

FORUM REVIEW ARTICLE

Mitochondria-Centric Review of Polyphenol Bioactivity in Cancer Models

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Abstract

Significance: Humans are exposed daily to polyphenols in milligram-to-gram amounts through dietary consumption of fruits and vegetables. Polyphenols are also available as components of dietary supplements for improving general health. Although polyphenols are often advertised as antioxidants to explain health benefits, experimental evidence shows that their beneficial cancer preventing and controlling properties are more likely due to stimulation of pro-oxidant and proapoptotic pathways.

Recent Advances: The understanding of the biological differences between cancer and normal cell, and especially the role that mitochondria play in carcinogenesis, has greatly advanced in recent years. These advances have resulted in a wealth of new information on polyphenol bioactivity in cell culture and animal models of cancer. Polyphenols appear to target oxidative phosphorylation and regulation of the mitochondrial membrane potential (MMP), glycolysis, pro-oxidant pathways, and antioxidant (adaptive) stress responses with greater selectivity in tumorigenic cells.

Critical Issues: The ability of polyphenols to dissipate the MMP ($\Delta\psi_m$) by a protonophore mechanism has been known for more than 50 years. However, researchers focus primarily on the downstream molecular effects of $\Delta\psi_m$ dissipation and mitochondrial uncoupling. We argue that the physicochemical properties of polyphenols are responsible for their anticancer properties by virtue of their protonophoric and pro-oxidant properties rather than their specific effects on downstream molecular targets.

Future Directions: Polyphenol-induced dissipation of $\Delta\psi_m$ is a physicochemical process that cancer cells cannot develop resistance against by gene mutation. Therefore, polyphenols should receive more attention as agents for cotherapy with cancer drugs to gain synergistic activity. *Antioxid. Redox Signal.* 29, 1589–1611.

Keywords: glycolysis, mitochondria, oxidative stress, polyphenols, protonophore, ROS signaling

Introduction

EPIDEMIOLOGICAL STUDIES HAVE revealed an inverse relationship between cancer risk and diets consisting of grains, fruits, and vegetables (12, 39, 218). It is more difficult to establish such a relationship with dietary constituents because food products contain thousands of phytochemicals with diverse actions on normal and abnormal physiology. Many published studies have promoted the popular view that dietary phytochemicals, especially polyphenols, act as antioxidants to repress oxidative stress. This postulate is oversimplified and needs refinement.

Oxidative stress, or the imbalance between the production and removal of reactive oxygen species (ROS), can result from

chronic inflammation and dysfunctional metabolism. NADPH oxidase, xanthine oxidase, and the electron transport chain (ETC) are prominent sources of superoxide, which is subsequently transformed by superoxide dismutase (SOD) into hydrogen peroxide (H_2O_2) and oxygen (52, 63). ROS play a crucial role in innate immunity to fight infections, but can also cause harm to the organism's own biomolecules. To protect cells from damage from excess ROS, the body relies on the antioxidant vitamins C and E to scavenge ROS or to interfere with the propagation of radical chain reactions, which are most likely to initiate in lipid bilayers (195).

Another important cellular antioxidant is glutathione (GSH) (63). Vitamin C and GSH are present in the cytosol at millimolar concentrations while dietary polyphenols or their metabolites

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might reach cellular concentrations in the low micromolar range. Therefore, it is unlikely that dietary polyphenols contribute significantly to any antioxidant activity *in vivo* in the sense that they scavenge ROS (62). This is not to say that dietary polyphenols exert no beneficial health effects *in vivo*. On the contrary, many studies have demonstrated that dietary polyphenols and other phytochemicals exert bioactivity through stimulating or interfering with cell signaling (86, 112). They affect cell signaling by interaction with nuclear receptors and other molecular targets, processes that require far smaller concentrations than the cellular concentrations of the endogenous antioxidants.

Mechanistic studies on the bioactivity of dietary polyphenols suggest that they can act as pro-oxidants *in vivo* (62). Under normal, physiological conditions, a pro-oxidant effect

will deplete cells of GSH, which triggers an adaptive stress response leading to GSH synthesis to protect cells from pre-existing ROS (140). In this scenario, polyphenols exert an indirect antioxidant effect (137, 186), sometimes referred to as “para-hormesis” (45, 46). Under pathophysiological conditions, such as existing in tumor cells where steady-state levels of ROS are abnormally high, dietary polyphenols may actually enhance oxidative stress and drive tumor cells into apoptosis (50).

Regardless of the outcome of polyphenol bioactivity, the chemical moiety responsible for any effect is invariably the hydroxyl group found in all polyphenol classes (Fig. 1). Oxidation of a phenol involves the loss of a proton and an electron to generate a phenoxy radical that then reacts with

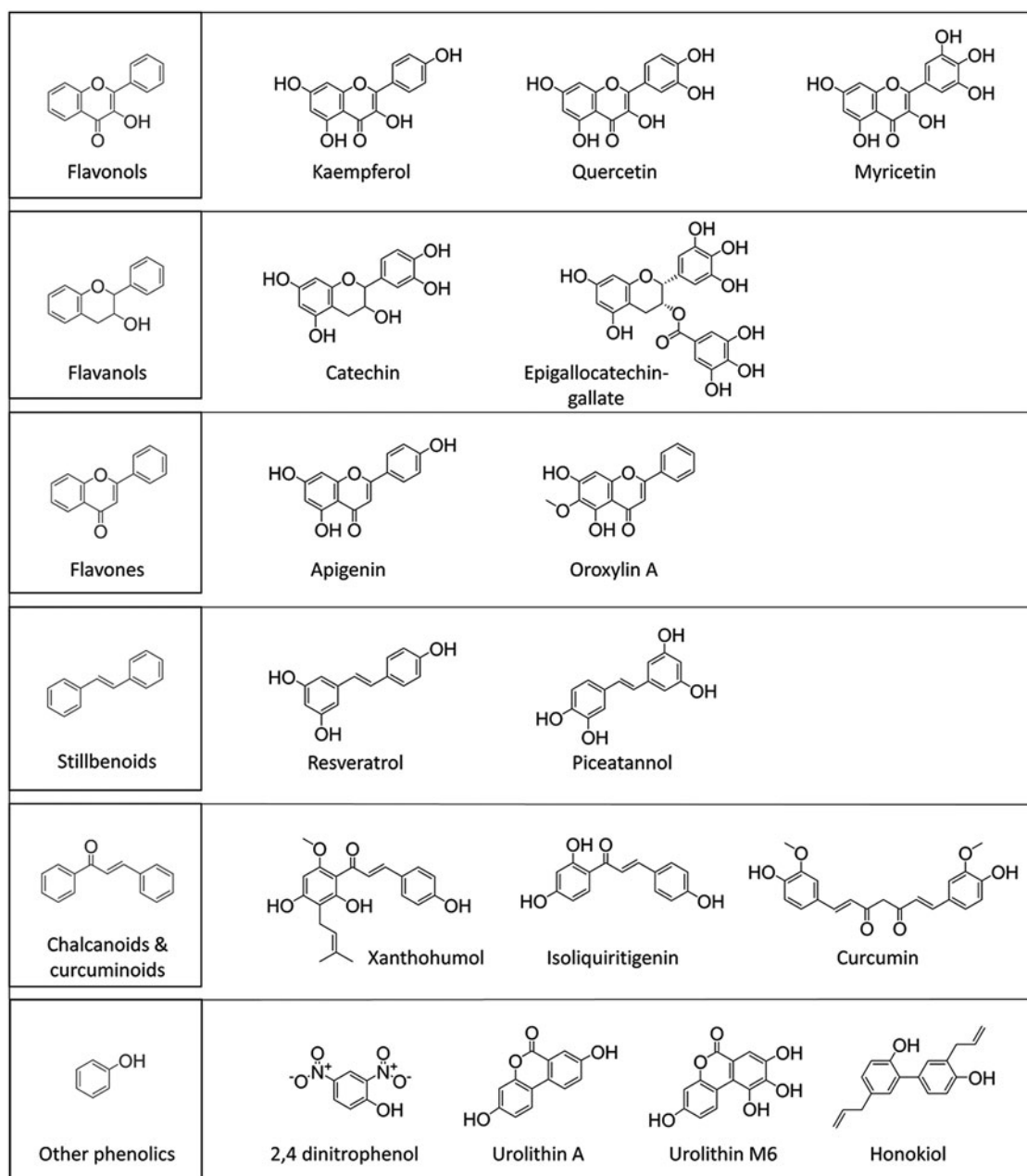


FIG. 1. Chemical structures of polyphenols discussed in this review.

ascorbate or GSH, or participates in radical propagation reactions (44). It can dissociate into a proton and a phenolate anion, which can dissipate the mitochondrial membrane potential (MMP) and cause uncoupling of the ETC from adenosine triphosphate (ATP) production (185).

When polyphenol molecules contain a catechol functionality, they are subject to oxidative conversion into *ortho*-quinones that can interfere with cellular processes through covalent interactions with various biomolecules (15, 77). Several comprehensive reviews provide examples of how *ortho*-quinones react with sulfhydryl groups in the Kelch-like ECH-associated protein 1 (Keap1) protein to induce the Keap1/antioxidant–nuclear factor (erythroid-derived 2)-like 2 (Nrf2) signaling pathway (5, 34, 130, 181). Other reviews (89, 104, 114, 141) describe how polyphenols interfere with the proinflammatory NF κ B (nuclear factor κ -light-chain-enhancer of activated B cells) pathway to exert an anti-inflammatory effect through repressing the release of proinflammatory cytokines from macrophages and other cell types (104, 114, 141).

Polyphenols act as cancer chemopreventive and controlling agents on mitochondrial targets at all stages of carcinogenesis (Fig. 2). This review focuses on the differential effects of polyphenols on mitochondria of normal *versus* cancer cells. Only a limited number of clinical trials have been so far conducted that used polyphenols extracts and/or purified compounds in cancer prevention strategies or as adjunctive in conventional cancer treatment protocols. Those studies have been reviewed recently elsewhere (75, 76).

Polyphenol Metabolism and Bioavailability

Polyphenols have been extensively studied for their health benefits related to cardiovascular diseases (14, 21) and cancer (123, 133, 160). Clinical trials testing efficacy of potential anticancer or chemopreventive phytochemicals have shown variable results [reviewed recently in Hosseini and Ghorbani (76), Kotecha *et al.* (97), Maru *et al.* (123), and Núñez-Sánchez *et al.* (146)]. While *in vitro* studies are crucial to decipher the mode of action of polyphenols at the cellular level, it is often difficult to translate the observations to *in vivo* bioactivity, because various factors influence the bioavailability of polyphenols. They include (i) water solubility (usually $<50 \mu\text{g/mL}$), (ii) route of administration, posology, and dosage formulation, and (iii) metabolic biotransformation [for a review, see Bohn (13)].

Polyphenol metabolism *in vivo* has long been considered as a major limitation for evaluation of clinical studies. Liver and gut metabolism generates a complex mixture of metabolites (methylated, hydroxylated, glucuronidated, sulfated, and hydrogenated forms), all of which contribute to the overall bioactivity of polyphenols. When comparing the physicochemical properties (lipophilicity, pK_a) and reduction potentials of polyphenols (Table 1) and their gut microbial metabolites (Table 2), it appears that metabolites of polyphenols are likely to be bioactive.

A large number of studies have often observed antioxidant and antiproliferative activities of products of polyphenol metabolism (135), such as hydroxyphenyl- γ -valerolactones (58), protocatechuic acid (115, 224), and urolithins (165).

