



Research
Animal Nutrition and Feed Science—Review

The Biofunctions of Phytochemicals and Their Applications in Farm Animals: The Nrf2/Keap1 System as a Target

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ARTICLE INFO

Article history:

Received 5 April 2017

Revised 21 April 2017

Accepted 18 May 2017

Available online 24 July 2017

Keywords:

Phytochemical

Biofunction

Nrf2/Keap1 system

Growth performance

Meat quality

Intestinal microbiota

ABSTRACT

Reactive oxygen species (ROS) can be caused by mechanical, thermal, infectious, and chemical stimuli, and their negative effects on the health of humans and other animals are of considerable concern. The nuclear factor (erythroid-derived 2)-like 2/Kelch-like ECH-associated protein 1 (Nrf2/Keap1) system plays a major role in maintaining the balance between the production and elimination of ROS via the regulation of a series of detoxifying and antioxidant enzyme gene expressions by means of the antioxidant response element (ARE). Dietary phytochemicals, which are generally found in vegetables, fruits, grains, and herbs, have been reported to have health benefits and to improve the growth performance and meat quality of farm animals through the regulation of Nrf2-mediated phase II enzymes in a variety of ways. However, the enormous quantity of somewhat chaotic data that is available on the effects of phytochemicals needs to be properly classified according to the functions or mechanisms of phytochemicals. In this review, we first introduce the antioxidant properties of phytochemicals and their relation to the Nrf2/Keap1 system. We then summarize the effects of phytochemicals on the growth performance, meat quality, and intestinal microbiota of farm animals via targeting the Nrf2/Keap1 system. These exhaustive data contribute to better illuminate the underlying biofunctional properties of phytochemicals in farm animals.

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1. Biofunctions of dietary phytochemicals in modulating the Nrf2/Keap1 system

1.1. Dietary phytochemicals

Phytochemicals are produced via primary or secondary plant metabolisms and originate in various kinds of fruits, vegetables, grains, and herbs, endowing them with the color, taste, smell, and other organoleptic properties of the plants [1]. They are produced to help plants thrive or to thwart competitors, predators, or pathogens. During the last two decades, dietary phytochemicals have been found to be strongly associated with human health and diseases through their biological functions [2,3]. More than 10 000 kinds of dietary phytochemicals have been classified into carotenoids, isothiocy-

anates, and polyphenols based on their chemical structure. Among these, the best-investigated category is that of polyphenols, which mainly include phenolic acids, flavonoids, and stilbenes/lignans. Many epidemiological investigations and lab-based studies have demonstrated that most polyphenols are conducive to the chemoprevention of several chronic diseases, including diabetes, cardiovascular diseases, neurodegenerative diseases, cancer, and other inflammatory diseases [4].

1.2. Phytochemicals as modulators for the Nrf2/Keap1 system

When phytochemicals are ingested by humans and other animals, they are recognized as xenobiotics. As a result, they stimulate the genes of a series of antioxidant and detoxifying enzymes (ADEs)

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to express. Most of these genes contain a specific conserved nucleotide sequence of 5'-TA/CANNA/GTGAC/TNNNGCA/G-3' in their promoters, named antioxidant response element (ARE)/electrophile-responsive element (EpRE) [5]. Nuclear factor (erythroid-derived 2)-like 2 (Nrf2) has been demonstrated to strongly activate ARE/EpRE to enhance the gene expressions of a series of ADEs [6], such as NAD(P)H quinone dehydrogenase 1 (NQO1), glutathione reductase (GSR), and solute carrier family 7 member 11 (SLC7A11) [7]. Nrf2 is a transcription factor (TF) transcribed by the *NFE2L2* gene in humans, with a basic leucine zipper (bZIP) protein that induces the gene expressions of phase II antioxidant proteins and detoxifying enzymes in order to protect against oxidative damage triggered by chronic inflammation and injury [2]. The crucial negative regulator of Nrf2 is Kelch-like ECH-associated protein 1 (Keap1), which maintains the dynamic balance of cytoplasmic Nrf2 by proteasomal degradation [8].

The molecular mechanisms of Nrf2-ARE activation are summarized in the schematic diagram in Fig. 1. As shown in this diagram, the mechanisms of the regulating Nrf2/Keap1 system can be divided into Keap1-dependent and Keap1-independent mechanisms. Under basal conditions, Keap1 inhibits Nrf2 by functioning as an E3 ubiquitin ligase with the cullin 3-RING box protein 1 (Cul3-Rbx1) system for the constant ubiquitination and proteasomal degradation of Nrf2. Under induced status, electrophiles, oxidants, or phytochemicals can influence the Keap1 structure/residues, in the forms of cysteine modification, ubiquitination, phosphorylation, and succination,

causing Nrf2 to escape from the Keap1-dependent ubiquitination system [9]. Alternatively, stress inducement may stimulate the phosphorylation of certain protein kinases, such as mitogen-activated protein kinases (MAPKs), phosphoinositide 3-kinase (PI3K), protein kinase C (PKC), PKR-like endoplasmic reticulum kinase (PERK), glycogen synthase kinase 3 (GSK3), or Nrf2 itself, thus regulating the activity of a series of TFs or certain nuclear proteins such as positive Brahma-related gene 1 (BRG1), nuclear receptor coactivator amplified in breast cancer 1 (AIB1), and Maf, as well as negative p53, p65, and cFos [6,8]. Moreover, phytochemicals may cause epigenetic modifications to affect the mRNA transcription of *NFE2L2* or *Keap1*, such as DNA methylation, histone modification, and microRNA tuning. All of the above result in the accumulation of Nrf2 in the nucleus to heterodimerize with small Maf or CREB-binding protein (CBP) and to bind to ARE, which finally activates the expression of its downstream ADEs genes [6,10].

1.3. Molecular mechanism underlying Nrf2 regulation by dietary phytochemicals

An extremely large number of studies performed *in vitro* and *in vivo* have revealed that many dietary phytochemicals have powerful abilities in regulating the Nrf2/Keap1 system [2–4]. However, the molecular mechanisms underlying this huge quantity of data are not well classified. Here, based on the current research status, we

