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The cancer chemopreventive actions of phytochemicals derived from glucosinolates

■ **Abstract** This article reviews the mechanisms by which glucosinolate breakdown products are thought to inhibit carcinogenesis. It describes how isothiocyanates, thiocyanates, nitriles, cyanoepithioalkanes and indoles are produced from glucosinolates through the actions of myrosinase, epithiospecifier protein and

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epithiospecifier modifier protein released from cruciferous vegetables during injury to the plant. The various biological activities displayed by these phytochemicals are described. In particular, their abilities to induce cytoprotective genes, mediated by the Nrf2 (NF-E2 related factor 2) and AhR (arylhydrocarbon receptor) transcription factors, and their abilities to repress NF- κ B (nuclear factor- κ B) activity, inhibit histone deacetylase, and inhibit cytochrome P450 are outlined. Isothiocyanates appear to alter gene expression through modification of critical thiols in regulatory proteins such as Keap1 (Kelch-like ECH-associated protein 1) or IKK ($I\kappa B$ kinase), causing activation of Nrf2 and inactivation of NF- κ B, respectively. Certain indoles act as ligands for AhR.

Isothiocyanates and indoles are also capable of affecting cell cycle arrest and stimulating apoptosis. The mechanisms responsible for these anti-proliferative responses are discussed.

■ **Key words** antioxidant response element apoptosis – arylhydrocarbon receptor - cytochrome P450 epithionitriles - gene induction glucosinolates - glutathione S-transferase – isothiocyanates – $NF-\kappa B - Nrf2 - quinone reduc$ tase - xenobiotic response element

Glucosinolates and their association with cancer chemoprevention

Regular consumption of cruciferous vegetables, such as broccoli, Brussels sprouts, cabbage, cauliflower, kale, swede and turnip, is associated with a reduced incidence of cancer [129, International Agency for Research on Cancer Workgroup [55]]. Furthermore, greater health benefit may be obtained from raw as opposed to cooked vegetables [72]. In man, these vegetables appear to protect against colorectal cancer [104], lung cancer [73], and possibly prostate cancer

[42]. In animals, feeding experiments have also suggested broccoli can protect against liver cancer [109]. Cruciferous vegetables uniquely contain glucosinolates at approximately 20 µmol/g dry mass of vegetable [23, 66], and it is thought that these phytochemicals are primarily responsible for the putative cancer chemoprevention conferred by eating diets that contain significant quantities of these vegetables

Glucosinolates are substituted β -thioglucoside *N*-hydroxysulfates, formed by the plant from any one of eight amino acids, namely, alanine, valine, leucine, isoleucine, phenylalanine, methionine, tyrosine and §

Fig. 1 Synthesis of Glucosinolates. The R group is derived from the original amino acid (i.e. Ala, Val, Leu, Ile, Phe, Met, Tyr or Trp) and is highly variable

tryptophan [55]. Over 115 naturally occurring glucosinolates have been identified. Each cruciferous vegetable contains a mixture of glucosinolates that varies according to the strain of the plant [23, 35, 74, 89, 110]. The glucosinolate content is primarily under genetic control, with the last step in the pathway being of particular importance [128], though it can also be influenced by environmental factors [15, 36]. Much of the diversity amongst glucosinolates arises from the addition of different sized alkyl groups to the side chain of amino acids, principally valine, phenylalanine and methionine, used in their biosynthesis; this variable elongation of amino acid side chains entails repetitive additions of methyl groups through a series of transamination, condensation, isomerisation and decarboxylation reactions [43]. As shown in Fig. 1, the synthesis of glucosinolates proceeds through the conversion of elongated amino acids to their oxime derivatives, catalysed by members of the cytochrome P450 (CYP) 79 family [3]. Subsequently, the oxime is metabolised to a thiohydroximate, which is in turn conjugated with glucuronic acid to form a desulfoglucosinolate before finally being sulfated to yield the glucosinolate [55].

The task of establishing a link between the ingestion of particular glucosinolates and their possible health benefits is not straightforward. This endeavour is simplified to some extent by the fact that relatively few glucosinolates are present in the human diet. The most common of these are the methylsulfinylalkyl glucosinolates glucoiberin and glucoraphanin, the olefinic glucosinolates sinigrin, gluconapin, glucobrassicanapin and progoitrin, and the aromatic glucosinolate gluconasturtiin (Table 1) [66, 118]. Glucoraphanin has been reported to be abundant in broccoli [66], though certain strains of this plant

Table 1 Trivial names of some glucosinolates with the corresponding sidechain (R) composition

Name	R side-chain	
Sinigrin Gluconapin Glucobrassicin Glucobrassicanapin Progoitrin Glucoiberin Gluconapoleiferin Glucocheirolin Glucoerucin Glucoberteroin	2-Propenyl 3-Butenyl 3-Indolylmethyl 4-pentenyl 2-Hydroxy-3-butenyl 3-Methylsulfinylpropyl 2-Hydroxy-4-pentenyl 3-Methylsulfonylpropyl 4-Methylthiobutyl 5-Methylthiopentyl	

also contain substantial amounts of glucoiberin [87]. Sinigrin has been reported to be the predominant glucosinolate in Brussels sprouts, cabbage, cauliflower and kale [66]; gluconapin is also found in high levels in Brussels sprouts [66]. Substantial amounts of progoitrin are present in many cruciferous vegetables [66]. The aromatic glucosinolate gluconasturtin is present in watercress. The indolyl glucosinolate glucobrassicin is present in Savoy cabbage, Brussels sprouts and cauliflower [66, 83], and whilst not abundant it can elicit distinct pharmacological effects.