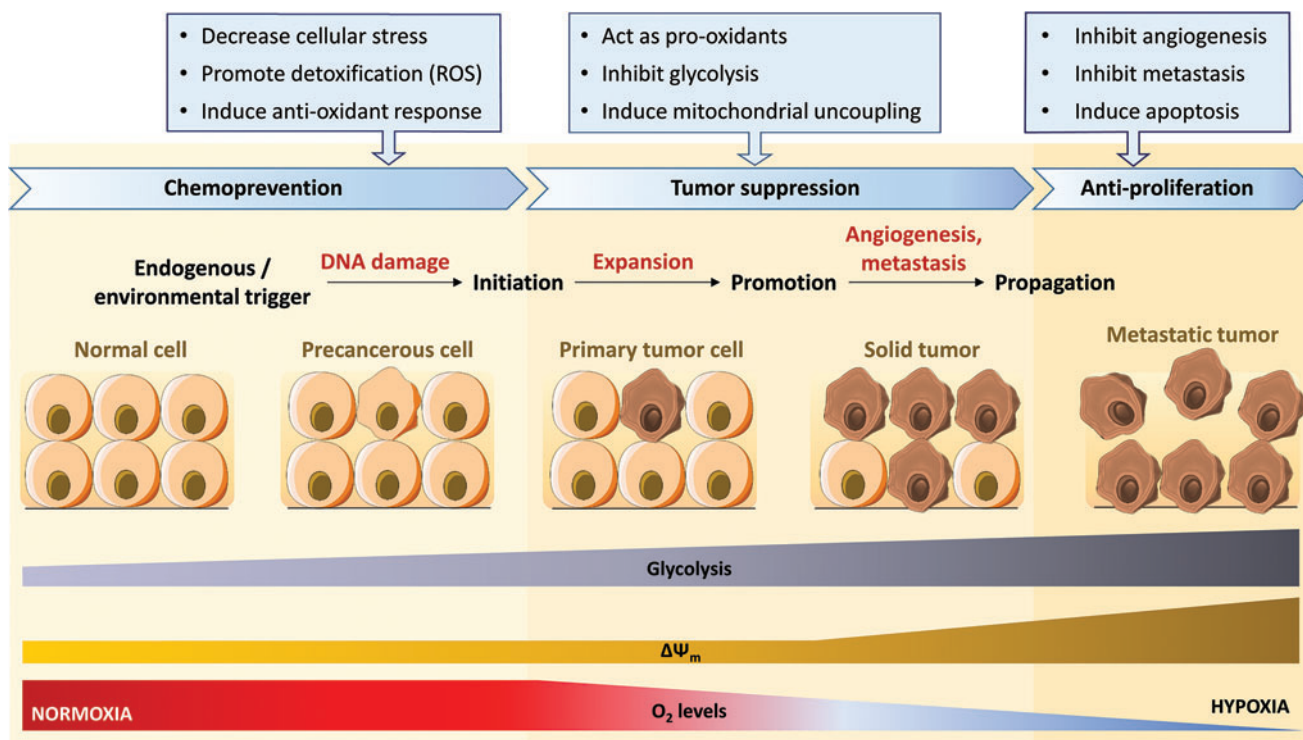


FIG. 2. The multistep process of carcinogenesis. Cancer development, effect of malignant transition on glycolysis rate, mitochondrial membrane potential ($\Delta\Psi_m$) and dioxygen (O_2) levels, and related actions of polyphenols at the different stages of carcinogenesis [adapted from Moga *et al.* (133) and Surh (190)]. ROS, reactive oxygen species. To see this illustration in color, the reader is referred to the web version of this article at www.liebertpub.com/ars

TABLE 1. PHYSICOCHEMICAL PROPERTIES OF POLYPHENOLS DISCUSSED IN THIS REVIEW

<i>Polyphenol</i>	pK_{a1} (most acidic)	pK_{a2}, pK_{a3}		$LogP$ ($logK_{OW}$)	<i>One-electron reduction potential (mV)</i>
2,4 Dinitrophenol	4.04 ± 0.22^a			1.715 ± 0.218^a	—
Apigenin	6.53 ± 0.40^a			2.127 ± 0.452^a	$>500^b$ (220)
Catechin	8.68 ± 0.23 (72)	9.70 ± 0.24 (72)	11.55 ± 0.20 (72)	0.610 ± 0.454^a	570 (83)
Curcumin	8.38 ± 0.04 (10)	9.88 ± 0.02 (10)	10.51 ± 0.01 (10)	3.071 ± 0.444^a	770 (81)
Epigallocatechin-gallate	7.69 (99) to 7.75 (80)	8.80 (80) to 10.70 (99)		0.639 ± 0.702^a	430 (80)
Honokiol	9.89 ± 0.48^a			4.560 ± 0.352^a	—
Isoliquiritigenin	7.50 ± 0.35^a			2.750 ± 0.340^a	—
Kaempferol	6.96 ± 0.09 (72)	8.78 ± 0.11 (72)	10.60 ± 0.12 (72)	2.685 ± 0.812^a	950 (83)
Myricetin	6.30 ± 0.40^a			1.206 ± 1.286^a	360 (9)
Piceatannol	9.17 ± 0.10^a			2.682 ± 0.359^a	—
Quercetin	7.10 ± 0.12 (72)	9.09 ± 0.11 (72)	11.02 ± 0.36 (72)	1.989 ± 1.075^a	330 (82)
Resveratrol	9.22 (27)	10.55 (27) 11.16 (27)		3.024 ± 0.267^a	653 (93)
Urolithin A	9.07 ± 0.20^a			2.311 ± 0.793^a	332^b (85)
Urolithin M6	7.34 ± 0.20^a			0.832 ± 1.274^a	—
Xanthohumol	7.59 ± 0.45^a			4.821 ± 0.416^a	—

^aData obtained from SciFinder (calculated using Advanced Chemistry Development (ACD/Labs) Software V11.02 (© 1994–2017 ACD/Labs).

^bValues given for anode peak voltage (E_{pa}).

Furthermore, conjugated metabolites accumulate in tissues, such as the liver (129), and form a metabolite pool for release of active metabolites. We observed that the magnitude of a biological effect correlates well with circulating levels of polyphenol glucuronides but poorly with circulating trace levels of free polyphenols (109), suggesting that plasma levels of glucuronides reflect tissue concentrations of bioactive forms better than circulating levels of free polyphenols.

Glucuronides are less lipophilic (lower partition coefficient octanol/water; $logP$) and need to be deconjugated by β -glucuronidase to accumulate in cells. β -Glucuronidase is present in the intestine (secreted by bacteria) and in other tissues confined in lysosomes and microsomes of healthy cells, but is extensively secreted by inflammatory cells surrounding tumor areas (16). Numerous anticancer prodrugs (glucuronide-conjugated drugs) have been designed

with the aim to be delivered specifically to the site of the tumor (197).

Polyphenol glucuronides might act as natural prodrugs, being a reservoir of the bioactive aglycone (147). Many studies conducted by others have shown that phase II metabolites can contribute to the pharmacological efficacy of polyphenols (135, 215). Patel *et al.* demonstrated that sulfate conjugates of resveratrol contribute to the regeneration of the intracellular pool of resveratrol and to the beneficial effects against colorectal cancer (155, 156). Moreover, conjugated metabolites of resveratrol, especially resveratrol-3-*O*-sulfate, showed greater potential to inhibit proliferation of human colon carcinoma cells (SW480 and SW62220) compared to the parent polyphenol (1). The advantage here is that phase II metabolites excreted into the bile or generated in the intestine are readily available to colon cells.

TABLE 2. PHYSICOCHEMICAL PROPERTIES OF METABOLITES OF POLYPHENOLS BIOTRANSFORMED BY THE GUT MICROBIOME

<i>Metabolite</i>	<i>Example of precursor</i>	pK_{a1} (most acidic)	$LogP$ ($logK_{OW}$)	<i>One-electron reduction potential (mV)</i>
5-(3',4'-Dihydroxyphenyl)- γ -valerolactone	Epicatechin/Epigallocatechin	9.73 ± 0.10^a	-0.002 ± 0.368^a	—
Caffeic acid	<i>P</i> -coumaric acid	4.58 ± 0.10^a	0.663 ± 0.286^a	534 (43)
Daidzein	Formononetin	7.01 ± 0.20^a	2.632 ± 1.134^a	500 (220)
Dihydrocurcumin	Curcumin	8.35 ± 0.60^a	3.244 ± 0.397^a	—
Gallic acid	Malvidin	4.33 ± 0.10^a	0.531 ± 0.325^a	550 (93)
Genistein	Biochanin A	6.51 ± 0.20^a	3.114 ± 1.137^a	790^b (42)
Hippuric acid	Caffeic acid	3.71 ± 0.10^a	0.757 ± 0.533^a	—
<i>O</i> -desmethylangolensin	Daidzein	7.63 ± 0.35^a	2.587 ± 0.362^a	—
Protocatechuic acid	Catechin	4.45 ± 0.10^a	1.010 ± 0.237^a	298 (219)
(S)-equol	Daidzein	9.94 ± 0.40^a	2.773 ± 0.319^a	—
Urolithin A	Ellagitannins	9.07 ± 0.20^a	2.311 ± 0.793^a	332^b (85)
Urolithin M6	Ellagitannins	7.34 ± 0.20^a	0.832 ± 1.274^a	—

Some compounds (such as caffeic acid, genistein, or daidzein) are also polyphenols in their own right.

^aData obtained from SciFinder (calculated using Advanced Chemistry Development (ACD/Labs) Software V11.02 (© 1994–2017 ACD/Labs).

^bValues given for anode peak voltage (E_{pa}).

In conclusion, the emerging knowledge of bioactive polyphenol metabolites argues in favor of including polyphenol metabolites in estimating overall bioavailability, in accordance with FDA recommendations (200). Pharmacokinetic studies conducted in recent years indicate that the combined bioavailability of polyphenols and their metabolites exceeds 30% (106a, 107, 151, 187). Nevertheless, phase II metabolism is a mechanism of detoxification aimed to remove phenolics from the body. A considerable fraction of the ingested polyphenols are excreted after conjugation. Oral consumption of polyphenols exerts beneficial effects against primary tumors in the gastrointestinal tract (156). Intravenous (IV) administration yields higher plasma levels at equal doses (107). Recently, IV administration has been proposed as the preferred route of administration of bioactive polyphenols for various oncotherapeutic applications (36).

The relatively high concentrations often used in cell culture studies raise the question whether *in vitro* findings bear relevance to the *in vivo* situation. Researchers usually perform cell assays after a single exposure of cells to treat compounds. A small fraction of the amount administered to cultured cells may actually be taken up by the cells due to polyphenol–protein binding in the growth medium surrounding the cells. Polyphenol–protein binding occurs in blood of animals and humans as well, but continuous *in vivo* dosing of a polyphenol or polyphenol mixtures may drive cellular uptake resulting in polyphenol accumulation in organs (129). These considerations suggest that comparisons between cell culture medium concentrations and plasma concentrations are not very meaningful to account for discrepancies between *in vitro* and *in vivo* bioactivity of polyphenols.

Many studies on the bioactivity of polyphenols involve mitochondria. For example, polyphenols extracted from pinecone of *Pinus koraiensis* inhibited tumor growth in sarcoma-bearing 180 mice (223), which the authors attributed to apoptotic pathways initiated by mitochondria. Although very few studies have verified mitochondrial uptake of polyphenols and their bioactive metabolites, their physicochemical properties (lipophilicity, pK_a ; Tables 1 and 2) actually favor their enrichment in mitochondria (142). Human T lymphocyte (Jurkat) cells accumulate quercetin inside the mitochondria, which led the authors to hypothesize that mitochondria represent a reservoir of biologically active quercetin (41). Taken together, the available evidence indicates that polyphenols have greater bioavailability than previously thought and they have the ability to reach potential sites of action in mitochondria. We now discuss the molecular mechanisms by which polyphenols affect mitochondrial function in cancer models (summarized in Table 3). Mechanistic information on the bioactivity of polyphenols primarily originated from *in vitro* studies.