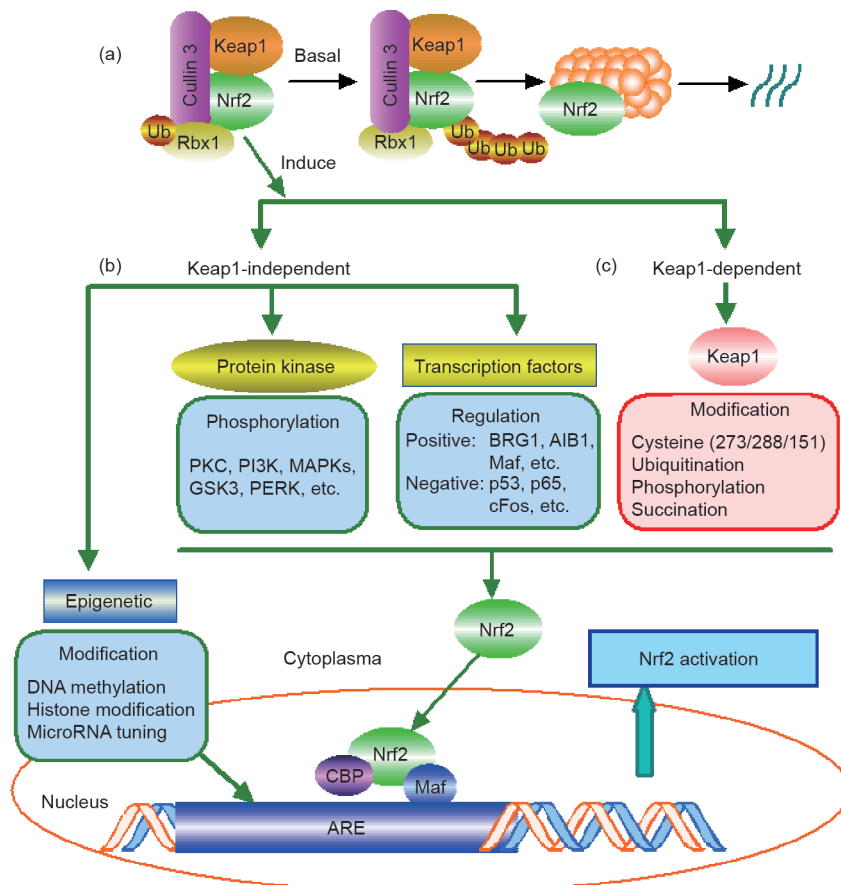


Fig. 1. Schematic diagram of the molecular mechanisms underlying the modulation of the Keap1/Nrf2 pathway. (a) Under normal/basal conditions, Nrf2 is inhibited by the Keap1-mediated Cul3-Rbx1 ubiquitination system for general proteasomal degradation. Under an induced state/stimulation, Nrf2 is activated by the Keap1-independent or Keap1-dependent Nrf2 pathway. (b) The Keap1-independent pathway. The protein kinases (PKC, PI3K, MAPKs, GSK3, and PERK) can phosphorylate Nrf2, and some transcription factors bind to ARE in order to positively or negatively regulate the expressions of Nrf2/ARE-mediated genes (positive regulators include BRG1, AIB1, and Maf, and negative regulators include p53, p65, and cFos). Epigenetic modifications include DNA methylations of promoters, histone modifications such as acetylations or methylations, and microRNA tuning by transcriptional regulations. (c) Keap1-dependent pathway. The cysteine modifications in the locations of Cysteine 273, 288, and 151, ubiquitination, phosphorylation, and succination of Keap1 are minimally involved.

review the molecular mechanisms of Nrf2 regulation by dietary phytochemicals and classify them into Keap1-dependent and Keap1-independent mechanisms.

1.3.1. Keap1-dependent pathway

Several models have been suggested to explain the inhibitory regulation of Nrf2 by Keap1. Most of the ARE inducers can target and modify the cysteines of Keap1 to affect Nrf2-ARE signaling. It is interesting that the location of the Keap1 cysteine that is targeted differs, depending on the type of the inducer [9,10]. The essential cysteine residues generally involve C288, C273, and C151 [11]. After the discovery of Keap1 as an E3 ligase substrate adaptor of the Cul3-Rbx1-containing ubiquitination system, the “Keap1 dissociation and Cul3-Rbx1 ubiquitination” model was developed to explain the major Nrf2 regulation mechanism [12]. Moreover, several other important models such as the “Keap1 hinge-and-latch,” “Keap1 phosphorylation,” “Keap1 ubiquitination,” and “Keap1 succination”

models reveal that modifications of Keap1 caused by a variety of stimuli constitute a primary mechanism in the modulation of the Nrf2/Keap1 system [13–17].

An enormous number of dietary phytochemicals have been found to modify the cysteines of Keap1 to regulate the Nrf2/Keap1 system. As displayed in Table 1 [18–80], sulforaphane, resveratrol, catechol estrogens, quercetin, carnosic acid, baicalein, glyceollins, oridonin, faltarindiol, piceatannol, xanthohumol, and 6-(methylsulfinyl)hexyl isothiocyanate were reported to activate the Nrf2/Keap1 system. Of these, quercetin works in the “Keap1 dissociation” model [18] and baicalein works in the “Keap1 ubiquitination” model; it is noteworthy that baicalein also works in the “Keap1 hinge-and-latch” model [26]. In addition, sulforaphane works in the “Keap1 hinge-and-latch” model in human Keap1, whereas it works in the “Keap1 dissociation” model in animal Keap1 [56–59]. These data suggest that Keap1 modification by phytochemicals varies, and that the cell model used is an important factor.

Table 1

The molecular mechanisms of Nrf2 regulation by phytochemicals.

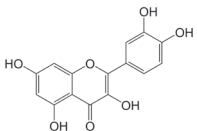
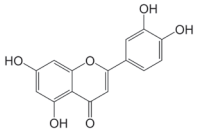
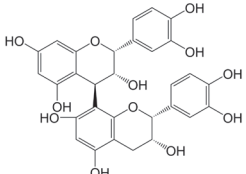
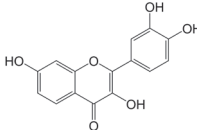
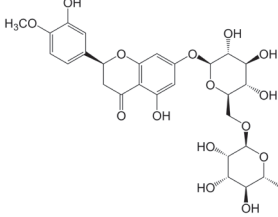
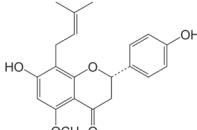
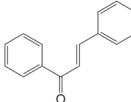
Classification	Origin	Compound	Structure	Dose	Time	Mechanism	Model	Refs.
Activation of Nrf2-ARE pathway								
Flavonoid-type polyphenols	Apple, tea, caper, lovage, onion	Quercetin		0–40 $\mu\text{mol}\cdot\text{L}^{-1}$	6 h	\uparrow Keap1 modification, Nrf2 stability	HepG2 cells	[18]
				100–200 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h, 48 h	\uparrow p38 MAPK and ERK	Human hepatocytes epithelial cells	[19]
	Celery, green pepper	Luteolin		0–20 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h, 72 h	\uparrow ERK1/2, HO-1, ARE binding	PC12 cells	[20]
	Cocoa, red wine	Procyanidin B2		10 $\mu\text{mol}\cdot\text{L}^{-1}$	20 h	\uparrow ERKs and p38 MAPK	Human colonic cells	[21]
	Strawberry	Fisetin		0–25 $\mu\text{mol}\cdot\text{L}^{-1}$	NM	\uparrow PKC- δ and p38 MAPK	Human umbilical vein endothelial cells	[22]
	Citrus fruits	Hesperidin		0–80 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow ERK1/2	Human hepatic L02 cells	[23]
	Hops	Xanthohumol		4 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow Modification of Keap1 cysteine	Murine Hepa1c7 cells	[24]
	Plant phenols	Chalcone		10–25 $\mu\text{mol}\cdot\text{L}^{-1}$	NM	\uparrow Nrf2, HO-1	Endothelial cells	[25]