Production of isothiocyanates, thiocyanates, nitriles, cyano-epithioalkanes and oxazolidine-2-thiones from glucosinolates

Inhibition of carcinogenesis by glucosinolates is not primarily attributable to this class of compound, but rather it appears to be due to certain of their breakdown products. Hydrolysis of these phytochemicals is catalysed by myrosinase (β -thioglucoside glucohydrolase, EC 3.2.3.1), an enzyme that is physically segregated from glucosinolates within the intact plant by virtue of the fact that it is sequestered in specialised "myrosin" cells [5]. Upon wounding of the vegetable, for example during harvesting, during freeze-thawing, during food preparation, or during chewing whilst being eaten, myrosinase is released from the "myrosin" cells and catalyses the hydrolysis of glucosinolates within the damaged plant. In addition, myrosinase activity may be present in human colonic microflora, suggesting that it is possible glucosinolates are hydrolysed in the gastrointestinal tract during digestion of food [6, 32, 65]. Myrosinase cleaves glucosinolates at the thioglycoside linkage to produce glucose and an unstable aglycone thiohydroximate-O-sulfonate that spontaneously rearranges to yield several breakdown products. The outcome of the reaction with myrosinase depends on the nature of the aglycone, as well as the reaction temperature, the pH and the presence of ferrous ions (Fig. 2).

The thiohydroximate-O-sulfates formed from methylsulfinylalkyl, olefinic and aromatic glucosinolates undergo a Lossen rearrangement, with the elimination of sulfate, to form their respective isothiocyanates (ITCs), thiocyanates or nitriles [6, 37]. Elemental sulfur is also formed in certain circumstances. At neutral pH, hydrolysis of glucosinolates with aliphatic or aromatic side chains gives rise primarily to ITCs. The glucosinolates glucoiberin, gluconapin, glucoraphanin, glucobrassicanapin and sinigrin yield 3-methylsulfinylpropyl-ITC, 3-butenyl-ITC, 4-methylsulfinylbutyl-ITC (sulforaphane), 4-pentenyl-ITC and 2-propenyl-ITC (allyl-ITC), respectively.

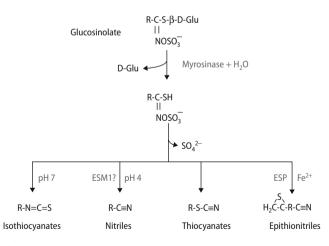


Fig. 2 Hydrolysis of Glucosinolates. At high or neutral pH the formation of isothiocyanates is favoured while at low pH the formation of nitriles is favoured. Epithiospecifier protein in the presence of Fe²⁺ ions interacts with myrosinase to promote the transfer of the sulfur to the alkenyl group from the S-Glucose of the terminally unsaturated Glucosinolate [6]

At low pH, the thiohydroximate-O-sulfates formed by myrosinase from glucosinolates with a side chain containing a double bond (e.g. sinigrin, gluconapin and glucobrassicanapin) may, in the presence of an epithiospecifier protein (ESP) and ferrous ions, give rise to a cyano-epithioalkane [102]. In this case, ESP interacts with myrosinase to promote sulfur transfer from the S-glycosyl unit to the alkenyl chain derived from the amino acid part of the aglycone [39]. Thus, at pH 4 and in the presence of Fe²⁺ ions, myrosinase and ESP convert sinigrin to 1-cyano-2,3-epithiopropane [68]; Fig. 3 shows hydrolysis products produced from sinigrin. Gluconapin can similarly be converted by the combined actions of myrosinase and ESP to 1-cyano-3,4-epithiobutane [17, 63]. Likewise, glucobrassicanapin can be hydrolysed to 1-cyano-4, 5-epithiopentane [17]. Progoitrin, a (2R)-hydroxy-3butenyl glucosinolate, is converted in the presence of myrosinase, ESP and Fe²⁺ ions to an epithionitrile [76]. In the case of epi-progoitrin [(2S)-hydroxy-3butenyl glucosinolatel, it can be hydrolysed by myrosinase to crambene (1-cyano-2-hydroxy-3-butene) [24, 40]. Two cDNAs for ESP have been cloned from Arabidopsis and broccoli, and the purified proteins characterized following their heterologous expression in E. coli [82, 139]. In Arabidopsis, an epithiospecifier modifier (ESM) gene has been reported that inhibits formation of the epithionitrile and favours production of nitrile [144].

If the aglycone generated by myrosinase is from a glucosinolate with a side chain lacking a double bond, the sulfur atom may be lost and a nitrile formed [69,

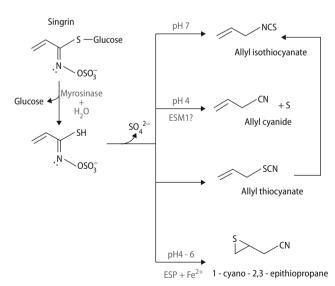


Fig. 3 Hydrolysis of Sinigrin. Following damage to the plant tissue, the glucosinolate sinigrin is hydrolysed by myrosinase resulting in the formation of four distinct compounds

Fig. 4 Production of Indoles from Glucosinolates. At neutral pH the hydrolysis of glucobrassican by myrosinase leads to the formation of an unstable isothiocyanate intermediate that degrades to form Indole-3-carbinol and a thiocyanate ion

80, 90]. This reaction may involve ESP, and is diminished by heating [81]. A few glucosinolates produce thiocyanates though the mechanism involved is unclear [5, 6]. Upon hydrolysis by myrosinase, those aglycones from glucosinolates that contain β -hydroxylated side-chains form oxazolidine-2-thiones, as a consequence of spontaneous cyclization. Examples of these include progoitrin, glucoconringiin and gluconapoleiferin [55].