Polyphenols as Mild Mitochondrial Uncouplers

Beneficial effects of mitochondrial uncoupling in obesity

A mild mitochondrial uncoupling activity of dietary polyphenols represents one mechanism pertaining to their beneficial effects on dysfunctional glucose and lipid metabolism in type 2 diabetes, nonalcoholic fatty liver disease, and obesity. Uncoupling of the ETC from ATP production is achieved by dissipation of the electrochemical (proton) gradient across the inner mitochondrial membrane (180), referred to as the MMP, $\Delta\psi_m$. When polyphenols move by

diffusion in cells, they encounter different pH values. In the cytosol (pH 7.4), polyphenols exist as a mixture of phenolate anions and neutral phenols. The proportion of phenol anions and neutrals is dictated by the pK_a of each phenolic group (Tables 1 and 2). Their propensity to cross the cell membrane and the outer and inner mitochondrial membranes depends on their lipophilicity.

Because many polyphenols have favorable pK_a values (pK_{a1} ranging between 6 and 9, *i.e.*, close to the physiological pH of the cytosol and mitochondrial compartments), and distribution coefficients (Table 1) (71, 180), they have the ability to reach the mitochondrial matrix and release a proton in the relatively basic environment (pH 7.8) of the matrix (17). After dissociation, phenolate anions migrate back down the electrochemical gradient to the relatively acidic environment of the intermembrane space (pH 6.8). Back-and-forth diffusion of the phenolate–phenol species across the inner mitochondrial membrane results in transport of protons from the inner membrane space to the mitochondrial matrix (Fig. 3), leading to dissipation of $\Delta\psi_m$ (180, 185).

To maintain $\Delta\psi_m$, cells respond by increasing the activity of the ETC, which requires biofuel. Dissipation of $\Delta\psi_m$ will activate the nutrient sensor AMP-activated protein kinase (AMPK) to stimulate β -oxidation of fatty acids, glycolysis, and the tricarboxylic acid (TCA) cycle, which deliver the necessary reducing equivalents (FADH₂ and NADH) to drive oxidative phosphorylation (150). The ultimate result is that mitochondria consume more fatty acid and glucose molecules to generate the same number of ATP molecules.

The capacity of polyphenols as proton carriers across the inner mitochondrial membrane is referred to as the protonophoric uncoupling or protonophore effect of polyphenols (17, 154, 162), to distinguish this mitochondrial uncoupling effect from effects of polyphenols on adenosine nucleotide translocases (ANTs) (Fig. 4) and uncoupling proteins (UCPs) (Fig. 5), both of which have the ability to leak protons across the inner mitochondrial membrane (18). The protonophore effect of polyphenols and the downstream signaling pathways (*e.g.*, AMPK activation) often explains their improvement of obesity-related dysfunctional glucose and lipid metabolism. In cancer cells, which constantly fight oxidative stress under hypoxic conditions by keeping $\Delta\psi_m$ high, severe dissipation of the $\Delta\psi_m$ will drive the cells into apoptosis (66).

2,4-Dinitrophenol (DNP) has attracted much attention as a potent mitochondrial uncoupler. It was a popular agent for inducing weight loss in the early 1900s, but in the late 1930s taken off the market in response to reports of severe morbidity and death (24). Its mechanism of action appears to be protonophoric in nature, but its acidic and hydrophilic properties ($pK_a = 4.09$ and $\text{Log}P = 1.7$, Table 1) suggest a different mechanism. It is more likely that DNP forms an ion-pair with an endogenous amphiprotic molecule that enhances DNP's lipophilicity and increases the complex's pK_a . McLaughlin has suggested that DNP binds to the lipid bilayer and forms a homodimeric complex consisting of one undissociated and one deprotonated molecule (DNP₂[−]) (125). DNP has been considered for cancer therapy, but deemed too toxic due to its narrow therapeutic window (202). While it is possible to attenuate DNP's potency as a protonophore by improving its pharmacokinetic properties using a slow-release formulation (158), recent interest has shifted to the development of mild mitochondrial uncouplers with better physicochemical and therapeutic properties (202).

TABLE 3. CANCER-RELATED BIOACTIVITIES AT THE MITOCHONDRIA LEVEL FOR POLYPHENOLS DISCUSSED IN THIS REVIEW

<i>Polyphenol</i>	<i>Type of cells</i>	<i>Effect described</i>	<i>References</i>
2,4 Dinitrophenol	Calu-6 human pulmonary adenocarcinoma	Proapoptotic; depolarizes the membrane, increases ROS, decreases GSH	Han <i>et al.</i> (64)
Apigenin Catechin	DU145 human prostate cells B16F-10 melanoma cells	Proapoptotic; inhibits ANT2 Anti-inflammatory, antiangiogenic; reduces mRNA levels of VEGF, reduces cytokines and nitric oxide	Oishi <i>et al.</i> (148) Guruvayoo-rappan and Kuttan (61)
Curcumin	<i>Escherichia coli</i> M15 bacteria SGC-7901 human adenocarcinoma cells T-REx-293 cells	Inactivates PTP complex, induces VDAC closure Proapoptotic; depolarizes the membrane Proapoptotic; induces VDAC oligomerization	Tewari <i>et al.</i> (193) Eggler <i>et al.</i> (33) Keinan <i>et al.</i> (91)
Epigallocatechin-gallate	HCT116 and HT29 human colorectal cancer cells MIA Paca-2 human pancreatic cells and human papillomavirus-18-(+) HeLa cells Human malignant mesothelioma cells	Proapoptotic; downregulates expression of HK-II Proapoptotic; depolarizes the membrane Proapoptotic; impairs ETC	Wang <i>et al.</i> (205) Qanungo <i>et al.</i> (161); Kuo <i>et al.</i> (103) Balogun <i>et al.</i> (6)
Honokiol	Neuro-2a neuroblastoma cells	Proapoptotic; depolarizes the membrane	Kensler and Wakabayashi (92)
Kaempferol	HeLa and HepG2 cells	Suppresses PDK2 and PDK4 mRNA expression	Wang <i>et al.</i> (204)
Myricetin	JAR and JEG-3 malignant trophoblast cells	Antiangiogenic, proapoptotic; increases ROS production, induces GSH depletion, depolarizes the membrane	Yang <i>et al.</i> (221)
Oroxylin A	MDA-MB-231 and MCF-7 human breast cancer A549 lung carcinoma cells	Enhances oxidative stress, activates SIRT3, deacetylates CypD, dissociates HK-II Antimetastatic; dissociates HK-II	Wei <i>et al.</i> (211, 212) Wei <i>et al.</i> (211)
Piceatannol Quercetin (and myricetin) Resveratrol	DU145 prostate cancer cells A549 adenocarcinoma human cells MDA-MB-231 and MCF-7 human breast cancer SW620 colon cancer cells	Proapoptotic; increases MMP Proapoptotic; increases ROS production, inhibits TrxR Stimulates SIRT1, inhibits ETC complex I Proapoptotic; downregulates UCP2	Müller and Hengstermann (139) Lu <i>et al.</i> (117) Deus <i>et al.</i> (28) Dinkova-Kostova and Abramov (30)
Urolithin A	C2C12 myotubes	Decreases membrane potential	Waseem and Parvez (208)
Urolithin M6	Raji cells	Antiproliferative; inhibits LDH, inhibits lactate production	Rupiani <i>et al.</i> (165)
Xanthohumol	C2C12 myotubes	Increases ROS, decreases respiration at concentration >25 μ M	Kirkwood <i>et al.</i> (95)

ANT, adenosine nucleotide translocase; CypD, cyclophilin D; ETC, electron transport chain; GSH, glutathione; HK, hexokinase; LDH, lactate dehydrogenase; MMP, mitochondrial membrane potential; mRNA, messenger RNA; PDK, pyruvate dehydrogenase kinase; PTP, permeability transition pore; ROS, reactive oxygen species; SIRT1, sirtuin 1; TrxR, thioredoxin reductase; UCP, uncoupling protein; VDAC, voltage-dependent anion channel; VEGF, vascular endothelial growth factor.

Attene-Ramos *et al.* (2) screened the Tox21 library of 10,000 compounds to identify chemicals and structural features that are associated with changes in the MMP in HepG2 cells. They found that 11% of the compounds decreased MMP after 1 h of treatment without affecting cell viability. By analysis of

structure/activity relationships of MMP-decreasing compounds using a self-organizing map algorithm, the authors found that nitrobenzenes, thiazoles, flavonoids, and phenols emerged as enriched clusters (2). Many dietary polyphenols meet the requirements of protonophores (Table 1), but their

FIG. 3. Polyphenols are protonophores. Polyphenols (represented by phenol for simplicity) dissociate to phenolate anions at cytosolic pH, diffuse through the outer mitochondrial membrane along the electrochemical gradient into the mitochondrial intermembrane space, where a more acidic environment facilitates their re-protonation and diffusion to the matrix. In the matrix, polyphenols deprotonate and the resulting phenolate anions diffuse back to the intermembrane space down the electrochemical gradient. They travel back and forth between the intermembrane space and the matrix, thereby transporting protons from the intermembrane space to the matrix. Dissipation of the mitochondrial membrane potential ensues. To see this illustration in color, the reader is referred to the web version of this article at www.liebertpub.com/ars

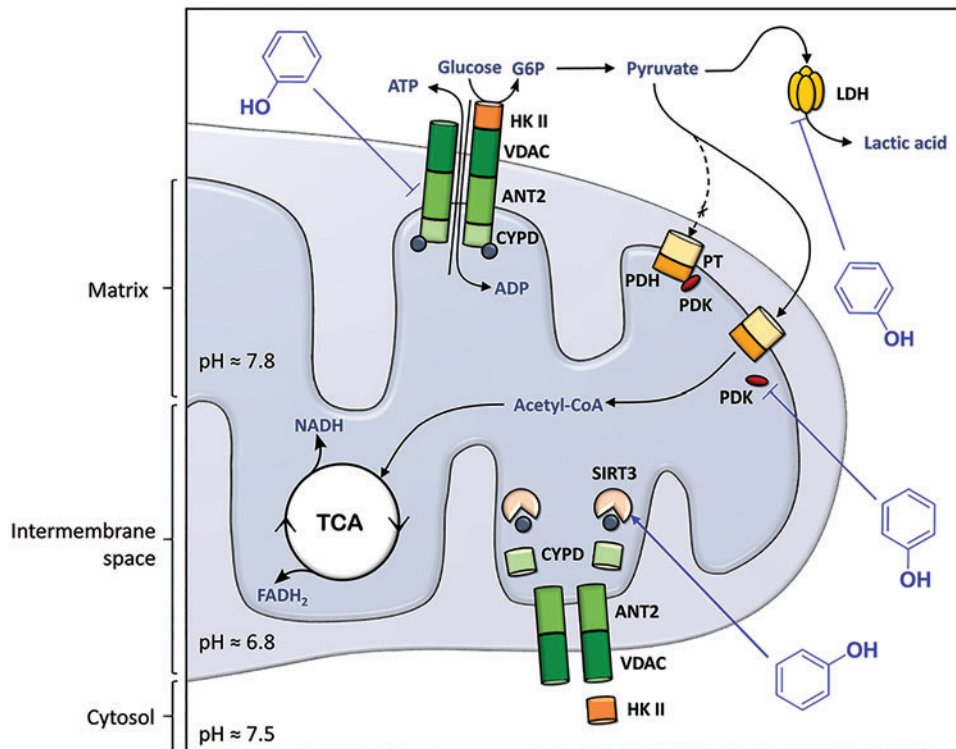
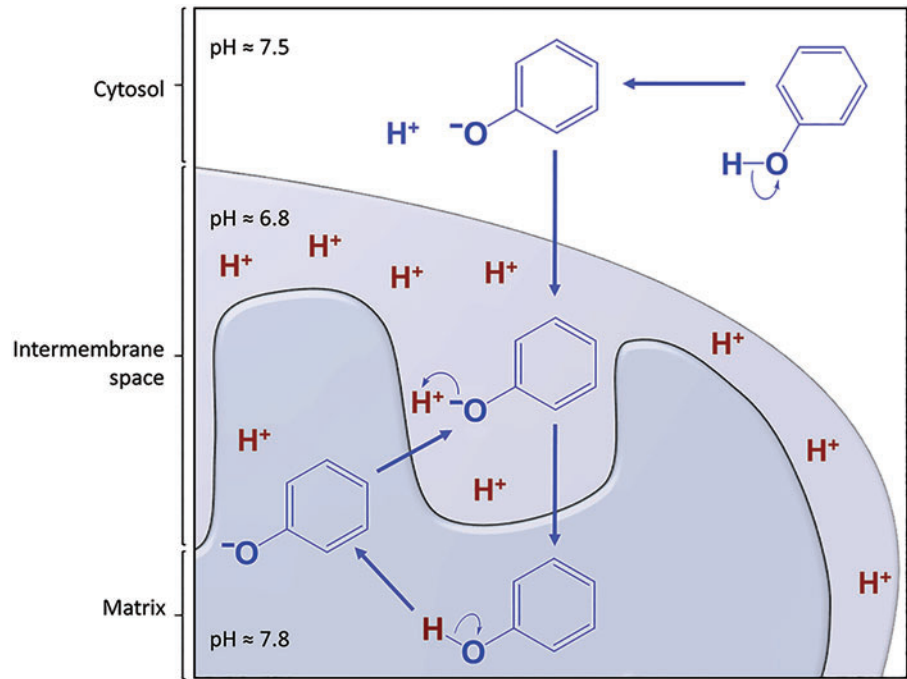


