetin	Red clover Apples, capers		
	Flavonol, 1,6-diphenol Flavonol, 1,2,6-triphenol	Kaempferol Fistein	Apples, broccoli, tea Acacia, mangos		
	Flavonol, phenol Trans-stilbene, phenol Trans-stilbene, 1 2-diphenol	Galangin Resveratrol Piceatannol	Alpinia officinarum Grape skins, berries Grape skins, berries		
	Withanolide, MA	18-Hydroxywithanolide D	Tomatillo		
	Withanolide, MA	Withaphysacarpin	Tomatillo Deprea subtriflora		
	Coumarin, MA(?)	Coumarin	Leguminosae spp.		
	Butenolide, MA(?)	β-Angelica lactone	Archangelica officinalis		
	Coumarinoid, MA(?)	Auraptene	Citrus		
	Terpenoid, furan	Catestol	Green coffee beans		
	Terpenoid MA	Zerumbone	Tropical ginger		
	Terpenoid, MA	Citral (geranial)	Lemongrass		
	Terpenoid, MA	Celastrol	Thunder of God vine (T. wilfordii)		
	Terpenoid (carotene)	Lycopene	Tomatoes		
	Phytosterol, MA	E-Guggulsterone	Commiphora mukul Sida acuta		
	Organosulfide	Allicin	Garlic		
	Organosulfide	Diallyl disulfide	Garlic		
	Organosulfide	Diallyl trisulfide	Garlic		
	Organosulfide	Dilallyl sulfide	Garlic		
	Organosulfide	S-allyl cysteine	Garlic		
i gluco- olates	Isothiocyanate	Sulforaphane Phonothyl ITC <sup>b</sup>	Broccoll		
	Isothiocyanate	6-Methylsulfinylhexyl ITC	Wasabi		
	Indoles	Indole-3-carbinol -> DIM <sup>c</sup>	Brussels sprouts, cabbages		
om	Indoles	Brassinin	Chinese cabbage		
Ē Ĩ	Epithionitrile	1-Cyano-2,3-epithiopropane	Cabbages		
	Epithionitrile	1-Cyano-3,3-epithiobutane	Cabbages		

<sup>a</sup>MA, Michael acceptor

<sup>b</sup>ITC, isothiocyanate

<sup>c</sup>As described in the text, indole-3-carbinol is a precursor for diindolylmethane (DIM), which is considerably less toxic.