Production of indoles from glucosinolates

The indolyl glucosinolates glucobrassicin and neoglucobrassicin are synthesised by the plant from tryptophan. The best studied of these is glucobrassicin. At neutral pH, hydrolysis of glucobrassicin by myrosinase does not generate an ITC, but rather gives rise to indole-3-carbinol and a thiocyanate ion (Fig. 4); this reaction probably proceeds through a Lossen rearrangement generating an unstable ITC intermediate [83]. At acidic pH, hydrolysis of glucobrassicin yields indole-3-acetonitrile, hydrogen sulfide and elemental sulfur [83]. Formation of indole-3-acetonitrile requires both myrosinase and ESP [9]. In the acidic environment of the stomach, indole-3-carbinol condenses to form various compounds including indolo[3,2-b]carbazole and 3,3'-diindolylmethane, both of which have potent pharmacological effects [7, 25]. It can also combine with ascorbic acid to form ascorbigen [105] (Fig. 5).

Fig. 5 Structures of indoles produced from Glucosinolates - 3,3'-Diindolylmethane (DIM), Indolo[3,2-b]carbazole (ICZ) and Ascorbigen (ASG)

Chemopreventive mechanisms stimulated by glucosinolate hydrolysis products

In view of the diverse spectrum of chemicals generated from glucosinolates by the actions of myrosinase, ESP and ESM, it is not surprising that a number of

distinct cancer chemopreventive mechanisms have been proposed to account for the putative anti-cancer properties of cruciferous vegetables. These include induction of antioxidant and detoxification genes through activation of Nrf2 (NF-E2 related factor 2) and AhR (arylhydrocarbon receptor), inhibition of pro-inflammatory reactions by repression of NF- κ B (nuclear factor-κB), inhibition of cytochrome P450 (CYP) enzyme activity, inhibition of histone deacetylase, and stimulation of cell cycle arrest and apoptosis. There is a dose-dependency in these responses: generally, induction of cytoprotective genes and inhibition of CYP activity occurs at relatively low concentrations of phytochemical, whereas activation of cell cycle arrest and apoptosis occurs at higher levels of phytochemical [7, 62]. A major problem exists in interpreting experiments utilizing vegetable extracts because of their inherent variable composition and thus uncertainty about attributing biological effects to specific phytochemicals. It is also apparent that uncertainties exist about the bioavailability of many glucosinolate breakdown products that further complicates interpretation of in vitro data [48]. A challenge in evaluating the literature arises from the emphasis placed on certain glucosinolate breakdown products and the dearth of data relating to others. Thus, there is a relative abundance of information about isothiocyanates and indoles, when compared with the data about thiocyanates, nitriles, cyanoepithioalkanes, and oxazolidine-2-thiones. Considerable evidence has, for example, been presented that the ITC sulforaphane can inhibit experimental chemical carcinogenesis in animal models at various sites including mammary gland [142], stomach [33] and skin [133]. Sulforaphane can also protect against familial adenomatous polyposis in the intestine of ApcMin/+ mice [113]. Also, broccoli extracts enriched with sulforaphane can prevent UV-light initiated skin carcinogenesis [29], and sulforaphane activates cell defences against UV radiation [121]. Comparatively little testing of the chemopreventive properties of other glucosinolate-derived phytochemicals in animal models has been reported, though there is a body of literature that indicates indoles can exert promoting effects in experimental chemical carcinogenesis (see ref [55] for a review).

Induction of gene expression mediated by Nrf2

Isothiocyanates increase the expression of antioxidant and detoxication proteins, such as glutathione S-transferase (GST) and NAD(P)H:quinone oxidoreductase 1 (NQO1), both in vivo and in vitro [96, 147]. Agents such as ITC that increase GST and NQO1

1
$$CH_{2}$$
 CH_{2} $N=C=S$

2 CH_{2} $N=C=S$

3 CH_{2} $N=C=S$

4 CH_{3} S CH_{2} $N=C=S$

5 CH_{3} S CH_{2} $N=C=S$

6 CH_{3} S CH_{2} S $N=C=S$

Fig. 6 Structures of isothiocyanates which induce NQ01: **1-** Allyl ITC, **2-** Phenethyl ITC, **3-** Benzyl ITC, **4-** 3-methylsulfinylpropyl ITC, **5-** 7-methylsulfinylpropyl ITC, **6-** 8-methylsulfinyloctyl ITC

enzymes without increasing arylhydrocarbon hydroxylase activity, catalysed by the phase I drugmetabolising enzyme CYP 1A1, are sometimes called mono-functional inducers [106]. Sulforaphane, 3-methylsulfinylpropyl-ITC, allyl-ITC, 7-methylsulfinylheptyl-ITC, 8-methylsulfinyloctyl-ITC, benzyl-ITC and phenethyl-ITC, induce NQO1 in the mouse Hepa-1c1c7 hepatoma cell line [27, 146] (Fig. 6). Many of these compounds induce GST P1-1 in the rat liver RL34 cells [91]. The mouse ngo1 gene contains a functional antioxidant response element (ARE, minimal enhancer 5'-A/GTGÂC/GNNNGCA/G-3'), sometimes called an electrophile response element (EpRE), in its 5'-upstream region [100], as does the rat GSTP1 gene (in which it was originally called glutathione transferase P enhancer I, GPEI) [44]. The induction of these two genes by ITCs is mediated by the Nrf2 bZIP (basic-region leucine zipper) transcription factor that is recruited to the ARE as a heterodimer with a small Maf protein, MafF, MafG or MafK [100]. As far as is known, all genes that are induced by ITCs contain an ARE in their promoters and are regulated by Nrf2. Examination of $nrf2^{-/-}$ and wild-type mice, suggests