FIG. 4. Polyphenols modulate mitochondrial metabolism. The PTP complex is composed of the VDAC in the mitochondrial outer membrane, the ANT in the mitochondrial inner membrane, and CypD in the matrix. Under physiological conditions, the PTP complex regulates exchange of ATP/ADP and small metabolites between the cytosol and the mitochondrial matrix, and it interacts with HK II in the cytosol that converts glucose to G6P, the first step of glycolysis. In cancer cells, HK II is overexpressed. Polyphenols downregulate ANT (and specifically the isoform 1) and activate the sirtuin 3 protein (SIRT3), responsible for deacetylation of CypD, resulting in detachment of CypD and HK II from the PTP complex. Pyruvate, the end product of glycolysis, is converted into acetyl-CoA by the complex pyruvate transporter (PT)/PDH to enter the TCA cycle. When PDK binds PDH, the conversion of pyruvate to acetyl-CoA is inhibited. In cancer cells, PDK is overexpressed, and pyruvate is converted into lactic acid by LDH in the cytosol. Polyphenols can inhibit PDK to allow conversion of pyruvate into acetyl-CoA by PDH, thereby limiting substrate availability for LDH, which results in inhibition of lactate production. Polyphenols are represented by phenol for simplicity. ADP, adenosine diphosphate; ANT, adenosine nucleotide translocase; ATP, adenosine triphosphate; CypD, cyclophilin D; G6P, glucose-6-phosphate; HK, hexokinase; LDH, lactate dehydrogenase; PDH, pyruvate dehydrogenase; PDK, pyruvate dehydrogenase kinase; PTP, permeability transition pore; TCA, tricarboxylic acid; VDAC, voltage-dependent anion channel. To see this illustration in color, the reader is referred to the web version of this article at www.liebertpub.com/ars

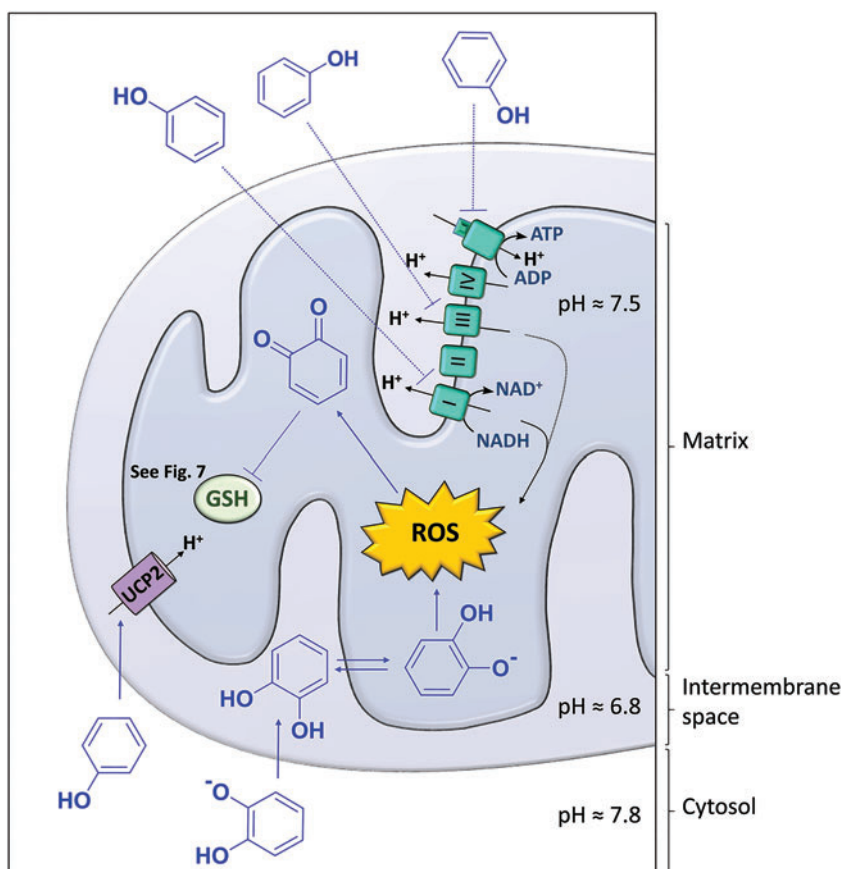


FIG. 5. Polyphenols modulate the mitochondrial redox status. Polyphenols target oxidative phosphorylation by interacting with the electron transport chain complexes (mainly complexes I and III, and ATP synthase) and by decreasing the membrane potential (acting as protonophores) or by activating the UCP2. ROS can convert polyphenols into their corresponding quinones, which can react with the thiol group of GSH. Polyphenols are presented as phenol or as catechol. GSH, glutathione; UCP2, uncoupling protein 2. To see this illustration in color, the reader is referred to the web version of this article at www.liebertpub.com/ars

development is hampered by their low bioavailability and rapid metabolism. On the contrary, many dietary polyphenols produce gut microbial or hepatic metabolites with (partial) retention of bioactivity (184).

The catabolic effects of mitochondrial uncouplers could indirectly suppress growth of metastatic tumor cells by depriving them of nutrients produced and supplied by surrounding cells. This “seed (cancer cell) and soil (tumor microenvironment)” theory offers a two-pronged approach to treat cancers by combination therapy of a mitochondrial uncoupler (e.g., DNP or a suitable polyphenol) to deplete macronutrients in the microenvironment together with a cancer chemotherapeutic to kill fewer and smaller metastatic cell colonies more effectively.

Mitochondrial uncoupling sensitizes cancer cells to apoptosis

Solid tumor cells are heterogeneous in terms of metabolic microenvironment and $\Delta\psi_m$. Heerdt *et al.* (67) found, by characterizing single-cell subclones from the SW620 human colonic carcinoma cell line, that there are subtle, distinct, and stable differences in the cells' $\Delta\psi_m$. While cell growth and proliferation were similar among cells with different $\Delta\psi_m$, the differences in $\Delta\psi_m$ showed a significant effect on levels of vascular endothelial growth factor (VEGF) and matrix metalloproteinase-7 levels. Both proteins play important roles in tumor progression, invasion, and metastasis. Therefore, there appears to be a positive relationship, associative or causative, between $\Delta\psi_m$ and metastatic potential of tumor cells. Polyphenols could conceivably repress the propensity for tumor progression through their protonophoric activity.

Few studies tested the hypothesis whether polyphenols have the ability to suppress the metastatic potential of cancer cells by lowering $\Delta\psi_m$. Guruvayoorappan and Kuttan injected B16F-10 melanoma cells to induce capillary formation in C57BL/6 mice and subsequently monitored the effect of administration of (+)-catechin on angiogenesis (61). They found that the tumor-induced angiogenesis in control animals was accompanied by enhanced serum levels of proinflammatory cytokines and VEGF. Catechin-treated animals showed fewer tumor-directed capillaries, indicating an antiangiogenic effect of catechin. Using an *in vitro* system, they found that catechin reduced messenger RNA (mRNA) levels of VEGF in B16F-10 melanoma cells. The authors contributed the observed antiangiogenic effects of catechin to an anti-inflammatory effect by reducing the production of proinflammatory cytokines and nitric oxide. The authors did not investigate whether catechin exerted its effects through dissipation of $\Delta\psi_m$.

Apigenin inhibits ANT-2

Tumor cells have a higher $\Delta\psi_m$ (≈ -210 mV) compared with normal cells (≈ -140 mV, more negative inside the mitochondrial matrix) (38). The $\Delta\psi_m$ is primarily regulated by the activity of the ETC, ATP synthase (complex V), UCPs, and ANTs. ANTs are embedded in the inner membrane of the mitochondria and are part of the ATP/adenosine diphosphate (ADP) exchange under physiological conditions (Fig. 4). The higher $\Delta\psi_m$ of cancer cells appears to be due, at least in part, to changes in the expression and activity of ANTs.

Tumor cells downregulate the expression of ANT1 (51, 122), which transports ATP from the matrix to the cytosol.

Downregulation of ANT1 expression leads to a lower ATP concentration in the cytosol and to AMPK-mediated stimulation of glycolysis. In contrast, the expression of ANT2 is overexpressed in proliferating cells and upregulated in tumor cells (22, 51, 106). Unlike ANT1, ANT2 appears to export ADP from the mitochondrial matrix to the cytosol in exchange for an equivalent of ATP. The ensuing elevated concentration of ATP in the mitochondrial matrix switches ATP synthase in reverse mode: it hydrolyzes ATP and pumps protons across the inner membrane thereby creating a state of hyperpolarization (high $\Delta\psi_m$) (22). In addition, the decreased cytosolic levels of ATP stimulate glycolysis.

Aside from stimulation of glycolysis, expression and activity of ANTs in tumor cells benefit from keeping $\Delta\psi_m$ high because relative inactivity of the ETC in combination with UCP/ANT-mediated proton leakage could dissipate $\Delta\psi_m$ to a threshold that triggers release of proapoptotic factors and initiation of apoptotic cell death (68, 70).

Thus, polyphenols that exert protonophoric activity or inhibit ANT2 would be expected to have cytotoxic (proapoptotic) potential that is more selective in cancer cells. Apigenin, a dietary flavonoid known to induce apoptosis, was recently identified as an inhibitor of ANT2 in cultured human prostate DU145 cells (148). The authors found that apigenin binds and inhibits ANT2, resulting in Apo 2 ligand (Apo2L)/tumor necrosis factor-related apoptosis-inducing ligand (TRAIL)-induced apoptosis by upregulation of death receptor (DR)-5. Apo2L/TRAIL is a cytokine that belongs to the tumor necrosis factor family, binds to DR5 thereby inducing apoptosis in various malignant tumors but not in normal cells. Silencing ANT2 expression in DU145 cells led to repression of Apo2L/TRAIL-induced apoptosis by apigenin. Genistein also induced apoptosis in this study. However, it did not induce DR5 expression nor did it inhibit ANT2 (148).

Other polyphenols that have been shown to induce apoptosis in cancer cell *in vitro* through increasing TRAIL expression/activity include epigallocatechin gallate (EGCG; gastric cancer cells) (149), human prostate carcinoma LNCaP cells (176), resveratrol (xenograft mouse model of prostate cancer) (47), and the rice bran polyphenol cycloartenyl ferulate (human colorectal adenocarcinoma cells) (96), but none of the studies investigated the role of ANT2 inhibition. One study reported that the polyphenols, chrysin, apigenin, and acacetin, inhibited TRAIL-receptor 1 on activated RAW264.7 macrophages (207). Sensitization of TRAIL-mediated apoptosis is an important approach for cancer therapy because a number of tumors have developed resistance to TRAIL (226).