Table 1 (continued)

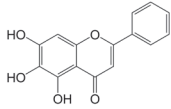
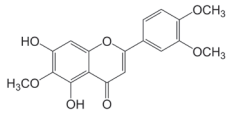
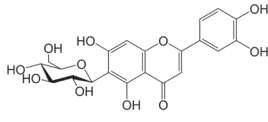
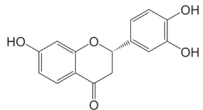
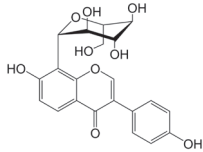
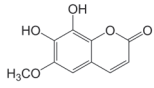
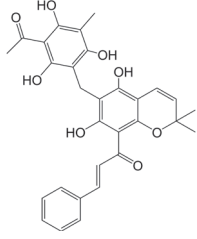
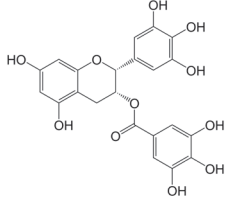
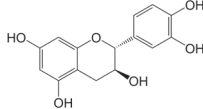
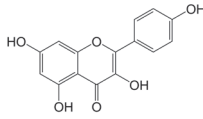
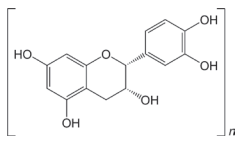
Classification	Origin	Compound	Structure	Dose	Time	Mechanism	Model	Refs.
	<i>Scutellaria baicalensis</i>	Baicalein		0–40 $\mu\text{mol}\cdot\text{L}^{-1}$	9 h, 24 h	\uparrow Nrf2, HO-1	HepG2 cells	[26]
	Artemisia	Eupatilin		0–150 $\mu\text{mol}\cdot\text{L}^{-1}$	16 h	\uparrow ERK	Feline ileal smooth muscle cells	[27]
	<i>Sasa borealis</i>	Isoorientin		5 $\mu\text{g}\cdot\text{mL}^{-1}$	0–6 h	\uparrow PI3K/Akt	HepG2 cells	[28]
	<i>Vernonia anthemintica</i> , <i>Dalbergia odorifera</i>	Butin		10 $\mu\text{g}\cdot\text{mL}^{-1}$	12 h, 24 h	\uparrow PI3K/Akt	Chinese hamster lung fibroblast (V79-4)	[29]
	<i>Inula helenium</i>	Phytoestrogen puerarin		0–100 $\mu\text{mol}\cdot\text{L}^{-1}$	2–18 h	\uparrow PI3K/Akt	Hepa1c1c7 cells	[30]
	<i>Fraxinus rhin-chophylla</i>	Fraxetin		30–100 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow Nrf2, HO-1	Vascular smooth muscle cells	[31]
	<i>Mallotus philippinensis</i>	Rottlerin		1–10 $\mu\text{mol}\cdot\text{L}^{-1}$	9 h	\uparrow ERK and p38 MAPK	HT29 cells	[32]
Tea		EGCG		20 $\mu\text{mol}\cdot\text{L}^{-1}$	48 h	\uparrow p38 MAPK and Akt	B lymphoblasts	[33]
				50 $\mu\text{mol}\cdot\text{L}^{-1}$	6 h	\uparrow ERK and PI3K/Akt	Bovine aortic endothelial cells	[34]
Cocoa, tea		Epicatechin		5–30 $\text{mg}\cdot\text{kg}^{-1}$ BW	1 h, 6 h, 18 h	\uparrow ERK and PI3K/Akt	Ischemic damaged mice	[35]
Tea, broccoli		Kaempferol		0–10 $\mu\text{mol}\cdot\text{L}^{-1}$	18 h	\uparrow JNK, HO-1, GCLC	Organ of Corti 1 (HEI-OC1) cells	[36]
Wild grape		Procyanidins		25 $\mu\text{g}\cdot\text{mL}^{-1}$	1 h	\uparrow p38 MAPK, PI3K/Akt	HepG2 cells	[37]

Table 1 (continued)

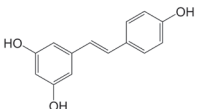
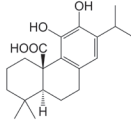
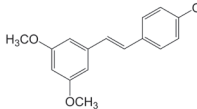
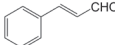
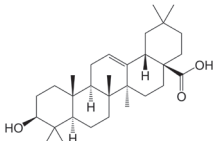
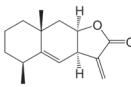
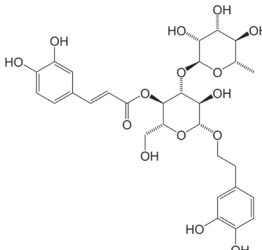
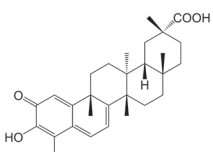
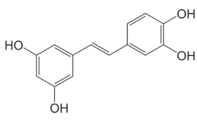
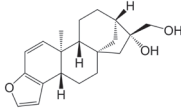
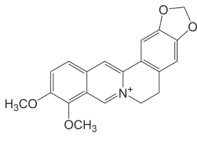
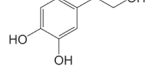
Classification	Origin	Compound	Structure	Dose	Time	Mechanism	Model	Refs.
Non-flavonoid-type polyphenols	Red grape	Resveratrol		10 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow Modification of Nrf2 and Keap1	A549 cells	[38]
			15 $\mu\text{mol}\cdot\text{L}^{-1}$	0–6 h	\uparrow ERK and PI3K	PC12 cells	[39]	
	Rosemary, common sage	Carnosic acid		1–20 $\mu\text{mol}\cdot\text{L}^{-1}$	0–1 h	\uparrow p38 MAPK		[40]
			10 $\mu\text{mol}\cdot\text{L}^{-1}$	1 h	\uparrow S-alkylation of Keap1		[41]	
	Blueberries, grapes	Pterostilbene		5 $\text{mg}\cdot\text{kg}^{-1}$ BW	6 weeks	\uparrow Nrf2, HO-1	Male BALB/c mice	[42]
	<i>Cinnamomum cassia</i> Presl	Cinnamaldehyde		50–100 $\mu\text{mol}\cdot\text{L}^{-1}$	0–12 h	\uparrow Nrf2, HO-1	Endothelial cells	[43]
	American pokeweed, garlic	Oleanolic acid		10–50 $\mu\text{mol}\cdot\text{L}^{-1}$	0–2 h	\uparrow Akt and ERK	Primary rat vascular smooth muscle cells	[44]
	<i>Inula helenium</i>	Alantolactone		0–10 $\mu\text{mol}\cdot\text{L}^{-1}$	NM	\uparrow PI3K and JNK	Hepa1c1c7 mouse hepatoma cells	[45]
	Scrophulariaceae	Acteoside		30 $\mu\text{mol}\cdot\text{L}^{-1}$	0–12 h, 6 h	\uparrow ERK and PI3K/Akt	PC12 cells	[46]
	<i>Tripterygium wilfordii</i>	Celastrol		0–1 $\mu\text{g}\cdot\text{mL}^{-1}$	0.5 h	\uparrow ERK and p38 MAPK	HaCaT cells	[47]
	<i>Euphorbia lagascae</i>	Piceatannol		30 $\mu\text{mol}\cdot\text{L}^{-1}$	0–12 h	\uparrow Akt and modification of Keap1	MCF10A cells	[48]
	Coffee	Kahweol		0–10 $\mu\text{mol}\cdot\text{L}^{-1}$	0–2 h	\uparrow Akt and p38 MAPK	SH-SY5Y cells	[49]
	<i>Rhizoma coptidis</i>	Berberine		1–10 $\mu\text{mol}\cdot\text{L}^{-1}$	0–2 h	\uparrow PI3K/Akt, phosphorylation of Nrf2	Rat brain astrocyte cell line (RBA-1)	[50]
	Olive	Hydroxytyrosol		50 $\mu\text{mol}\cdot\text{L}^{-1}$	0–1 h	\uparrow PI3K/Akt, MEK1/2-ERK1/2	Vascular endothelial cells	[51]
				0–200 $\mu\text{mol}\cdot\text{L}^{-1}$	2–24 h	\uparrow JNK	Human retinal pigment	[52]