adaptor protein [64, 143], and carbamylation of

certain critical residues (most frequently thought to

be Cys-273 and Cys-288, though there is not a con-

sensus view on this point, c.f. [31, 49, 126, 141]) in

the central intervening region (IVR) of Keap1 leads

to failure of the Cullin-3:Rbx1/Keap1 complex to ubiquitylate Nrf2. Within the cell, Keap1 is a dimeric

zinc-containing protein [28] and binds Nrf2 through

both of its C-terminal Kelch-repeat domains inter-

acting simultaneously with a low-affinity DLG motif

and a high-affinity ETGE motif in the N-terminal

Neh2 domain of the bZIP factor [86]; see Fig. 7 for a cartoon of the two-site interaction between the Keap1

dimer and Nrf2. As a consequence of this two-site

interaction, the lysine acceptor sites in Nrf2 that lie

between the DLG and ETGE motifs are immobilized

and presented to Cullin-3:Rbx1 in an orientation that

allows their ubiquitylation. Under normal homeo-

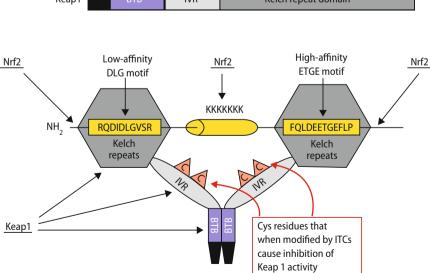
CNC-bZIP

AREs are present in the promoters of at least 100 genes. The battery of genes regulated by Nrf2 includes those for antioxidant proteins, drug-metabolising enzymes, drug efflux pumps, heat shock proteins, and α and β subunits of the 26S proteasome [54, 67, 84, 124]. Specifically, some of the most inducible genes in rodent and human cells are those encoding aldo-keto reductase (AKR), carboxyl esterase, ferritin, glutamate cysteine ligase catalytic (GCLC) and modifier (GCLM) subunits, GST, heme oxygenase 1, NQO1, metallothionein, microsomal epoxide hydrolase, multidrug resistance-associated protein, thioredoxin, thioredoxin reductase and UDP-glucuronosyl transferase [26, 54, 67, 84, 124]. Many of these genes are induced by sulforaphane in vivo in an Nrf2-dependent fashion in the stomach, small intestine and liver of rodents [34, 51, 54, 84, 124]. Importantly, feeding broccoli seed to mice increased the levels of GCLC, GST and NQO1 in the gastrointestinal tract in an Nrf2-dependent fashion [87].

It is thought that ITCs possess the ability to induce ARE-driven gene expression because they are thiolactive [30]. Through this characteristic, it is highly probable that dietary ITCs modify cysteine residues in many proteins. They also react with glutathione [140] and therefore presumably produce redox stress by altering the intracellular GSH:GSSG ratio. Most significantly in terms of eliciting an adaptive response to such stress, sulforaphane forms adducts with cysteines in Keap1 (Kelch-like ECH-associated protein 1) [31, 49], a Cullin-3:Rbx1 E3 ubiquitin ligase substrate

static conditions, this process is highly efficient and results in rapid proteasomal degradation of the transcription factor. It appears most likely that modification of Keap1 by sulforaphane causes a conformational change in the substrate adaptor that prevents it from binding both the low-affinity DLG and high-affinity ETGE motifs in Nrf2. Through modification by ITC, it seems likely the Cullin-3:Rbx1/Keap1/Nrf2 complex becomes "stalled", with Nrf2 only being bound to one of the Kelch-repeat domains via its high-affinity ETGE motif [86]. This will probably result in Keap1 becoming saturated, thereby allowing newly translated Nrf2 to avoid Neh2 Nrf2 **IVR** Kelch repeat domain Keap1

Fig. 7 The domain structures of Nrf2 and Keap1, and the complex formed between the two proteins. The cartoon shows interaction between the two Kelchrepeat domains found in the dimeric Keap1 protein with the low-affinity DLG motif and the high-affinity ETGE motif found in the Neh2 domain of Nrf2 [86]. The seven lysine residues located between the DLG and ETGE motifs in the Neh2 domain that serve as ubiquitin acceptor sites are shown. The Cvs residues in the IVR of Keap1 that are modified by ITCs are also depicted



capture by Keap1 and accumulate within the cell. Consistent with this view, exposure of RL34 cells or HepG2 cells to sulforaphane causes stabilisation and rapid accumulation of Nrf2 [57, 85]. Surprisingly, allyl-ITC does not increase Nrf2 stability [57] and therefore other factors may be involved in enzyme induction by these phytochemicals. Treatment of cells with benzyl-ITC causes a rapid increase in the level of reactive oxygen species, and this could also contribute to gene induction [97].