Polyphenols induce apoptosis and mitochondrial membrane depolarization

Many studies have demonstrated that polyphenols induce apoptosis and dissipate $\Delta\psi_m$ in cultured cancer cells. Curcumin induced apoptosis in human gastric adenocarcinoma SGC-7901 cells, which was associated with dissipation of $\Delta\psi_m$ (33). Honokiol, a neolignan-type biphenol from the bark of the *Magnolia* tree and available as a dietary supplement, induced apoptosis and decreased $\Delta\psi_m$ of neuroblastoma neuro-2a cells at 40 μM exposure (92).

The tea polyphenols, EGCG and theaflavins, caused a concentration- and time-dependent inhibition of proliferation of human papilloma virus-18-positive HeLa cervical cancer

cells, apoptosis, and a decrease of $\Delta\psi_m$ (103). Qanungo *et al.* (161) observed that, in human pancreatic MIA PaCa-2 cells treated with 200 μM of EGCG, mitochondrial membrane depolarization preceded apoptosis.

Kim and coworkers examined the mechanism by which the grape stilbenoid, piceatannol, inhibited proliferation of cultured androgen-insensitive DU145 prostate cancer cells at concentration in the 1–10 μM range. Under these conditions, the cells exhibited increased mitochondrial membrane permeability and apoptosis (139).

These studies have in common that structurally unrelated polyphenols induced apoptosis and cause $\Delta\psi_m$ dissipation in various cultured cancer cells at low micromolar concentrations. We hypothesize that the protonophoric effect of these polyphenols forms a “universal” contributing mechanism for the polyphenol-induced dissipation of $\Delta\psi_m$ and apoptosis. The attractive aspect of the protonophore effect is that it is biophysical in nature and thus not sensitive to development of tolerance resulting from mutation-related changes of specific molecular targets.

Polyphenols as modulators of the ETC-complexes

Because cancer cells prefer glycolysis and circumvent oxidative phosphorylation, the inhibition of ETC complexes seems less promising as a cancer treatment strategy. However, cancer cells exhibit enhanced oxidative stress levels and production of ROS (compared with normal cells) due to their high energy demands (56, 196). Hence, enhanced generation of ROS by inhibiting the ETC complexes may overburden the antioxidant systems and make cancer cells more susceptible to cell death. Current reports are mostly limited to testing of polyphenols as ETC complex modulators in isolated mitochondria.

For instance, resveratrol was reported to inhibit complex I activity (136), while the isoflavone, genistein, inhibited complex III in isolated rat liver mitochondria (167). Flavonoids from a French maritime pine bark decreased activity of complexes I, II, and III and reduced cytochrome C reversibly in rat liver mitochondria (134). Interaction of polyphenols with ETC-complexes seems to stimulate generation of ROS (Fig. 5) (59, 168). Interestingly, resveratrol and its mitochondria tropic derivatives, that is, conjugates with (4-triphenyl-phosphonium)butyl group attached to either 4'- or 3-OH, in conjunction with capping of its free hydroxyl groups inhibited the F_0F_1 -ATPase, accumulated in energized mitochondria as expected and showed cytotoxicity against fast growing but not against slower growing cells (11).

Valenti *et al.* (201) reported that EGCG negatively modulated oxidative phosphorylation by impairing ETC activity (particularly complex I, II, and ATP synthase) in human malignant pleural mesothelioma cells and induced growth arrest and apoptosis. EGCG did not inhibit the growth of normal mesothelial cells.

We postulate that inhibition of ETC constituents by polyphenols will lead to a nonfunctional ETC that is incapable of maintaining $\Delta\psi_m$. Because cancer cell survival depends on high $\Delta\psi_m$, any mechanism that leads to dissipation of $\Delta\psi_m$ will promote activation of apoptotic pathways.

Polyphenols affect UCP2 levels

Several cancer cell types have relatively high UCP2 expression compared with normal cells (Fig. 5) (74). Glycolysis-

utilizing cells benefit from elevated UCP2 expression because the ensuing dissipation of $\Delta\psi_m$ triggers the reverse action of ATP synthase, which increases ATP hydrolysis and leads to a lower cytosolic ATP/ADP ratio (4), favoring glycolysis and production of NADPH by the pentose phosphate pathway (PPP) to keep oxidative stress low. Polyphenols have received little attention to target UCP2 in cancer cells: most studies investigating the effects of polyphenols on UCP2 expression support the hypothesis that polyphenols stimulate UCP2 to reduce ROS and confer protective effects against diabetes and cardiovascular disease.

In rats treated by intraperitoneal administration of resveratrol, Della-Morte *et al.* (26) found that resveratrol significantly decreased UCP2 levels in the hippocampus by 35% compared to sham-treated rats. They determined that the decrease was mediated by sirtuin 1 (SIRT1) based on the observation that treatment with the SIRT1-specific inhibitor, sirtinol, abolished the decrease in UCP2 levels. Deus *et al.* (28) found that resveratrol was toxic to breast cancer cells and that the cytotoxicity was similarly related to the stimulation of SIRT1 (the role of UCP2 was not examined). Furthermore, they observed that resveratrol decreased mitochondrial respiration, which they attributed to complex I inhibition. SW620 colon cancer cells responded to resveratrol treatment (10 μM) by downregulation of UCP2 expression, apoptosis, and cell growth inhibition (30).

By contrast, in rats fed a high-fat diet and treated orally with resveratrol, hepatic expression of UCP2 was significantly increased by 76% compared with rats fed the high-fat diet only (159). While resveratrol shows antiproliferative effects in cancer cells by inducing apoptosis (101, 192), the precise role of UCP2 in these studies remains to be determined. Stimulation of UCP2 activity or expression by polyphenols, if sufficiently potent, may result in antitumor effects by collapsing $\Delta\psi_m$, thereby driving cells into apoptosis.

Polyphenols Reverse Metabolic Reprogramming in Cancer Cells

A significant hurdle in cancer control strategies is that tumors are heterogeneous, harboring different cell communities with different oncogenic genotypes. Therefore, treatment strategies that are designed for targeting a single gene or a distinct kinase cascade have been moderately successful. There is a growing body of work indicating that polyphenols display biological activities as metabolic modulators that have potential in cancer prevention and control.

In general, cancer prevention and chemotherapies aim to exploit differences between cancer cell phenotypes and their normal counterparts. Maybe, in part, stimulated by the renaissance of metabolomics, the fourth omics realm, there is a growing body of research that indicates that malignant cells exhibit a dependence on cytoplasmic glycolysis under aerobic conditions for covering their ATP demands, a phenomenon broadly known as the Warburg effect (54). Although aerobic glycolysis generates fewer ATP molecules than aerobic respiration (2 vs. 38 ATP molecules per molecule of glucose consumed), the lower efficiency of glycolysis is made up for by higher metabolic flux.

The metabolic advantages of the metabolic switch are multiple. Glycolysis does not require oxygen. Glycolysis produces fructose-6-phosphate and glyceraldehyde-3-phosphate that can

be used as substrates in the PPP to generate building blocks for the biosynthesis of macromolecules that are needed to sustain cell division (*e.g.*, ribose-5-phosphate and other C₅ sugars for RNA and DNA synthesis). Glycolysis requires NAD⁺, which is provided by the conversion of pyruvate into lactate. The resultant extracellular acidosis creates a selection medium for growing cells, from which adapting cells with upregulated glycolysis and acid resistance emerge as a pool of cells that promote proliferation and invasion (48).

Polyphenols shift glycolysis to fatty acid β -oxidation

Urolithins are formed by gut microbial metabolism of ellagitannins (105), a group of bioavailable polyphenols that are abundantly present in walnuts and pomegranate (35, 127). The physicochemical characteristics of urolithins would give them potential protonophoric properties (Table 2). Urolithin A indeed acutely lowered $\Delta\psi_m$ in C2C12 myotubes, but the effect was not a result of inhibition of the ETC or mitochondrial uncoupling (166).

Instead, Ryu *et al.* (166) attributed the effect to a decrease in basal respiration. By challenging the urolithin A-treated myotubes with intermediates or precursors of glycolysis/TCA cycle, the researchers were able to reveal a urolithin-induced downregulation of aerobic glycolysis, resulting in indirect inhibition of complex I respiration. By exposing the cells to a medium containing palmitate that stimulates fatty acid β -oxidation (FAO), they demonstrated that the cells can partially compensate for this effect by shifting to FAO-driven respiration. Although the study was aimed at elucidating the effects of urolithin A on mitophagy in *Caenorhabditis elegans* and on muscle function in rodents, this reprogramming of energy metabolism would deprive tumor cells of using aerobic glycolysis for their energy needs.

del Mar Blanquer-Rosselló *et al.* (25) studied the effects of resveratrol in SW620 colon cancer cells. In these cells, resveratrol treatment (10 μM) resulted in decreased UCP2 levels, increased $\Delta\psi_m$, increased H₂O₂ levels, and increased respiration with greater ATP production compared to vehicle-treated cells. These changes resulted in apoptosis and inhibition of cell proliferation, which the authors, substantiated with experimental data, attributed to a resveratrol-induced metabolic switch from glycolysis to FAO.

Polyphenols inhibit lactate dehydrogenase

The switch from aerobic respiration to glycolysis as the primary ATP-producing pathway makes tumor cells highly dependent on glucose. The lactate dehydrogenase (LDH)-mediated conversion of pyruvate into lactate is essential in tumor cells because it generates sufficient NAD⁺ equivalents to sustain glycolysis (38). The reliance on LDH makes it a potential target for starving tumor cells, especially in combination with cancer therapeutics that target other pathways to overcome resistance. Galloflavin and the gut microbial metabolite of ellagitannin, urolithin M6, inhibited human LDH-A with IC₅₀ values of 70 and 77 μM in a biochemical assay (165). They inhibited proliferation of Raji cells (IC₅₀ 33 and 25 μM) and inhibited lactate production in these cells with IC₅₀ values of 62 and 36 μM . While these IC₅₀ are high, these ellagitannin derivatives provide lead compounds for further development and they could reach effective concentrations in the colon to kill colon cancer cells.

Polyphenols modulate pyruvate dehydrogenase

Pyruvate dehydrogenase (PDH) activity is a key determinant for metabolic flux from pyruvate to acetyl-CoA, an important entry metabolite of the TCA cycle. PDH activity is under control of PDH kinase (PDK) (98). PDK, which exists as multiple isoforms, has emerged as potential target in cancer therapy (191). PDK isoforms have been identified as molecular switches that downregulate the PDH complex by reversible phosphorylation in mitochondria (90). Phosphorylating the E1 subunit of PDH by PDK results in inhibition of PDH activity, thereby inhibiting oxidative decarboxylation of pyruvate to acetyl-CoA and thus facilitating cytosolic conversion of pyruvate into lactate (Fig. 4).

In cancer cells, PDK has been found activated and augmenting the glycolytic phenotype of cancer cells. Therefore, PDK is a potential therapeutic target because its inhibition would result in enhanced PDH activity, thereby facilitating the conversion of pyruvate into acetyl-CoA and reducing substrate availability for the conversion of pyruvate into lactate. The overall result is an increase in oxidative phosphorylation at the expense of aerobic glycolysis that underlies the Warburg phenotype.

Dichloroacetate (DCA) inhibits PDK by promoting local conformational changes that are communicated to both the nucleotide-binding and lipoyl-binding pockets of PDK1, leading to inactivation of kinase activity (90). This in turn increases the flux of pyruvate into the mitochondria, thereby stimulating oxidative phosphorylation over glycolysis. DCA has shown in preclinical work to reverse glycolysis-related suppression of mitochondrial apoptosis in cancer cells (20, 216) and boosted the effectiveness of hypoxia-specific chemotherapies *in vitro* (65, 164) and in animal models (19). Resveratrol was reported recently to enhance PDH activity, likely *via* activation of AMPK, which in turn led to a shift from glycolysis to oxidative phosphorylation in HTC116 colon cancer cells (171). Importantly, the resveratrol concentration (10 μ M) used in the cell culture study to elicit the reversal of the Warburg effect was in the range that was detected in the serum of resveratrol-treated human subjects (8).