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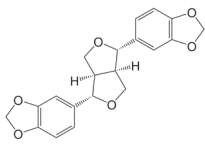
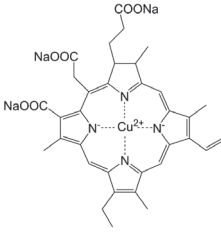
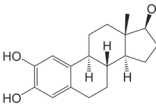
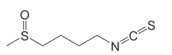
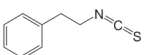

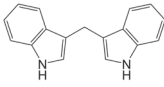
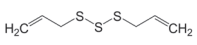
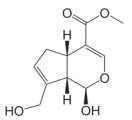
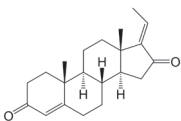
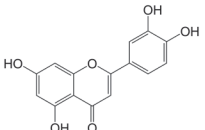
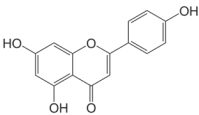
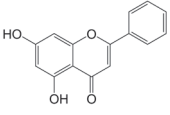
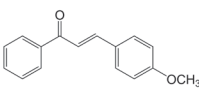
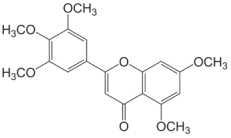
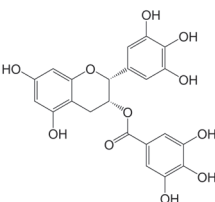
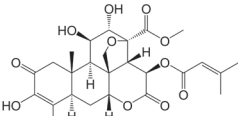
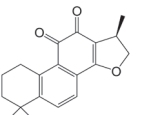
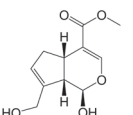
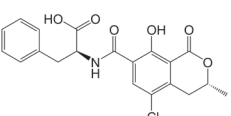
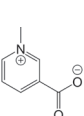
Classification	Origin	Compound	Structure	Dose	Time	Mechanism	Model	Refs.
	Sesame seeds	Sesamin and episesamin		0–10 $\mu\text{mol}\cdot\text{L}^{-1}$	0–2 h	\uparrow p38 MAPK	Rat pheochromocytoma PC12 cells	[53]
	Spinach, green leafy vegetables	Chlorophyllin		50 $\mu\text{mol}\cdot\text{L}^{-1}$	0–2 h	\uparrow PI3K/Akt	Human umbilical vein endothelial cells	[54]
	Soybean	Catechol estrogens		10 $\mu\text{mol}\cdot\text{L}^{-1}$	3 h	\uparrow Modification of Keap1	RAW264.7 cells	[55]
Isothiocyanates and other phytochemicals	Cruciferous vegetables	Sulforaphane		0–200 $\mu\text{mol}\cdot\text{L}^{-1}$	2 h	\uparrow Cysteine thioacetylation of Keap1	Human Keap-1-transfected HEK293 cells	[56]
				20 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow p38 MAPK isoforms	HepG2 cells	[57]
				20 $\mu\text{mol}\cdot\text{L}^{-1}$	1 h	\uparrow ERK and PI3K	Caco-2 cells	[58]
				0–2.5 $\mu\text{mol}\cdot\text{L}^{-1}$	5 d	\uparrow CpGs, demethylation of Nrf2 promoter, Nrf2, NQO1; \downarrow DNMT1/3a, HDAC1/4/5/7	TRAMP C1 cells	[59]
Cruciferous vegetables		PEITC		5 $\mu\text{mol}\cdot\text{L}^{-1}$	12 h	\uparrow ERK and JNK	PC-3 cells	[60]
Cruciferous vegetables		I3C		6.25 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow JNK	HepG2-C8 cells	[61]
Cruciferous vegetables		DIM		0–5 $\mu\text{mol}\cdot\text{L}^{-1}$	NM	\uparrow CpGs, demethylation of Nrf2 promoter, Nrf2, NQO1, JNK	TRAMP-C1 cells, TRAMP prostate tumors	[62]
Garlic, onion		Diallyl trisulfide		100 $\mu\text{mol}\cdot\text{L}^{-1}$	1 h	\uparrow Calcium-dependent signaling, ERK, p38 MAPK	HepG2 cells	[63]
<i>Gardenia jasminoides</i>		Genipin		0–100 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h	\uparrow PI3K-JNK1/2	RAW264.7 macrophages	[64]
<i>Commiphora mukul</i>		Guggulsterone		25 $\mu\text{mol}\cdot\text{L}^{-1}$	0–2 h, 6 h	\uparrow PI3K/Akt	Human mammary epithelial cells	[65]
Inhibition of Nrf2-ARE pathway								
Flavonoid	Celery, green pepper	Luteolin (Lut) ^a		20 $\mu\text{mol}\cdot\text{L}^{-1}$	24 h, 48 h	\uparrow Nrf2 mRNA degradation	A549, HCT116-OX, SW620OX, MDA-MB 231 cells	[66,67]

Table 1 (continued)