Up-regulation of thioredoxin and thioredoxin reductase (TrxR1) by sulforaphane is controlled at several levels [2, 145]. Induction of mRNA for TrxR1 by sulforaphane occurs through an ARE in its gene promoter, presumably mediated by Nrf2. However, as TrxR1 is a seleno-protein, the consequence of increased gene transcription can be augmented at the translational level by sodium selenite, and other forms of selenium, through providing an adequate supply of SeCys for incorporation into protein and also by delaying degradation of TrxR1 mRNA [145].

It has been found that administration of cranbene to Fischer 344 rats causes an elevation in hepatic GST and NQO1 enzyme activities, but not CYP1A1, suggesting it is a mono-functional inducer [78]. By comparison with sulforaphane, cranbene was found to be an approximately equally potent inducer in the rat. However, it was not a particularly effective inducer of NQO1 activity in Hepa-1c1c7 cells suggesting that the relatively high potency of induction observed in vivo is due to bio-transformation of cranbene to a thiol-active metabolite [61]. The identity of this metabolite is not known.

The indole-containing glucobrassicin breakdown products can activate gene expression by several mechanisms. Indole-3-carbinol is a modest inducer of Nrf2-dependent ARE-driven gene expression in the liver and small intestine of mice [12, 84]. Although indole-3-acetonitrile has not been studied in Nrf2 knockout mice, it is a good inducer of GST enzyme activity in wild-type mouse liver and small intestine [45], a fact that infers it works through Nrf2.

The activity of the Nrf2 transcription factor can be inhibited by retinoic acid through a protein-protein interaction between Nrf2 and retinoic acid receptor alpha that prevents the bZIP protein from binding to the ARE [127]. Similarly, the nuclear orphan receptor estrogen-related receptor beta also negatively regulates Nrf2 [149]. At present it is not known whether dietary retinoids and their precursors, or dietary estrogens, can block the benefits of ITCs in vivo, but this will be an important point to clarify as the presence of high levels of retinoic acid could represent a confounding factor in clinical intervention studies.

Induction of gene expression mediated by AhR

The major effect indole-3-carbinol has on gene expression arises because it can condense in acid conditions to form indolo[3,2-b]carbazole and 3,3'diindolylmethane. Both these condensation products induce CYP1A1 genes via the xenobiotic response element (XRE, 5'-TA/TGCGTGA/C-3') in their promoter regions because they are ligands for the arylhydrocarbon receptor (AhR), a basic helix-loop-helix transcription factor. Examination of the dose of indole required to double XRE-driven reporter gene expression showed that indolo[3,2-b]carbazole is a much more potent inducing agent than 3,3'-diindolylmethane, indole-3-carbinol or ascorbigen [4, 7]. Amongst genes for drug-metabolising enzymes, mouse, rat and human CYP1A1 are prototypic XREregulated genes. This cytochrome has O-deethylase activity towards ethoxyresorufin, and there is abundant evidence from enzyme assays, western or northern blotting that CYP1A1 is inducible by indolo[3,2-b]carbazole and 3,3'-diindolylmethane [55]. The promoters of other genes including rat and human NQO1, rat ALDH-3, rat GSTA2, rat UGT1A1, and rat UGT1A6 contain an XRE, as does the mouse BAX gene, and it is therefore anticipated that activation of AhR by glucobrassicin-derived indoles will influence significantly the metabolism of xenobiotics; for a review, see [99] and for a more recent report of gene expression profiling to identify AhR target genes, see

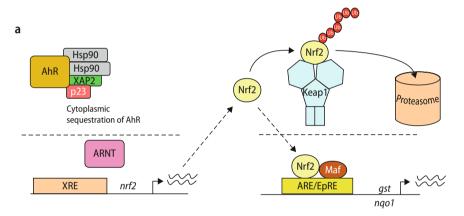
Significantly, the mouse *nrf2* gene promoter also contains an XRE, and AhR ligands such as dioxin produce an increased production of Nrf2 mRNA [88] (Fig. 8). As Nr2 is regulated primarily at the level of protein stability [85], the increase in mRNA will not in itself cause an increase in ARE-driven gene expression unless the rate of translation of Nrf2 is sufficiently high to saturate the Cullin-3:Rbx1/Keap1 complex. However, the combination of an AhR ligand, such as indolo[3,2-b]carbazole, plus a redox stressor, such as an ITC, should result in a more marked increase in Nrf2 protein than occurs through redox stress alone; Fig. 9 shows how indoles and ITCs could act synergistically. Whether indolo[3,2-*b*]carbazole and 3,3'-diindolylmethane can increase both Nrf2 mRNA levels and the stability of Nrf2 protein is not known, but unless these phytochemicals are metabolised by CYPs, and thereby generate reactive oxygen species, this seems unlikely [98].

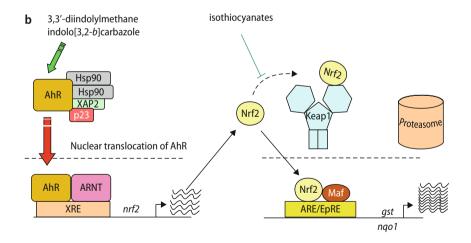
In addition to up-regulation of CYP1A1 by indoles, CYP1B1 and CYP19 are inducible [112], as are the drug-metabolising enzymes AKR, GST T1-1, sulfotransferase and UGT1 [71]. Furthermore, 3,3'-diindolylmethane can induce the transcription

Fig. 8 Presence of an XRE in the promoter of the *nrf2* gene. The sequence data for the XRE in the promoter of *nrf2* are taken from ref [88]. The other XRE-containing sequences are taken from ref [99]