There is limited published work hinting toward a role of polyphenols as inhibitors of PDKs. Wang *et al.* (204) reported that kaempferol, a flavonol found in many plants and plant-derived foods, functions as an estrogen receptor (ER)- β agonist and as an estrogen-related receptor (ERR)- α and ERR- γ inverse agonist in HeLa and HepG2 cells. ERRs play important roles in oxidative phosphorylation, mitochondrial biogenesis, and energy metabolism (49). The authors attributed the kaempferol-induced inhibition of PDK to the existence of multiple isoforms of PDK whose expressions are regulated by ERRs. Noteworthy, ERR- α and Nrf (both isoforms 1 and 2) are targets of peroxisome proliferator-activated receptor γ coactivator 1- α (PGC-1 α) transcriptional coactivators (179). ERR- α is a recognized prognostic marker in breast, colon, and prostate cancers (182).

Polyphenols modulate voltage-dependent anion channel and hexokinase activity

In response to hypoxia, orchestrated, at least in part, by hypoxia-inducible factor-1 (HIF-1), mitochondrial ATP production shifts to glycolysis. Mitochondrial dysfunction has been linked to downregulation of the catalytic F1 subunit of mito-

chondrial ATP synthase in human carcinomas (116). Malignant cells display reduced mitochondrial vitality and compromised competency to orchestrate apoptosis. The mitochondrial phenotype of cancer cells and associated alteration of mitochondrial biology have been reviewed in Gogvadze *et al.* (52).

Highly malignant cancer cell phenotypes have acquired resistance to apoptotic cell death and have been associated with poor clinical prognosis. The intrinsic apoptotic pathway involves permeabilization of the outer mitochondrial membrane (OMM) and subsequent release of cytochrome c (and other proteins) from the intermembrane space of mitochondria. Cytosolic cytochrome c associates with apoptotic protease activating factor-1, Apaf-1, and initiates recruitment and activation of caspases leading to protein degradation and ultimately cell death. The permeabilization of the OMM is critically involved in initiating mitochondrial apoptosis (53).

There is a growing body of work that suggests that adaptation to hypoxia associated with metabolic reprogramming is linked to enhanced resistance of malignant cells to apoptosis. Gogvadze *et al.* discussed the role of Bcl-2 family proteins in OMM permeabilization (52). In malignant cells, imbalance exists between proapoptotic (*e.g.*, Bax and Bak) and anti-apoptotic proteins (*e.g.*, Bcl-2, Bcl-XL, and others). Gogvadze *et al.* argue that binding of Bcl-2 to the OMM-embedded voltage-dependent anion channel (VDAC) favors the open state preventing initiation of the apoptotic cell death program in cancer cells (54).

As an important gatekeeper of mitochondrial metabolic flux, VDAC interacts with ANT, forming a link between the outer and inner mitochondrial membranes (Fig. 4).

The VDAC family of pore-forming proteins consists of three isoforms, VDAC1-3. VDAC1 is overexpressed in many cancer types in line with the increased energy demands of malignant cells. VDAC is located in the OMM. VDAC1 controls the metabolic cross talk between mitochondria and the rest of the cell by the regulating the metabolite flux in the mitochondrion and thus cellular energy production. VDAC1 is also a key player in mitochondria-mediated apoptosis through its involvement in the release of apoptotic proteins and interaction with antiapoptotic proteins (175).

Curcumin, the main polyphenol of turmeric (*Curcuma longa*), has been reported to exert cancer preventive and anticancer activities (100). Human VDAC1, embedded in planar lipid bilayers, showed reduced conductance when treated with curcumin (193). Molecular docking in conjunction with site-directed mutagenesis experiments showed that curcumin is capable of interacting with residues in the N-terminal α -helical region and the channel wall of VDAC1. Based on the results, the authors proposed that curcumin stabilizes the closed state of VDAC1. There is an ongoing dispute as to which state of the channel, closed or open, is proapoptotic. However, stabilization of the closed state will presumably disrupt normal metabolite flux and hence likely induce apoptosis. VDAC1 closure by curcumin may therefore contribute to the chemopreventive and anticancer activity often attributed to this polyphenolic compound (193).

VDAC1 displays binding to multiple proteins, including proteins that regulate apoptosis, such as Bcl2-family members and hexokinase (HK) II, which catalyzes the ATP-dependent phosphorylation of glucose to glucose-6-phosphate (G6P), the rate-limiting step in glycolysis. HK-II binding to VDAC1 protects cancer cells against apoptosis and promotes cell survival

(175). Curcumin ($40\ \mu\text{M}$) also induces VDAC oligomerization and apoptosis in T-Rex293 cells (91). The effectiveness of curcumin to act as a proapoptotic agent is in agreement with earlier reports (32).

In many types of tumors, HK-II has been found to be up-regulated and to promote enhanced aerobic glycolysis. Mitochondria-bound HK-II has been touted as both facilitator and gatekeeper of malignancy (124). Binding of HK-II to VDAC has been proposed to be critical for preventing induction of apoptosis in malignant cells. Gogvadez *et al.* suggested that HK-II antagonizes the binding of proapoptotic proteins and hence prevents induction of apoptotic pathways (54).

Oroxylin A, an *O*-methylated flavone, occurs in the root of the Chinese traditional medicinal plant *Scutellaria baicalensis*, and has been identified as a HK-II inhibitor. Oroxylin A treatment resulted in detachment of HK-II from VDAC and in redistribution of HK-II to the cytosol in the human breast cancer cell lines MDA-MB-231 and MCF-7 (211). Oroxylin A increased SIRT3 activity, which in turn caused deacetylation of cyclophilin D (CypD) and detachment from ANT2, which caused HK-II dissociation from the mitochondria and HK-II inhibition (211, 212). The same authors also reported that oroxylin A caused dissociation of HK-II from the mitochondria in human A549 lung carcinoma cells (210). These findings indicate that oroxylin A acts to enhance HIF-1 α destabilization *via* SIRT3 activation, CypD deacetylation, and dissociation of HK-II from the mitochondria, resulting in an overall regain of apoptotic capacity in cancer cells (119).

Curcumin has been reported to downregulate the expression and activity HK-II thereby inhibiting aerobic glycolysis and inducing mitochondria-mediated apoptosis in human colorectal cell lines HCT116 and HT29 (205).

Antioxidant and Pro-oxidant Activities of Polyphenols

In vitro antioxidant and pro-oxidant chemistry of phenolic compounds

Radical scavenging, transition metal chelation, and activation of the antioxidant stress response represent diverse antioxidant mechanisms of polyphenols (Fig. 6). The radical scavenging properties and structure/function relationships have been reviewed previously (69, 163). Experimental *in vitro* studies often measure their antioxidant capacity. Gas-phase calculations indicate that radical scavenging by polyphenols can occur *via* two 1-electron mechanisms: (i) H-atom transfer from a polyphenol to a free radical R^\bullet and (ii) the free radical abstracts an electron from the polyphenol (110) [Fig. 6, schema (1)]. The reduction potential of a compound is another way to assess its ability to scavenge free radicals. It is calculated by voltammetry techniques and dependent on different parameters, including concentration.

Published reduction potentials of polyphenols are given in Tables 1 and 2. The lower the reduction potential, the better hydrogen donor the polyphenol (antioxidant) is. The reduction potentials of GSH and vitamin C are $-170\ \text{mV}$ (128) and $282\ \text{mV}$ (214), respectively. Polyphenols, with reduction potential values ranging between 300 and 900 mV, have the possibility to act as radical scavengers, but they are less potent compared to GSH and vitamin C. Moreover, *in vivo* direct radical scavenging by polyphenols is believed to be a minor antioxidant mechanism in the sense that the rate of this reaction is limited by the concentration of the antioxidant

(45). Polyphenols could act as antioxidants in the intestine where polyphenol concentrations can reach millimolar concentrations after consumption of fruits and vegetables (172).

The antioxidant properties of polyphenols have also been associated with their capability of chelating metal ions, which would prevent Fenton-type reactions that produce highly reactive hydroxyl radicals. For instance, Leopoldini *et al.* (111) studied the iron(II) chelation mechanism of quercetin using density functional theory computation. These calculations indicate that neutral and deprotonated quercetin forms complexes with Fe^{2+} with 1:1 and 1:2 metal/ligand stoichiometry. Complex formation engages the oxygen atoms at the 3 and 4, and at the 5 and 4 carbons. The architecture of the chelator complex was also confirmed by electrospray ionization mass spectrometry studies (170). The chelating properties of polyphenols depend on the presence of a suitable chelation site, such as found in flavonoids (40).

In vitro pro-oxidant activity of polyphenolics is dependent on the total number of hydroxyl groups and their substitution patterns. Polyphenols with *ortho* or *para* substituted di- and trihydroxy groups display significantly increased production of hydroxyl radicals and readily undergo auto-oxidation, thereby forming conjugated dicarbonyl compounds (quinones) as reactive intermediates [Fig. 6, schema (2)].

The higher rate of production of ROS in cancer cells compared with normal cells puts cancer cells at a disadvantage to resist apoptosis when exposed to pro-oxidant polyphenols. Several studies have shown that polyphenols of the flavanol type (catechins and EGCG) and proanthocyanidin type, which are abundantly present in green tea and berries, induce apoptosis in colon cancer cells presumably by producing H_2O_2 in the gut lumen. Although oligomeric catechins are poorly taken up by cells, the H_2O_2 they produce extracellularly can diffuse into colon epithelial cells. This mechanism does not require uptake of polyphenols by cancer cells that face the gut lumen.

Chung *et al.* studied the effects of hop proanthocyanidins on human colon cancer HT-29 cells and found that the mixture of polyphenols induced apoptotic death in a dose-dependent manner (23). They attributed apoptosis to proanthocyanidin-mediated H_2O_2 production in the medium but noted that the increased ROS production could not fully account for the observed protein carbonylation. A satisfactory explanation for the observation that proanthocyanidins promote protein carbonylation can be inferred from the work by Gosse *et al.* (57), who studied the effects of apple proanthocyanidins in the human adenocarcinoma SW620 cell line. They found that apple proanthocyanidins produced ROS, triggered apoptosis, and promoted polyamine catabolism, which produces H_2O_2 as well as the reactive aldehyde, acrolein (183). Weh *et al.* (209) reported that cranberry proanthocyanidins induced autophagic cell death in apoptosis-resistant esophageal adenocarcinoma (EAC) cells. They found that the cranberry proanthocyanidins increased ROS levels in EAC cells and also in a medium in the absence of cells, indicating that proanthocyanidins can produce ROS extracellularly.