Classification	Origin	Compound	Structure	Dose	Time	Mechanism	Model	Refs.
	Parsley, celery, celeriac	Apigenin (Api) ^a		20 μmol·L ⁻¹	14 d	↓ p-Akt	Tumor of mice	[68]
	<i>Passiflora incarnata</i>	Chrysin (Chry) ^a		10–20 μmol·L ⁻¹	24 h	↓ p-Akt, p-ERK1/2, Nrf2 protein levels	BEL-7402/ADM cells	[69]
		4-methoxychalcone ^a		5 μg·mL ⁻¹	3–24 h	↓ p-Akt (Thr308)	A549 cells	[70]
	Tangerine peel	3',4',5',5',7-pentamethoxyflavone ^a		10–25 μmol·L ⁻¹	24 h	↓ p-ERK	A549 cells	[71]
	Tea	(EGCG) ^a		100 μmol·L ⁻¹ , 200 μmol·L ⁻¹	24 h	↓ Nrf2 protein level; ↑ apoptosis	A549 cells	[72]
	<i>Brucea</i>	Brusatol (Bru)		10–300 nmol·L ⁻¹	2 h	↓ Nrf2 mRNA translation	A549, Hep- pa1c1c7 cells	[73]
	<i>Salvia</i>	Cryptotanshinone		5–10 μmol·L ⁻¹	24 h	NM	H1299 cells	[74]
		Metformin (Met)		1–5 mmol·L ⁻¹	24 h	↓ pRaf, p-ERK1/2; ↑ microRNA-34a; ↓ Nrf2	HepG2, HeLa, A549, MCF-7 cells	[75,76]
		Mycotoxin ochratoxin A		5 μmol·L ⁻¹	1 d, 3 d	↓ Nuclear import of Nrf2; ↓ DNA binding; ↑ microRNA-32; ↓ Nrf2	Human primary proximal tubule cells	[77,78]
	Leguminosae extract of fenugreek	Trigonelline (Trig)		0.0001–1 mmol·L ⁻¹	3 h	↓ Nuclear import of Nrf2	Panc1, Colo357, MiaPaca2 cells	[79,80]

CpG: 5'-C-phosphate-G-3'; DIM: 3,3'-diindolylmethane; DNMT: DNA methyltransferase; EGCG: epigallocatechin-3-gallate; ERK: extracellular signal-regulated kinase; GCLC: glutamate-cysteine ligase catalytic subunit; HDAC: histone deacetylase; HO-1: heme oxygenase 1; I3C: indole-3-carbinol; JNK: c-Jun N-terminal kinase; MEK: mitogen-activated protein kinase kinase; PEITC: phenethyl isothiocyanate; BW: body weight; NM: not mentioned in the reference.

^a indicates that the compound has a dual role in the regulation of the Nrf2-ARE pathway, including activation and inhibition.

1.3.2. Keap1-independent pathway

Aside from Keap1, a large number of other factors have been proven to play significant roles in the regulation of the Nrf2/Keap1 system. As shown in Fig. 1, these factors can be mainly classified into epigenetic modifications, the phosphorylation of protein kinases, and the regulation of TFs.

As shown in Table 1, the phosphorylation of extracellular signal-regulated kinase (ERK) can be promoted by quercetin [19], sulfora-

phane/phenethyl isothiocyanate (PEITC) [58,60], hydroxytyrosol [51], resveratrol [39], luteolin [20], procyanidin B2 [21], hesperidin [23], oleanolic acid [44], epigallocatechin-3-gallate (EGCG) [34], epicatechin [35], eupatilin [27], rottlerin [32], acteoside [46], and celastrol [47]. The activation of p38 MAPK can occur from treatments of quercetin [19], procyanidins [37], sulforaphane [57], procyanidin B2 [21], fisetin [22], rottlerin [32], carnosic acid [40], celastrol [47], sesamin/episesamin [53], EGCG [33], and kahweol [49]. The activity of c-Jun

N-terminal kinase (JNK) has been reported to be induced by treatments of alantolactone [45], hydroxytyrosol [52], PEITC [60], kaempferol [36], genipin [64], and indole-3-carbinol/3,3'-diindolylmethane [61,62]. The activity of PI3K can be stimulated by procyanidins [37], sulforaphane [58], hydroxytyrosol [51], resveratrol [39], chlorophyllin [54], genipin [64], isoorientin [28], butin [29], guggulsterone [65], alantolactone [45], phytoestrogen puerarin [30], berberine [50], acetoide [46], EGCG [34], and epicatechin [35].

Several lines of research have found that the Nrf2/Keap1 system can be regulated by dietary phytochemicals through modulation of other transcriptional factors or nuclear proteins. Jun dimerization protein 2 (JDP2) was found to be strongly associated with sulforaphane-induced Nrf2 activation, and it was shown that JDP2 positively promotes Nrf2-ARE activation caused by sulforaphane [81]. Another study reported that sulforaphane inhibited the Nrf2 signaling pathway at the transcription level via nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B); NF- κ B promotes histone deacetylase 3 (HDAC3) to cause local hypoacetylation and competes against Nrf2 transactivation with CBP to inhibit Nrf2 signaling [82].

Some dietary phytochemicals are considered to be potent epigenetic modifiers, including isothiocyanates, tea polyphenols, genistein, and curcumin [83]. Sulforaphane, 3,3'-diindolylmethane, curcumin, and Z-ligustilide were shown to inhibit the expressions of DNA methyltransferase (DNMT) and HDAC, resulting in the demethylation of Nrf2 promoter and the reactivation of Nrf2 signaling in the prostate of TRAMP mice or in TRAMP C1 cells [59,62,84,85]. Moreover, apigenin, sulforaphane, and tanshinone IIA were reported to demethylate excessively methylated 5'-C-phosphate-G-3' (CpG) sites in the Nrf2 promoter region in mouse skin epidermal JB6 P+

cells, which was associated with the reactivation of Nrf2 signaling, the expression of Nrf2 target genes, the suppression of TPA-induced transformation, and the inhibition of protein expression of DNMTs and HDACs [86–88]. These findings suggest that phytochemicals can regulate Nrf2 expression epigenetically; however, the exact effects of these special Nrf2 modulators on cancer and other chronic diseases need to be clarified by further study.

It is interesting that several lines of study reported that some flavonoids work as inhibitors of the Nrf2/Keap1 system in certain cancer cell lines and play an important role in overcoming cancer drug resistance (Table 1) [66–80]. For example, luteolin, apigenin, chrysin, 4-methoxychalcone, pentamethoxyflavone, and EGCG have been found to play different roles in Nrf2-ARE regulation in normal cells and in cancer cells. In normal cells, they work as activators for Nrf2-ARE regulation to prevent chronic diseases, whereas in cancer cells, they work as inhibitors for Nrf2-ARE regulation to overcome cancer drug resistance. This dual action of phytochemicals on the Nrf2/Keap1 system in normal and cancer cells is attracting considerable attention regarding its health benefits [89].

2. The effects of phytochemicals on the growth performances, meat quality, and intestinal microbiota of farm animals by targeting the Nrf2/Keap1 system

The source of phytochemicals for farm animals is generally agroindustrial byproducts, such as skins, stems, seeds, pomace, nuts, hulls, and waste from the production of juice, wine, or beer, in order to reduce feed cost [90]. Table 2 summarizes the effects of phytochemicals on the growth performances, meat quality, and intestinal microbiota of farm animals [91–135].

Table 2

The effects of phytochemicals on the growth performances, meat quality, and intestinal microbiota of farm animals.