Species	Gene	Element	5'-USR	enhancer	3'-USR	3' nucleotide
mouse	nrf2	XREL1	gtttg	c AGCGTG g	actca	-706
mouse	ngo1	XRE	tcccc	TAGCGTGC	aaagg	-374
rat	NQO1	XRE	tcccc	TTGCGTGC	aaagg	-360
human	NQO1	XRE	attac	ag GCGTGA	gcacc	-730
rat	ALDH-3	XRE	gccgc	CTGCGTGA	ctgca	-376
mouse	cyp1a1	DRE1	gagge	TAGCGTGC	gtaag	-903
mouse	cyp1a1	DRE2	ccage	TAGCGTGA	cagca	-1052
mouse	cyp1a1	DRE3	cggag	TTGCGTGA	gaaga	-976
rat	CYP1A1	XRE1	cggag	TTGCGTGA	gaaga	-1020
rat	CYP1A1	XRE2	ccage	TAGCGTGA	cagca	-1079
human	CYP1A1	XRE1	aggcg	TTGCGTGA	gaagg	-991
human	CYP1A1	XRE2	ccccc	TCGCGTGA	ctgcg	-1048
rat	GSTA2	XRE	gcatg	TTGCGTGC	atccc	-893
rat	UGT1A1	XRE	agaat	gTGCGTGA	caagg	-123
human	UGT1A6	XRE	agaac	TOGCGTGC	agcag	-1492
XRE 'core' consensus				TAGCGTGA		
				T C		

Fig. 9 Cross-talk between the AhR-XRE pathway and the Nrf2-ARE pathway. **a** Under normal homeostatic conditions, Nrf2 is primarily ubiquitylated and degraded through the 26S proteasome, with only a relatively small fraction of the total translated protein being recruited to the promoters of ARE-driven genes. **b** The combined effect of i) indoles inducing transcription of the *nrf2* gene through AhR and ii) isothiocyanates preventing ubiquitylation of the increased load of translated Nrf2 protein, by inhibition of Keap1, could theoretically result in hyper-induction of ARE-driven genes





factors ATF3, c-Jun and NF-IL6 as well as genes involved in cell growth such as growth arrest and DNA damage (GADD) GADD34, GADD45 and GADD153 [1, 11]. Also, p21 is induced by indole-3-

carbinol [19]. It is not however clear whether the genes for ATF3, c-Jun, and NF-IL6, and the various GADD genes, are regulated through XREs in an AhR-dependent fashion.

It has been argued that induction of CYP1A1 by indoles is potentially deleterious to the cell because the cytochrome can activate polycyclic aromatic hydrocarbons to ultimate carcinogens. This viewpoint is probably an oversimplification and does not take into account the multiple changes in gene expression that indoles affect. Bonnesen et al. [7] have reported that treatment of human colon LS-174 cells with indolo[3,2-b] carbazole before exposure to benzo[a]pyrene provides a small measure of protection against DNA damage as measured by a Comet assay. Most importantly, prior treatment of the LS-174 cells with both indolo[3,2-b]carbazole and sulforaphane before exposure to benzo[a]pyrene was found to confer substantial protection against genotoxicity, and this protection was greater than was achieved by either phytochemical alone [7].

It is interesting to note that some cross-talk appears to exist between the Nrf2 and AhR pathways insofar as AhR requires the presence of Nrf2 in order to induce the expression of NQO1 by dioxin or 3-methylcholanthrene [75, 101]. As mentioned above, the promoter of mouse *nrf2* contains several XREs [88] and the promoter of mouse *ahr* contains an ARE [114]. Thus AhR regulates Nrf2 mRNA levels, and Nrf2 regulates AhR mRNA levels. It remains to be tested whether combined exposure of cells to AhR ligands such as indolo[3,2-b]carbazole along with stabilizers of Nrf2 protein such as ITCs will have synergistic effects on both ARE-driven gene expression and XRE-driven gene expression.

Inhibition of pro-tumourigenic processes by glucosinolate breakdown products

Inflammation is a well-recognized risk factor in carcinogenesis. Isothiocyanates possess anti-inflammatory activity through inhibiting NF- κ B [47, 119]. The ability of sulforaphane and phenethyl-ITC to inhibit the transcriptional activity of NF- κ B is a consequence of the phytochemicals antagonising phosphorylation of I κ B, the inhibitor of NF- κ B, which is carried out by IKK [134]. Inhibition of NF- κ B prevents transcriptional activation of genes such as cyclin D1, VEGF, Bcl-X_L, COX2 and MMP-9. Although these gene products influence various biological processes, inhibition of NF- κ B generally appears to make cells more sensitive to apoptosis. Interestingly, circumin and cyclopentenone prostaglandins, such as 15-deoxy- $\Delta^{12,14}$ -prostaglandin J_2 , also antagonize NF- κ B by inhibiting IKK [16, 103]. As circumin and cyclopentenone prostaglandins contain an α,β -unsaturated carbonyl group they are, like ITCs, thiolreactive. It has been proposed that 15-deoxy-∆12,14prostaglandin J_2 inactivates IKK by modifying Cys-179 [111], and it would be interesting to know if ITCs modify the same cysteine residue.

Certain isothiocyanates can block the activation of several carcinogens to their ultimate carcinogenic forms. Tumorigenesis caused by the carcinogens 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone and N-nitrosobis-(2-oxopropyl)amine can be prevented by phenethyl-ITC and this involves inhibition of activation of pro-carcinogens by CYP isoenzymes [46, 95]. Inhibition of CYP can also be achieved by sulforaphane [18, 77]. In the case of the nitrosamine 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone, the ITCs with highest lipophilicity and low reactivity of their NCS group had the greatest ability to inhibit lung tumourigenesis [58].