Valenti *et al.* (201) addressed the EGCG-induced extracellular production of H_2O_2 by adding SOD and catalase, both cell impermeable, to the culture medium. Under these conditions, human malignant mesothelioma cells showed characteristics of cell growth arrest and apoptosis, which the authors explained by EGCG-induced impairment of ETC

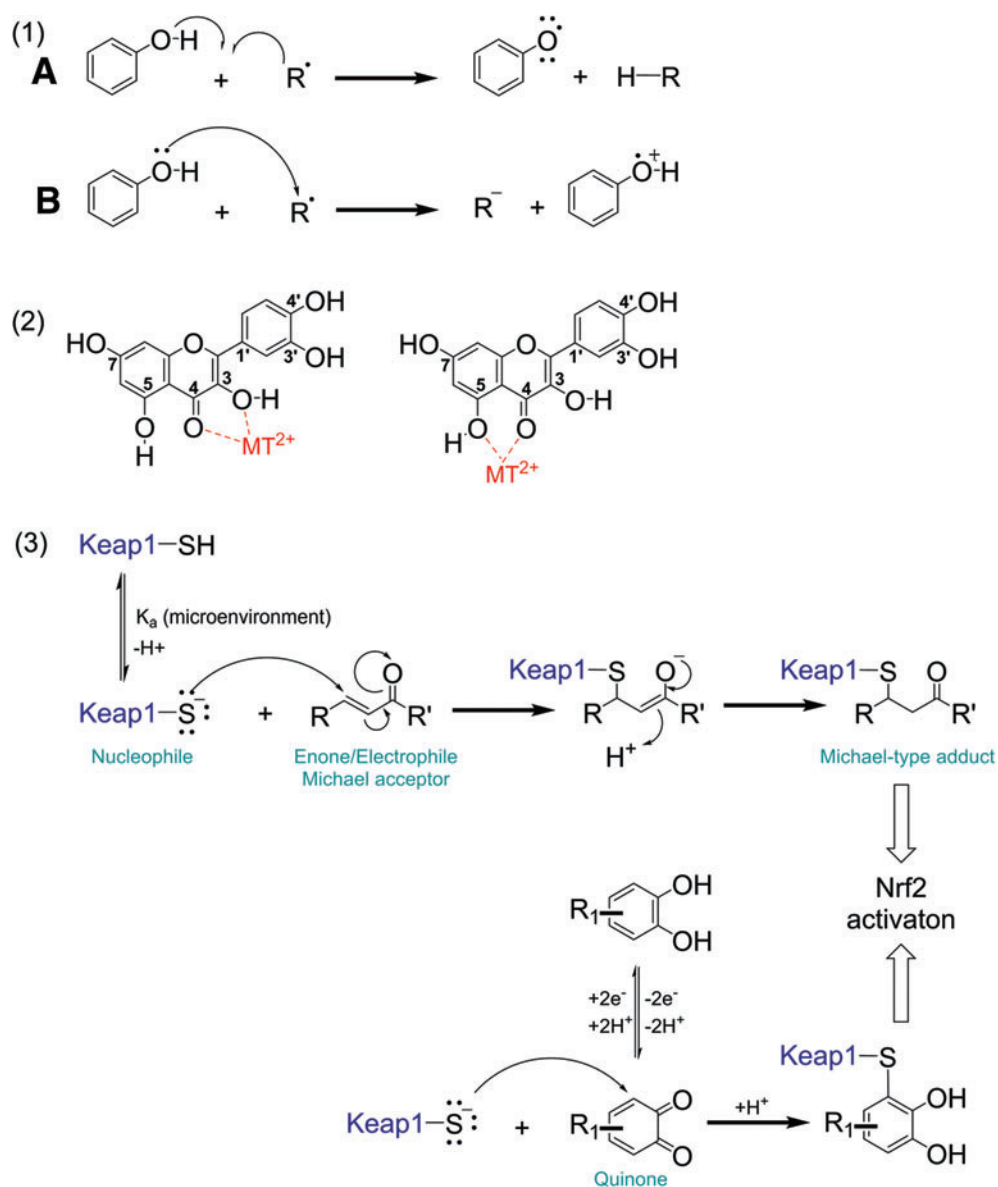


FIG. 6. Antioxidant mechanisms of polyphenols. (1) Hydrogen donation to a radical species R^\bullet to form RH (A), electron donation to a radical species R^\bullet (B), (2) transition metal chelation, (3) induction of the antioxidant Keap1 - Nrf2 pathway by Michael-type attack of Keap1 sulfhydryl groups at electrophilic moieties in polyphenol-derived quinones and, as shown, chalcone-type flavonoids. Keap1, Kelch-like ECH-associated protein 1; Nrf2, nuclear factor (erythroid-derived 2)-like 2. To see this illustration in color, the reader is referred to the web version of this article at www.liebertpub.com/ars

complexes, resulting in reduction of oxidative phosphorylation without a compensatory increase in glycolytic capacity. This study demonstrates that metabolic reprogramming may not be related to a pro-oxidant effect of EGCG.

Our laboratory investigated the role of intracellularly produced ROS by the hop prenylated polyphenol, xanthohumol, on oxidative phosphorylation. At low concentrations ($<5 \mu\text{M}$), xanthohumol increased respiration in C2C12 myotubes, whereas treatment at higher concentrations ($>25 \mu\text{M}$) resulted in a decrease. By pretreating the cells with the cell-permeable ROS scavengers, SOD/peroxynitrite dismutase mimetic MnTMPyP and the SOD/catalase mimetic EUK 134, we were able to demonstrate that ROS play a role in the xanthohumol-induced mitochondrial uncoupling (95).

Another prenylated flavonoid, artocarpin, appears to exert cytotoxic effects on nonsmall-cell lung carcinoma (NSCLC; A549), breast adenocarcinoma (MCF-7), ovarian carcinoma (1A9), and glioblastoma (U87-MG) cells (206). Tsai *et al.* found that the artocarpin-induced cytotoxicity (1–10 μM) in the A549 cell line was due to artocarpin's ability to produce

ROS by activating NADPH oxidase *via* the Nox2/p47^{phox} pathway, thereby triggering apoptosis (199).

Polyphenols act as inhibitors of the thiol redox system

Cancer cells are forced to respond to increase in ROS production by increasing levels of antioxidant proteins to preserve redox homeostasis (Fig. 7). Thioredoxin (Trx), Trx reductase (TrxR), and NADP are the constituents of the Trx system, which are critical for maintenance and control of the redox homeostasis by removing ROS and replenishing antioxidants (Fig. 7A). In parallel to maintaining Trx redox homeostasis, the GSH system is active: the oxidized form of glutaredoxin is reduced by oxidation of GSH, which in turn is regenerated by GSH reductase (GR) (73).

In general, malignant cells with high levels of antioxidant enzymes display resistance to chemotherapies. For example, MCF-7 cells that overexpressed TrxR-1 were resistant to the proapoptotic synthetic flavonoid LW-214, a chrysin derivative (152). Gorklach *et al.* extensively reviewed polyphenols

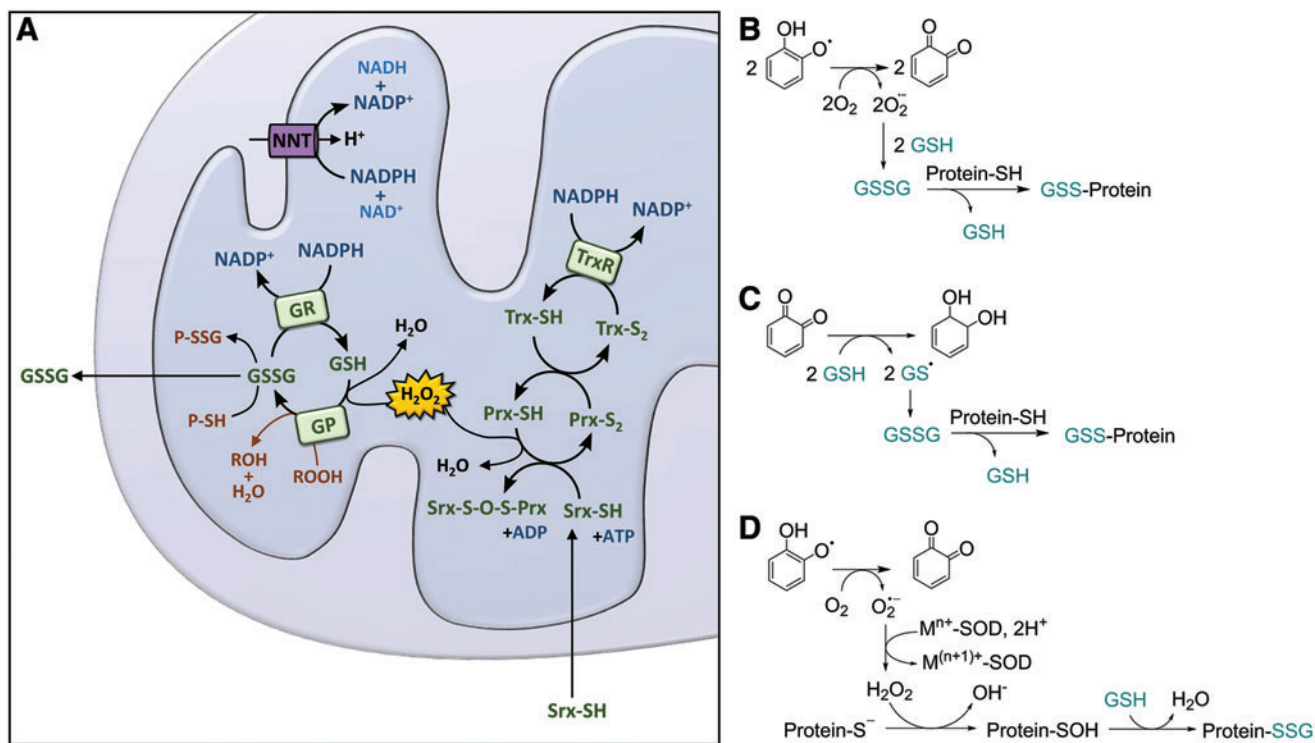


FIG. 7. Redox homeostasis in the mitochondrion. (A) ROS such as H₂O₂ can be metabolized in the conversion of GSH to GSSG by GP. GSSG can be reduced to GSH by reacting with protein sulfhydryl groups or by the action of GR. Under high oxidative stress, GSSG can also be excreted out of the mitochondria by GSH transporters, as stated (198). The antioxidant response can also be regulated by activating TrxR. Trx is oxidized by the action of Prx, and the oxidation of Prx depletes the H₂O₂ pool in the mitochondria to release H₂O. Under high oxidative stress, Prx is inactivated. Srx is translocated to mitochondria to reactivate Prx (145). Protein glutathionylation by polyphenols can principally occur *via* the following mechanisms: (B) the pro-oxidant effect of polyphenols promotes depletion of GSH under formation of GSSG potentially promoting S-glutathionylation; (C) protein glutathionylation can also occur by free radical reactions. This assumes the formation of a thiyl radical (here on GSH, GS[•], but this can also be a protein thiol, PS[•]) in agreement with the mechanisms suggested in Batinic-Haberle and Spasojevic (7) and (D) *via* protein sulfinylation. GP, glutathione peroxidase; GR, glutathione reductase; GS, glutathione synthesis; GSSG, oxidized glutathione; H₂O₂, hydrogen peroxide; NNT, nicotinamide nucleotide transhydrogenase; Prx, peroxiredoxin; Srx, sulfiredoxin; Trx, thioredoxin; TrxR, thioredoxin reductase. To see this illustration in color, the reader is referred to the web version of this article at www.liebertpub.com/ars

that act on the Trx system (55). In most reports, it is not specified which of the Trx systems, the cytosolic Trx1/TrxR-1 or the mitochondrial Trx2/TrxR-2 system, is inhibited. Mitochondrial targeting of polyphenols by mitochondriotropic derivatization has been used as an experimental strategy to determine their mitochondria-specific effects on the Trx and other redox systems (168, 169, 177).

Principally, Trx or TrxR can both be targeted by polyphenols. However, polyphenols have been most promising as inhibitors of the Trx system by targeting TrxR. TrxR activity is irreversibly inhibited on modification of the C-terminal redox active site by curcumin, thereby turning the modified enzyme into a pro-oxidant that stimulates ROS production *via* an acquired NADPH oxidase activity (37). Similarly, TrxR inhibition has been reported for myricetin and quercetin (117). Another mode of action of hijacking the thiol redox system to promote proapoptotic mechanisms in cancer cells is GSH depletion by polyphenols, shown in cell-free systems (3, 84) and in several cancer cell culture studies (84, 102).

The GSH/oxidized GSH (GSSG) system is tightly regulated (Fig. 7). In the mitochondrion, GSH plays a major role in the elimination of H₂O₂ in interplay with GSH peroxidase

(GP) or peroxiredoxin (Prx) III (isoform III is specific to the mitochondrion). The conversion of GSSG back into GSH is facilitated by the NADPH-dependent GSSG reductase. However, under conditions of increased production of ROS, a shift to elevated concentration of GSSG is proposed, which may lead to mixed disulfide formation with protein thiols and GSSG is excreted out of the mitochondria by GSH transporters (Fig. 7A).

Mailloux and Willmore suggest that S-glutathionylation plays a role in the control of mitochondrial metabolism (121). It has also been proposed that the severity of GSH depletion and the extent and kinetics of distinct protein S-glutathionylation events may determine whether cells initiate apoptosis or undergo necrosis (29). Given the properties of polyphenols, different mechanisms of polyphenol-mediated glutathionylation can be considered (Fig. 7B–D). However, the ability of polyphenols to modulate protein glutathionylation levels has not yet been explored.