Function classification	Phytochemical	Concentration	Animal/meat tested	Effect	Refs.
Growth performances					
	Resveratrol and resveratrol-rich grape extract	100 mg·(kg·d) ⁻¹	Pigs	Lower fat deposition, improve myocardial function or glucose metabolism, prevent development of atherosclerotic lesions and coronary heart disease	[91–93]
	Polyphenol-rich grape seed and grape marc meal		Pigs	No change in Nrf2/Keap1 pathway	[94,95]
	Grape seed proanthocyanidin extract		Broilers	Improve weight gain and lower mortality of broilers infected with <i>Eimeria tenella</i>	[96]
	Thymol, tannic acid, or gallic acid	200 mg, 5 g·kg ⁻¹ diet	Broilers	Improve the feed utilization and final BW	[97]
	Grape pomace	60 g·kg ⁻¹ diet	Broilers	Improve feed efficiency	[98]
	Green tea polyphenols		Broilers	Improve the feed conversion ratio and impair feed efficiency without corticosterone treatment	[99]
	Resveratrol	1% of diet	Broilers	Impair body weight gain and feed conversion ratio	[100]
	Quercetin	0.2–0.6 g·kg ⁻¹ diet	Hens	Increase laying rate, decrease feed-to-egg ratio	[101]
	Tea polyphenols	5–15 mg·kg ⁻¹ diet	Laying hens	Prevent the adverse effect of vanadium on egg quality	[102]
	Pomegranate-extract polyphenols	5–10 g·d ⁻¹	Dairy cows	Decreased the digestibility of protein and fat	[103]
	Polyphenol-rich grape seed and grape marc meal extract		Dairy cows	Improve milk performance	[104]
	Green tea and curcuma extract		Dairy cows	Cause a reduction of fat content in the liver and an increase in milk performance	[105]
Meat quality					
Antioxidant	Quercetin, a flavonoid; ampelopsin, isoflavones, a polyphenols mix	10 mg·(kg·d) ⁻¹ , 1 g·kg ⁻¹ diet	Pigs	Reduce plasma lipid peroxidation and lower MDA level	[94,106,107]
	Tea polyphenols, grape seed proanthocyanidin extract	1000 mg·kg ⁻¹ diet	Broilers and laying hens	Reduction of MDA and TBARS concentrations, induction of GPx activity	[96,98,102]
	Extracts of rosemary, grape skin, green tea, and coffee	50–200 ppm	Pork patties	Reduce lipid oxidation, reduce values of TBARS and hexanal	[108]

Table 2 (continued)

Function classification	Phytochemical	Concentration	Animal/meat tested	Effect	Refs.	
Anti-inflammatory	Extracts of white peony, red peony, moutan peony, sappan wood, rehmannia, and angelica	0.5%–2.0%	Raw and cooked goat meat patties	Reduce lipid oxidation	[109]	
	Extracts of olive leaf, date pits, and rosemary leaf		Raw beef patties, ground beef, and buffalo meat patties	Reduce TBARS value, lipid oxidation, and oxymyoglobin oxidation	[110–112]	
	Adzuki bean extract and grape seed extract		Pork and beef sausages	Reduce lipid oxidation and TBARS values	[113,114]	
	Garlic juice	1% and 3%	Emulsified sausage	Decrease peroxide value, TBARS, and residual nitrite	[115]	
	Sage essential oil	3%	Raw pork	Decrease the TBARS value	[116]	
	Oregano essential oil	3%	Pork and beef	Lower levels of oxidation	[116]	
	Grape seed and grape marc meal extract or hop extract		Growing pigs	Downregulation of various pro-inflammatory genes	[95]	
	Cocoa powder	2.5 g, 10 g, 20 g	Pigs	Decrease gene expression of TNF- α and Toll-like receptors	[117]	
	Tea polyphenols	0.03–0.09 g·kg ⁻¹ BW	Broilers	Downregulation of the genes of IL-1 β , IL-4, IL-10, TNF- α , and IFN- γ	[118]	
	Pomegranate-extract polyphenols	5–10 g·d ⁻¹	Pigs	Increase the secretion of IFN- γ and IL-4, improve total IgG response	[119]	
Sensory	Grape seed and grape marc meal extract		Dairy cows	Downregulation of the marker of endoplasmic reticulum stress, FGF-21, and fat accumulation in the liver	[104]	
	White peony extract	0.5%–2.0%	Raw and cooked meat patties	Increase the redness value (a^* value)	[109]	
	Rosemary extract	300–500 ppm	Raw frozen sausage	Maintain the red color	[120]	
	Green tea extract	300 mg·kg ⁻¹ meat	Raw patties	Decrease a^* value	[121]	
			Cooked patties	Delay rancid flavor development	[122]	
	Grape seed extract	0.01%–0.02%	Beef patties	Reduce visual green discoloration	[123]	
	Myrtle extract	10%	Beef patties	Prevent color changes	[124]	
	<i>Eleutherine americana</i> extract	2.7–10.8 mg·(100 g) ⁻¹	Cooked pork	Increase a^* value	[125]	
	Adzuki bean extract	0.2%	Cured and uncured cooked pork sausages	Increase a^* value but decrease lightness (L^* value) and yellowness (b^* value)	[126]	
	Green tea extract	500–6000 ppm	Raw and cooked goat meat	Increase a^* value	[127]	
	Grape seed extract			Decrease a^* value	[128]	
	Pepper extract		Cooked pork	Maintain a^* value	[128]	
	Curry leaf extract	5 mL·(500 g) ⁻¹ meat	Raw ground pork	Decrease L^* value and a^* value while increasing b^* value	[129]	
	Rosemary leaf extract	130 ppm	Raw and cooked ground buffalo meat patties	Stabilized color	[130]	
	Plum products		Variety of meat and poultry products	Minor effect on flavor but caused color change	[131,132]	
	Intestinal microbiota	Grape seed extract		Meat products	Significant change in color	[133]
		Cocoa powder		Pigs	Increase the abundance of <i>Lactobacillus</i> , <i>Bifidobacterium</i> spp., <i>Bacteroides-Prevotella</i> , and <i>Faecalibacterium prausnitzii</i>	[117,134]
Grape pomace concentrate			Broilers	Increase the abundance of <i>Enterococcus</i> and decrease that of <i>Clostridium</i>	[98]	
Quercetin			Laying hens	Decrease the total aerobes and coliforms and increase the abundance of <i>Bifidobacterium</i>	[101]	
Tea polyphenols			Pigs	Increase the amount of lactobacilli and decrease that of the total bacteria, Bacteroidaceae, and <i>Clostridium perfringens</i>	[102]	
			Calves	Decrease <i>Bifidobacterium</i> spp., <i>Lactobacillus</i> spp., and <i>Clostridium perfringens</i>	[135]	

FGF: fibroblast growth factor; GPx: glutathione peroxidase; IFN: interferon; IgG: immunoglobulin G; IL: interleukin; MDA: malondialdehyde; TBARS: thiobarbituric acid reactive substance; TNF: tumor necrosis factor; BW: body weight.

2.1. Growth performances

Several lines of study have reported the effects of phytochemicals on the growth performances of farm animals including pigs, poultry, and cattle.