Inhibition of histone deacetylase by isothiocyanates

The isothiocyante sulforaphane has been shown to inhibit histone deacetylase (HDAC) [93, 94]. This function is likely to alter gene expression substantially. It may also have profound implications for cell fate as a change in the balance between histone acetyl transferase (HAT) and HDAC could alter tumourigenesis. Indeed, recognition of this possibility has lead to considerable recent interest in the ability of HDAC inhibitors to act as both chemopreventive and chemotherapeutic agents.

Within the cell, DNA is tightly coiled around an octamer of histone proteins in a structure known as a nucleosome, the basic structural unit of chromatin. Each of the histone proteins contains an evolutionary conserved amino tail protruding from the nucleosome, which can determine the accessibility of the DNA to transcription factors. The tail is also subject to many post-translational modifications including acetylation. The addition of an acetyl group to the histone tail results in a conformational change which enables the tail to move away from the DNA allowing transcription factors access to interact with the DNA. Conversely, removal of acetyl groups causes the tail to wrap tightly around the DNA thereby preventing interaction with the transcription machinery. The addition and removal of the acetyl groups is carried out by HAT and HDAC, respectively. In pre-cancerous and cancerous cells, tumour suppressor genes are associated with deacetylated histones resulting in the inactivation of these genes. Inhibition of HDAC may prevent the removal of acetyl moieties from histones, thus allowing transcription of the tumour suppressor genes.

Sulforaphane has been shown to diminish HDAC activity with a concomitant rise in histone acetylation

in prostate cancer cells [93], human embryonic kidney cells [94] and human colorectal cancer cells [94]. The link between inhibition of HDAC activity and the resultant increase in transcription of tumour suppressor genes has been reported for p21 [92, 93], p53 [38] and Bax [92, 93]. Equally, mammalian HDAC is capable of down-regulating p53 function, by deactylation of the *p53* gene, resulting in a reduction in its transcriptional activity. In addition, sulforaphane has been reported to cause a G₂/M phase delay with an increase in apoptotic cell fraction in a time and dose dependent fashion [60].

The ability of sulforaphane, along with other dietary HDAC inhibitors such as diallyl disulfide [22], to alter chromatin structure is likely to be of considerable biological significance but its contribution to chemoprevention is poorly understood.

Stimulation of cell cycle arrest and apoptosis by isothiocyanates

Many of the naturally occurring ITCs can suppress the growth of cultured tumour cells by modulating multiple targets that influence cell cycle arrest, apoptosis and differentiation [148]. However, the majority of the studies into mechanisms by which this class of chemical inhibit cell growth have focussed on sulforaphane and phenethyl-ITC. For example, Apc Min/+ mice treated with sulforaphane at either 300 or 600 ppm in their diet have been reported to develop fewer and smaller polyps in their small intestine than Apc Min/+ mice on a control diet; this was associated with a higher level of apoptosis and lower cell proliferation in animals on the ITC-containing diet [52].

In human PC-3 prostate cancer cells, treatment with sulforaphane or phenethyl-ITC causes an arrest in G₂/M phase of the cell cycle that is associated with a decrease in levels of cyclin B1 and cell division cycle (Cdc) 25B and Cdc25C proteins [115, 130]. The loss of Cdc25C was reported to be due to proteasomal activity, and was accompanied by its translocation from the nucleus to the cytoplasm [115]. Relocation of Cdc25C was controlled by its phosphorylation at Ser-216, mediated through activation of checkpoint kinase 2 (Chk2). Cell proliferation by ITCs may also be achieved by distrupting cytoskeletal structure and tubulin polymerisation [56, 117].

Administration of ITCs to cells at growth suppressive concentrations results in the rapid generation of reactive oxygen species (ROS), within 1 h of exposure, which appears to be necessary for cell death [115, 116]. The generation of ROS by ITCs is accompanied by depletion of intracellular GSH and is

achieved through the rapid export of ITC-glutathione and ITC-cysteinylglycine conjugates via MRP1 and Pgp-1 efflux pumps [10]. Consistent with the view that production of ROS is necessary for apoptosis, overexpression of catalase suppresses ITC-initiated apoptosis, as does pre-treatment with N-acetylcysteine [115]. Furthermore, addition of GSH subsequent to ITC treatment can block apoptosis [62, 136]. Treatment with ITCs leads to a loss of mitochondrial membrane potential and release of cytochrome c from mitochondria [123]. There is evidence that ITCs can activate both the intrinsic and extrinsic caspase cascades, though this may be cell specific. For example, in PC-3 cells, sulforaphane can increase Fas protein levels and activate caspase-8 whilst simultaneously targeting mitochondria and activating caspase-9 [115]. In human bladder cancer UM-UC-3 cells, benzyl-ITC and phenethyl-ITC are more effective in activating caspase-9 than caspase-8 [122]. By contrast, in human leukaemia HL60 cells, caspase-8 plays a major role in apoptosis stimulated by phenethyl-ITC [135]. The levels of pro-apoptotic proteins Bak and Bax, which neutralize the antiapoptotic effects of Bcl-2, are increased by phenethyl-ITC and sulforaphane in PC-3 prostate cancer cells, and this may lead to induction of Apaf-1 [115, 130, 132]. Furthermore, the pro-apoptotic proteins Bok and Bim EL are also induced by ITCs, and this is thought to amplify the effects of Bak and Bax [115]. Besides increasing the levels of these pro-apoptotic proteins, ITCs downregulate the anti-apoptotic proteins Mcl-1 and Bcl-x₁, though the effect is cell-specific [132]; for a review of regulation of apoptotic pathways by BCL-2 family proteins see [138]. Various ITCs have been shown to activate c-Jun N-terminal kinase (JNK) [13, 137], and this is mediated by extracellular signal-regulated kinases, ERK1/2 [131]. The use of inhibitors has indicated that JNK is essential for phenethyl-ITC to cause cytochrome c release and caspase-3 activation in human HT-29 colon adenocarcinoma cells [53]. However, the mechanism by which JNK activates caspases remains unclear.