In normal cells, an increased level of GSH is associated with elevated rates of proliferation, an increase in detoxification capacity, and resistance to oxidative stress. This stress response, however, allows cancer cells to adapt to elevated

levels of ROS generation under conditions of high proliferation. GSH synthesis is regulated by glutathione synthase (GS) activity, and GS expression is regulated by Nrf2. In many cancer cells, Nrf2 is upregulated, which in turn endows cancer cells even further with enhanced capacity to counteract ROS. Enhanced Nrf2 activity and elevated levels of GSH promote malignancy, cancer cell aggressiveness, and resistance to many anticancer drugs. Targeting Nrf2 therefore promises new strategies in cancer prevention and adjuvant chemotherapies (87, 126).

In the context of exploiting redox modulation to combat cancer, Prxs have emerged as interesting targets (143). Prxs catalyze the reduction of H₂O₂ and alkylperoxides to water and the respective alcohols (88, 143). Prx levels are elevated in many human cancers in line with the concept that cancer cells need to adapt to elevated ROS production. Elevated levels of Prxs in cancer cells have been correlated with a more aggressive phenotype (94, 144, 157, 189, 213, 227).

Prx III is specific to the mitochondrion. Prx III together with Trx2 and TrxR2 plays a major role in the metabolism of H₂O₂ (Fig. 7A). Prx III is sensitive to hyperoxidation at elevated levels of H₂O₂ and hyperoxidation leads to inactivation. Inactivation can be reversed by sulfiredoxin (Srx), and mitochondrial translocation of Srx in response to oxidative stress results in resistance to apoptosis (145). Thus, targeting Prx III and the mitochondrion-specific Trx2, TrxR2, and Srx in combination with established anticancer drugs may potentially sensitize cells to radiation and established chemotherapeutic agents (178, 225).

Studies exploring potential roles of polyphenols in modulating ROS levels in cancer cells are emerging. For instance, quercetin enhanced radiosensitivity of HeLa and MCF-7 tumor cells, and sensitized tumors to irradiation in a DLD-1 colorectal tumor xenograft model (113). Similarly, isoliquiritigenin increased the radiosensitivity of HepG2 cells (188). On the contrary, polyphenols (and all other antioxidants) may positively or negatively affect the efficacy of chemotherapy or radiation treatment (138, 194, 222). Further research is required to establish beneficial effects of distinct polyphenols or well-characterized botanical extracts in cancer management and treatment strategies.

Polyphenols as modulators of the Keap1/Nrf2 system in cancer

Nrf2 activation is beneficial in cancer prevention. The Keap1–Nrf2 system is considered as the master redox regulator in mammalian cells and orchestrates transcription of genes associated with regulating mitochondrial biogenesis, oxidative phosphorylation, and genes whose products have been linked to oxidative and xenobiotic stress responses (30).

Keap1 is a highly sensitive redox sensor protein that assembles with the Cul3 protein to form a Cullin-RING E3 ligase complex to facilitate the degradation of Nrf2. Under basal conditions, the levels of Nrf2 protein are kept low by the E3 ubiquitin ligase Keap1, which ubiquitinates Nrf2 in the cytoplasm and targets it for proteosomal degradation. Under conditions of oxidative stress, or in the presence of electrophilic xenobiotics, the activity of Keap1 is diminished and Nrf2 can translocate to the nucleus, bind to the antioxidant response element (ARE), which in turn induces the expression of its target genes, including many that code for

antioxidant and detoxification enzymes, stress response proteins, and ABC transporters. Nrf2 activity also regulates mitochondrial biogenesis *via* PGC-1 α and mitochondrial transcription factor A, Tfam, activation (30, 173).

Hence, Nrf2-dependent gene expression may be beneficial in cancer prevention and control strategies targeting the early stage of cancer initiation and progression. However, in late stages of malignancy, Nrf2 activity is detrimental because Nrf2 activation increases chemoresistance (87). In this context, polyphenols seem to have efficacy as chemopreventive agents in normal cells or early stages of cancer development but seem less useful in late stages of cancer due to aberrant Nrf2/Keap1 signaling and redox homeostasis at elevated ROS levels (153).

Keap1 functions essentially as sensor of cellular oxidative or electrophile stress. Keap1 harbors a large number of cysteine residues, which display reactivity toward polyphenol-derived quinones and chalcone-type flavonoids. The cysteine modification patterns affect Keap1's structural conformation and function, and, in turn, affect inhibition of Nrf2 ubiquitination (174). Human Keap1 contains a total of 27 cysteine residues (87), seven of which have been identified as highly reactive toward ROS and electrophiles by *in vitro* proteomics studies using recombinant Keap1, namely Cys151, Cys257, Cys273, Cys288, Cys297, Cys434, and Cys613 (120, 174). However, one should keep in mind that *in vitro* adduction experiments using recombinant proteins are methodology dependent and do not necessarily reflect true *in vivo* behavior.

Adduct formation depends on (i) the nucleophilicity of the cysteine thiol, which is determined by the protein microenvironment, and (ii) by the electrophilicity of the xenobiotic (203). Many plant phenolics are Michael acceptors featuring α,β -unsaturated carbonyl functionalities and are capable of forming Michael-type adducts with distinct Keap1 cysteine residues (34, 131) [Fig. 6, schema (3)]. For instance, the electrophilic chalcones, isoliquiritigenin and xanthohumol, both feature an α,β -unsaturated ketone moiety that forms Michael-type adducts with cysteine thiols of human Keap1 *in vitro*. Distinct adduction sites have been reported for these chalcones (33, 118).

As mentioned above, polyphenolic compounds exert *in vitro* antioxidant and radical scavenging activities. In addition, polyphenolic compounds are able to induce the Keap1–Nrf2 signaling pathway and the expression of ARE-response genes after oxidation to their corresponding electrophilic quinones, which contain Michael-type acceptor moieties (31). Substituted 1,2- and 1,4-diphenols but not the mono- or 1,3-dihydroxy regio-isomers can be biotransformed to the corresponding quinonoids (34). For instance, quercetin will only exhibit ARE induction after oxidation to its quinone methide (78). Similarly, EGCG contains several aromatic 1,2-dihydroxy units that can form quinones capable of activation of the Keap1/Nrf2 pathways.

In cancer patients, serum and tumor copper levels are elevated compared to healthy subjects. Copper levels positively correlate with tumor incidence, burden, malignant progression, and recurrence in many human and animal cancers (31,103). Copper promotes formation of hydroxyl radicals *via* Fenton-type chemistry, redox cycling, and formation of quinones. Dinkova-Kostova and Wang (31) hypothesized that endogenous copper levels are sufficient to enhance oxidation of polyphenols to quinone-type species that activate the Keap1/Nrf2/ARE pathway.

Aberrant Nrf2 activation promotes anabolic pathways in cancer. The Keap1-Nrf2 system in cancers might be considered a double-edged sword. In normal cells, Nrf2 activity protects cells and makes them resistant to oxidative and electrophilic stresses in line with the chemoprevention idea. However, in malignant cells, mutations and epigenetic modifications affecting the regulation and fate of Nrf2 can lead to constitutive, dominant hyperactivation of signaling pathways. They stimulate cancer cell survival and proliferation, enhance resistance to stresses, and render cancer therapies less effective (60, 92, 103, 139).

A recent study by Mitsuishi *et al.* (132) showed that, in human lung carcinoma A549 cells, Nrf2 modulated metabolic pathways that enhanced cell proliferation in the presence of active PI3K-Akt signaling. Nrf2 directly activates genes whose products are involved in the PPP and in NADPH production (namely, glucose-6-phosphate dehydrogenase, phosphogluconate dehydrogenase, transketolase, transaldolase, and malic enzyme 1) and GS. Thus, in proliferating cells, Nrf2 promotes anabolic metabolism and reinforces metabolic reprogramming. This antioxidant mechanism explains why aberrant high expression of Nrf2 in many types of tumors is associated with a poor prognosis in the clinic (79).

There is currently a large interest in deciphering factors determining the balance between a chemopreventive and the protumorigenic role of Nrf2. In this context, research is needed to further clarify to what extent polyphenols or diets rich in polyphenols might exert beneficial health effects or produce opposite effects promoting tumor growth and progression, and how they impact chemotherapeutic efficacy and radiation treatments (126).

Conclusion and Perspective

Polyphenols interfere with the function of the ETC by mild mitochondrial uncoupling through a protonophoric or an ROS-producing effect, or both. Such uncoupling effect is beneficial to promote energy expenditure in conditions of obesity-related dysfunctional metabolism. Their ability to undergo oxidation and form electrophilic quinones that stimulate the antioxidant Keap1-Nrf2 pathway suppresses the cells' transition to malignancy. Indeed, the totality of the polyphenol research indicates that polyphenols hold promise as cancer chemopreventive agents to induce adaptive stress responses that provide cytoprotection and maintenance of normal metabolic homeostasis.

Cancer cells rely less on oxidative phosphorylation than normal cells, and therefore, cancer monotherapy with polyphenols will be moderately successful at best. However, polyphenol-induced mitochondrial uncoupling can promote β -oxidation of fatty acids and anaplerotic feeding of the TCA cycle, resulting in an overall catabolic effect to counteract the glycolytic, anabolic phenotype of cancer cells. Such metabolic reprogramming might have synergistic outcomes when used in combination with chemotherapy, targeting both the anabolic phenotype and mitotic division of proliferating tumor cells.

Many of the polyphenol effects discussed in this review relate to the physicochemical properties of polyphenols (protonophore and ROS-producing effects that decrease $\Delta\psi_m$) rather than to their individual structure-specific pharmacophores targeting specific catalytic enzymes, channels,

transporters, or receptors (with moderate potency). The general protonophoric and ROS-producing effects of polyphenols hold promise for combination chemotherapy, because tumor cells cannot develop resistance against these general MMP effects. Furthermore, the MMP effects of polyphenols reverse the antiapoptotic phenotype often seen in drug-resistant cancer cells (217).

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Abbreviations Used

ADP = adenosine diphosphate
 AMPK = AMP-activated protein kinase
 ANT = adenosine nucleotide translocase
 Apo2L = Apo 2 ligand
 ARE = antioxidant response element
 ATP = adenosine triphosphate
 CypD = cyclophilin D
 DCA = dichloroacetate
 DNP = 2,4-dinitrophenol
 DR = death receptor
 EAC = esophageal adenocarcinoma
 EGCG = epigallocatechin gallate
 ERR = estrogen-related receptor
 ETC = electron transport chain
 FAO = fatty acid β -oxidation
 GP = glutathione peroxidase
 GR = glutathione reductase
 GS = glutathione synthesis
 GSH = glutathione
 GSSG = oxidized glutathione
 H₂O₂ = hydrogen peroxide
 HIF-1 = hypoxia-inducible factor-1
 HK = hexokinase
 IV = intravenous
 Keap1 = Kelch-like ECH-associated protein 1
 LDH = lactate dehydrogenase
 MMP = mitochondrial membrane potential
 Nrf2 = nuclear factor (erythroid-derived 2)-like 2
 OMM = outer mitochondrial membrane
 PDH = pyruvate dehydrogenase
 PDK = pyruvate dehydrogenase kinase
 PGC-1 α = peroxisome proliferator-activated receptor γ coactivator 1- α
 PPP = pentose phosphate pathway
 Prx = peroxiredoxin
 PTP = permeability transition pore
 ROS = reactive oxygen species
 SIRT1 = sirtuin 1
 SOD = superoxide dismutase
 Srx = sulfiredoxin
 TCA = tricarboxylic acid
 TRAIL = tumor necrosis factor-related apoptosis-inducing ligand
 Trx = thioredoxin
 TrxR = thioredoxin reductase
 UCP = uncoupling protein
 VDAC = voltage-dependent anion channel
 VEGF = vascular endothelial growth factor