In pigs, stilbenoid resveratrol and grape extract with a rich concentration of resveratrol were found to lower fat deposition, improve glucose metabolism and myocardial function, and prevent the progression of atherosclerotic lesions and coronary heart disease [91–93]. Although an improvement in growth performance was found in pig feed based on polyphenol-rich grape seed and marc meal, the activity of TF Nrf2 and the expressions of ARE-associated antioxidant genes or detoxifying enzymes showed no significant change [94,95].

In broilers and laying hens, phytochemicals were found to have a significant effect on improving growth performances. The supplementation of proanthocyanidin extract from grape seed was reported to lower the mortality of broilers infected with *Eimeria tenella* and improve their weight gain [96]. The broiler diet, which contained thymol (200 mg·kg⁻¹ diet), gallic acid (5 g·kg⁻¹ diet), and tannic acid (5 g·kg⁻¹ diet), was found to improve the feed utilization and final body weight [97]. Grape pomace concentrate (60 g·kg⁻¹ diet) was found to improve feed efficiency [98]. Green tea polyphenols in the broiler diet improved the feed conversion ratio in liver and muscle treated with corticosterone, but impaired the feed efficiency without corticosterone treatment [99]. Resveratrol (1% of diet) impaired the body weight gains of broiler birds as well as their feed conversion ratios [100]. Dietary quercetin (0.2–0.6 g·kg⁻¹ diet) was found to increase the laying rate and decrease the feed-to-egg ratio [101]. Tea polyphenols (5–15 mg·kg⁻¹ diet) were reported to prevent the adverse effect of vanadium on egg quality [102].

In dairy cows, pomegranate-extracted polyphenols decreased the digestibility of protein and fat due to the suppression of these nutrients by high tannins content [103]. The supplementation of dairy cow feed with polyphenol-rich grape seed and marc extract was found to improve milk performance [104]. Plant products were shown to cause a reduction of fatty liver formation and an improvement in milk performance in cows [105].

Although direct proof of the link between phytochemical-caused improvements on the growth performance of farm animals and the Nrf2/Keap1 system has not yet been fully established, significant improvements in antioxidant and anti-inflammatory properties caused by phytochemicals-based feedings have been extensively observed in many studies, and may be strongly associated with the Nrf2/Keap1 system.

2.2. Meat quality

A very large number of studies were performed to study the effects of phytochemicals on meat quality, with a focus on antioxidant properties, anti-inflammatory properties, and sensory performances such as color, texture, and flavor [106–133].

2.2.1. Antioxidant properties

The antioxidant properties of phytochemicals in farm animals, meat, and meat products have been extensively studied, forming a basis for an understanding of other functions of phytochemicals.

Phytochemicals supplementation was reported to improve the redox status and reduce excessive oxidative stress in pigs treated by peroxidation, by reducing plasma lipid peroxidation and lowering malondialdehyde (MDA) level. However, phytochemicals had no such effects in the case of non-pro-oxidative treatment [94,106,107]. Plant phytochemicals in the diet also moderately improved the antioxidant status in broilers and laying hens through the reduction of MDA and thiobarbituric acid reactive substance (TBARS) concen-

trations, and the induction of glutathione peroxidase (GPx) activity [96,98,102]. However, the antioxidant status was found to be less influenced in dairy cattle by phytochemicals supplementation, although the activity of superoxide dismutase (SOD) increased occasionally [104,105].

In meat and meat products, lipid oxidation is found to be the primary cause of quality loss. During the digestion–absorption–metabolism process, a number of oxidative compounds and stresses emerge and accumulate in the organism or tissues, adversely limiting the shelf-life and affecting the quality of the meat or meat products, including texture, color, flavor, nutritive value, and safety [136]. The toxic effects of synthetic antioxidants and consumers' interest in natural products have accelerated the development of natural phytochemicals as better choices than additives [137]. For example, the addition to pork patties of phytochemicals, such as extracts of grape skin, green tea, rosemary, and coffee, was observed to reduce lipid oxidation and the values of TBARS and hexanal at doses of 50–200 ppm [108]. In raw or cooked goat meat patties, extracts of red peony, white peony, moutan peony, rehmannia, sappan wood, and angelica were found to reduce lipid oxidation, at doses of 0.5%–2.0% [109]. In raw beef patties, ground beef, and buffalo meat patties, extracts of olive leaf, date pits, and rosemary leaf were found to reduce TBARS value, lipid oxidation, and oxymyoglobin oxidation, respectively [110–112]. In pork and beef sausages, adzuki bean extract and grape seed extract were found to reduce lipid oxidation and TBARS values [113,114]. The antioxidant properties of green tea extract, rosemary extract, and grape seed extract are well studied and their application in meat and meat products has been reviewed in a report [130].

2.2.2. Anti-inflammatory properties

Diets containing grape seed, marc extract, and hop extract were found to downregulate the expressions of various pro-inflammatory genes in the small intestine of growing pigs [95]. Cocoa powder in pig feed also decreased the gene expressions of Toll-like receptors and tumor necrosis factor (TNF)- α [117].

The anti-inflammatory effect of tea polyphenols on poultry was reported by investigating the expressions of a series of pro-inflammatory cytokines in the intestine of broilers. The results showed that tea polyphenols (0.03–0.09 g·kg⁻¹ body weight) caused a down-regulation of the gene expressions of TNF- α , interleukin (IL)-4, IL-10, IL-1 β , and interferon (IFN)- γ [118].

Feeding cattle with pomegranate-extract polyphenols (5–10 g·d⁻¹) increased the secretion of IL-4 and IFN- γ in peripheral blood mononuclear cells and improved the total immunoglobulin G (IgG) responses to the vaccination of ovalbumin [119]. Feeding dairy cows grape seed and marc extract stimulated a significant downregulation of the marker of endoplasmic reticulum stress, fibroblast growth factor (FGF)-21, as well as decreasing fat accumulation in the liver [104].

2.2.3. Sensory performance

Sensory performance is generally used to evaluate the color, flavor, and taste of meat or meat products. Phytochemicals have been found to affect the sensory performance of meat significantly.

For example, 0.5%–2.0% of white peony extract increased the redness value (a^* value) in raw and cooked meat patties [109]; 300–500 ppm of rosemary extract maintained the red color of raw frozen sausage [120]; 300 mg·kg⁻¹ meat of green tea extract decreased the a^* value in raw patties and eliminated rancid flavor in cooked patties [121,122]; 0.01%–0.02% of grape seed extract reduced visual green discoloration of beef patties [123]; 10% of myrtle extract prevented color changes in beef patties [124]; 2.7–10.8 mg·(100 g)⁻¹ of *Eleutherine americana* extract increased a^* value in cooked pork [125]; 0.2% of adzuki bean extract increased a^* value but decreased lightness (L^*

value) and yellowness (b^* value) in cured or uncured cooked pork sausages [126]; 500–6000 ppm of green tea extract increased a^* value whereas grape seed extract decreased a^* value in raw and cooked goat meat, and pepper extract was helpful in maintaining a^* value in cooked pork [127,128]; 5 mL·(500 g)⁻¹ meat of curry leaf extract decreased the L^* value and a^* value while increasing the b^* value in raw ground pork [129]; and 130 ppm of rosemary leaf extract stabilized the color in raw and cooked ground buffalo meat patties [130].