Stimulation of cell cycle arrest and apoptosis by indoles

Treatment of human MCF-7 breast cancer cells with $100 \mu M$ indole-3-carbinol inhibits proliferation through affecting a G_1 cell cycle arrest [20]. This may in part be due to 3,3'-diindolylmethane rather than indole-3-carbinol as significant quantities of the indole spontaneously condense to the dimer in culture conditions [120]. Cell cycle arrest at G_1 occurs as a consequence of indole-3-carbinol inhibiting both

cyclin-dependent kinase (CDK) 2 and CDK6. In the case of CDK6, expression of the gene is reduced because indole-3-carbinol attenuates recruitment of the Sp1 transcription factor to the CDK6 promoter [21]. Furthermore, in HaCaT keratinocytes, treatment with 400 µM indole-3-carbinol induces the CDK4/6 inhibitor p15 INK4b mRNA and protein causing hypophosphorylation of Rb protein [79]. It therefore appears that cyclin D-CDK6 activity can be inhibited by dual mechanisms, though the concentrations of indole involved seem rather high. In the case of CDK2, indole-3-carbinol has been reported to decrease the kinase activity in MCF-7 cells and inhibit phosphorylation of Rb protein [11]. The reduction in CDK2 activity is attributed to a selective alteration in the size of the complex in which it is contained, from an active form within a 90 kDa complex to a lower activity form within a 200 kDa complex [41]; the 90 and 200 kDa complexes include forms of cyclin E that differ in size, and the larger complex also contains an additional 75 kDa cyclin E immunoreactive protein. Furthermore, the reduction in CDK2 activity is accompanied by redistribution of the kinase in the 200 kDa complex from the nucleus to the cytoplasm, suggesting that indole-3-carbinol can influence the nucleocytoplasmic shuttling of the kinase [41].

In cells that contain wild-type p53, such as MCF-10A, treatment with 300 μ M indole-3-carbinol or 30 μ M 3,3'-diindolylmethane has been found to result in activation of the ATM signalling pathway, an increase in p53 protein levels, and induction of p21 [8]. These changes result in prevention of the CDK2-mediated G₁/S transition in the cell cycle [8].

Indole-3-carbinol, but not 3,3'-diindolylmethane, was found to inhibit expression of the androgen receptor in human lymph node carcinoma of prostate (LNCaP) cells as well as the probable downstream target gene prostate specific antigen [50]. It is possible that down-regulation of this receptor represents an antiproliferative mechanism in prostate cells.

Indoles can affect apoptosis in breast and prostate cancer cells. Treatment of PC-3 prostate cancer cells with 60 μ M indole-3-carbinol inhibits the EGF-induced autophosphorylation of PI3K and Akt [14]. Thus, the Akt/PI3K cell survival pathway appears to be targeted by indole-3-carbinol. Also, nuclear translocation of NF- κ B is inhibited by 3,3'-diindolylmethane through a reduction in phosphorylation of I κ B α [107, 108].

In HCT-116 human colon cells, indole-3-carbinol can induce nonsteroidal anti-inflammatory drugactivated gene-1 (NAG-1), a TGF- β family member associated with pro-apoptotic activities [70]. This may also mediate the anti-tumour effects of indoles.

Concluding comments

It is becoming clear that glucosinolate breakdown products can influence the initiation and progression of carcinogenesis. They also appear to influence apoptotic responses to chemotherapeutic agents, such as tamoxifen [19]. A major impediment to our understanding of the chemopreventative mechanisms stimulated by glucosinolates is that relatively little is known about the biological effects of glucosinolate breakdown products other than isothiocyantes and the indole-containing derivatives. Specifically, there are little data about chemopreventative activities of thiocyanates, nitriles, cyano-epithioalkanes and oxazolidine-2-thiones. It is unclear whether formation of thiocyanates, nitriles, cyano-epithioalkanes and oxazolidine-2-thiones from glucosinolates, at the expense of forming isothiocyanates, is undesirable from a cancer chemoprevention perspective. It is unclear whether the activity of ESP, which reduces the formation of isothiocyanates from glucosinolates, is undesirable. If so, ESP should possibly be eliminated by genetic means from commercial crops. Furthermore, relatively little is known about the pharmacokinetic properties of glucosinolate breakdown products in the human, and without this information it is difficult to relate responses of cells in culture to certain concentrations of phytochemical to the in vivo situation. These are areas that warrant further examination.

Mammalian cells display marked dose responsiveness to phytochemicals: at low doses of phytochemical, cytoprotective adaptive responses are activated, whereas at higher doses cell cycle arrest and apoptosis occurs. It is presently unclear how these different types of response are co-ordinated by the cell and how decisions about whether adaptation, growth arrest or apoptosis is chosen as the appropriate response are determined. Identification of mechanisms that control such outcomes will be useful.

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