In addition, plum products exhibited minor effect on flavor but caused color change in many meat and poultry products, and grape seed extract led to a significant change in color in meat products [137].

2.3. Intestinal microbiota

Studies focusing on the effect of phytochemicals on the intestinal microbiota *in vivo* have increased markedly in recent years. It is considered that intestinal microbiota are the first targets of dietary phytochemicals, and that they show many links to health. Thus, many health-promoting effects of phytochemicals may be attributed to their modulation of the intestinal microbiota [138]. For example, only 5%–10% of polyphenols can be absorbed in the small intestine; 90%–95% enter the colon and are bio-transformed with the aid of the enzymatic colon microbiota into a series of polyphenolic metabolites [90]. The polyphenolic metabolites are able to partially re-absorb into the systematic circulation after conjugation once again in the liver and the enterocyte, and partially serve as antimicrobial substances or growth-promoting substrates. On the other hand, polyphenols or their metabolites can affect the composition and density of the colon microbiota in a profitable manner, including promotion of the growth of beneficial bacteria in a prebiotic-like manner and inhibition of certain pathogenic bacteria [139,140].

Limited studies were performed to specifically investigate the effects of polyphenols on the intestinal microbiota in farm animals. Cocoa powder feeding was found to increase the abundance of several bacterial strains in pigs, including *Lactobacillus*, *Bifidobacterium* spp., *Bacteroides-Prevotella*, and *Faecalibacterium prausnitzii* [117,134]. A few studies revealed that polyphenols may exhibit favorable effects in the intestine of broilers. Grape pomace concentrate supplementation in broiler feed was found to have a beneficial effect on the intestinal microbial population by increasing the abundance of *Enterococcus* and decreasing that of *Clostridium* [98]. Quercetin feeding in laying hens was reported to improve the caecal microflora status by decreasing the total number of aerobes and coliforms and increasing that of *Bifidobacterium* [101]. Tea polyphenols were found to increase the amount of lactobacilli and decrease that of the total bacteria, Bacteroidaceae, and *Clostridium* (*C.*) *perfringens* in pigs; however, they decreased *Bifidobacterium* spp., *Lactobacillus* spp., and *C. perfringens* in calves [102,135].

A recent review summarized the impact of polyphenols on the intestinal microbiota in rat and human models, and revealed that polyphenols or polyphenol-rich sources are able to affect the relative abundance of different bacterial groups by reducing the abundances of the potential pathogens *C. perfringens* and *C. histolyticum*, as well as that of Gram-negative *Bacteroides* spp., and by increasing the populations of certain beneficial strains, such as clostridia, bifidobacteria, and lactobacilli [108].

Based on the effects of phytochemicals on bacterial strains in several lines of studies, the antioxidant and anti-inflammatory properties of phytochemicals may be linked to improvements in gut health [141,142].

2.4. Detrimental effects of phytochemicals in farm animals

Although the biofunctional properties of phytochemicals are

powerful and promising for farm animals, detrimental effects of phytochemicals have also been reported in some cases. For example, high consumption of polyphenols can inhibit the absorption of nutrients [143,144] and cause toxic effects. Moreover, a high dose of quercetin was observed to be related to chronic nephropathy in rats and to reduce the life expectancy in mice [145]. Excess administration of green tea polyphenols was reported to disrupt kidney function in mice [146] and enhance tumor development in the colon of male rats [147]. Excess intake of caffeic acid caused kidney and gastrointestinal tumors in mice and rats [148]. Although these data were obtained from experimental animals, they suggest that the administration of a high dose of phytochemicals should be avoided in farm animals.

Although the anti-inflammatory, antioxidative, and cytoprotective properties of phytochemicals have been less studied in farm animals, an extremely large number of such studies have been done using human and experimental animal models. Thus, the biofunctions of phytochemicals in farm animals are also considered to occur through modulating the Nrf2/Keap1 system, which is a central modulator in combating oxidative stress and chronic inflammation [90].

3. Future perspectives

Several lines of studies have reported that the Nrf2/Keap1 system can regulate the general energy metabolism system by inhibiting gluconeogenesis [149]; modulate the activities of several lipases involved in the degradation of triglycerides/phospholipids [150] as well as enzymes involved in fatty acid oxidation, lipid biosynthesis, fatty acid desaturation, and fatty acid transport [150]; affect redox-sensitive metabolic systems such as the AMP-activated protein kinase pathway [151]; and adjust mitochondrial metabolism processes such as glucose oxidation and substrate entry, and ATP production [152,153]. EGCG has been shown to affect the general energy metabolism system in rats and in humans [154]. However, no data are available regarding farm animals, and the underlying molecular mechanism remains unclear. Thus, further studies are required to clarify the impact of phytochemicals on the energy metabolism system in farm animals.

Although extensive data have been accumulated on the biofunctions of phytochemicals in humans and in experimental animals, most of these data focus on the chemopreventive effects of phytochemicals on chronic diseases such as cancer, cardiovascular disease, and metabolic syndrome. The molecular data have been deeply mined to clarify how dietary phytochemicals modulate signaling pathways and gene expressions for homeostasis. On the other hand, the biofunctions of phytochemicals in farm animals have been paid a great deal of attention regarding growth performance, meat quality, and the use of phytochemicals as an antibiotic replacer or substituter, although the molecular data on mechanisms that have been obtained from farm animals are fewer than those obtained from humans and experimental animals. It appears to be difficult to compare the differences in the mechanisms of phytochemicals in farm animal nutrition and in human nutrition. However, the results from studies on the antioxidant properties and mechanisms of phytochemicals in farm animals are almost the same as the results of similar studies in humans, with the Nrf2/Keap1 system acting as an axis. Therefore, it is possible to take advantage of the phytochemical data from humans and experimental animals and apply them to farm animals.

Phytochemicals have multiple biofunctions for human and other animal health. The modulation of the Nrf2/Keap1 system by phytochemicals may play a central role in their multiple biofunctions because the Nrf2/Keap1 system is linked to antioxidant functions, anti-inflammation functions, and many other functions. The relatively low absorption ratio of most phytochemicals in the small intestine

shifts the research field from a focus on direct antioxidant properties to a focus on indirect pro-oxidant properties, biotransformation, signaling transduction, and gene expression regulation. Although the limited studies on the effects of phytochemicals on the intestinal microbiota of farm animals are currently insufficient to show the significant improvements in growth performance, antioxidant parameters, and inflammatory parameters, these findings will pave the way for further studies to understand the health-promoting effects of dietary phytochemicals.

Acknowledgements

This work was financially supported by funds from the Core Research Program 1515 of Hunan Agricultural University, the National Natural Science Foundation of China (31101268), and Scholar Research of Kagoshima University of Japan (for De-Xing Hou).

Compliance with ethics guidelines

Si Qin and De-Xing Hou declare that they have no conflict of interest or financial conflicts to disclose.

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