

Role of Phytochemicals in Skin Photoprotection *via* Regulation of Nrf2

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Ethnopharmacological studies have become increasingly valuable in the development of botanical products and their bioactive phytochemicals as novel and effective preventive and therapeutic strategies for various diseases including skin photoaging and photodamage-related problems abnormal skin including pigmentation and inflammation. Exploring the roles of phytochemicals in mitigating ultraviolet radiation (UVR)-induced skin damage is thus of importance to offer insights into medicinal and ethnopharmacological potential for development of novel and effective photoprotective agents. UVR plays a role in the skin premature aging (or photoaging) or impaired skin integrity and function through triggering various biological responses of skin cells including apoptosis, oxidative stress, DNA damage and inflammation. In addition, melanin produced by epidermal melanocytes play a protective role against UVR-induced skin damage and therefore hyperpigmentation mediated by UV irradiation could reflect a sign of defensive response of the skin to stress. However, alteration in melanin synthesis may be implicated in skin damage, particularly in individuals with fair skin. Oxidative stress induced by UVR contributes to the process of skin aging and inflammation through the activation of related signaling pathways such as the mitogen-activated protein kinase (MAPK)/activator protein-1 (AP-1), the phosphatidylinositol 3-kinase (PI3K)/protein kinase B (Akt), the nuclear factor kappa B (NF-kB) and the signal transducer and activator of transcription (STAT) in epidermal keratinocytes and dermal fibroblasts. ROS formation induced by UVR also plays a role in regulation of melanogenesis in melanocytes via modulating MAPK, PI3K/Akt and the melanocortin 1 receptor (MC1R)-microphthalmia-associated transcription factor (MITF) signaling cascades. Additionally, nuclear factor erythroid 2related factor 2 (Nrf2)-regulated antioxidant defenses can affect the major signaling pathways involved in regulation of photoaging, inflammation associated with skin barrier dysfunction and melanogenesis. This review thus highlights the roles of phytochemicals potentially acting as Nrf2 inducers in improving photoaging, inflammation and hyperpigmentation via regulation of cellular homeostasis involved in skin integrity and function. Taken together, understanding the role of phytochemicals targeting Nrf2 in photoprotection could provide an insight into potential development of natural products as a promising strategy to delay skin photoaging and improve skin conditions.

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INTRODUCTION

The skin is the largest organ in the body, and one of its main functions is to protect the body from environmental stressors including ultraviolet radiation (UVR), which can result in dermatological disorders, such as skin premature aging, abnormal pigmentation and inflammatory reactions. UVR has been generally known to have both beneficial and detrimental effects on human health. While UVR plays a role in natural synthesis of vitamin D, melanin, and various peptides in the skin that have positive biological implications (Slominski et al., 2000; Lin et al., 2016), excessive exposure to UVR can lead to acute and chronic adverse effects on the health of skin and is involved in the pathogenesis of premature aging (or photoaging) and increased risk of photocarcinogenesis. In addition, UVR is accepted as human carcinogen through oxidative mechanisms accountable for increased risk of developing skin cancers including nonmelanoma and melanoma skin cancers (Tran et al., 2008). The incidence of skin cancer has dramatically risen in particular among fair-skinned populations, primarily due to lifestyle changes and increased recreational exposure to UVR including outdoor activities and sunbathing for cosmetic purposes (Narayanan et al., 2010). Both male and female independent of age are affected by dermatological concerns and increasingly interested in rejuvenation and skin cancer prevention. While the use of sunscreens is recommended in order to minimize the risk of photoaging and other UV-related skin disorders, sunscreen alone does not provide sufficient protection against deleterious effects of UVR. Oxidative stress plays a crucial role in UVRinduced photodamage via mediating multiple biological responses including apoptosis, DNA damage, mitochondrial dysfunction, inflammation, abnormal pigmentation and upregulation of matrix metalloproteinases (MMPs) (such as MMP-1) in related skin cell types including keratinocytes, fibroblasts and melanocytes (Liebel et al., 2012; Denat et al., 2014; Silva et al., 2017; Lohakul et al., 2021b). Therefore, cellular and molecular regulation of antioxidant defenses to combat oxidative stress and promote redox balance could be a potential therapeutic and preventive strategies for photodamaged skin.

Nuclear factor erythroid 2-related factor 2 (Nrf2) is an important transcription factor controlling antioxidant responses in various tissues including the skin and plays a cytoprotection against chemical major role in and environmental insults including UVR (Ikehata and Yamamoto, 2018). Targeting Nrf2 could improve photoaging, wound repair and dyspigmentation as well as prevent photocarcinogenesis via regulation of cellular homeostasis involved in skin integrity and function (Saw et al., 2011; Gegotek and Skrzydlewska, 2015). Phytochemicals, which are ubiquitously present in plant-based diets and are active ingredients in several botanical drugs, have thus gained remarkable attention as promising candidates for effective photoprotective agents due to their abilities to activate Nrf2 signaling-regulated redox balance and subsequently maintain cellular homeostasis involved in skin integrity and function. Much attention has been focused on the role of dietary polyphenols in the repair of photodamaged skin and

prevention of solar-induced skin diseases (Surh and Na, 2008; Saw et al., 2011; Dunaway et al., 2018). In this review, we provide an overview of the promising roles of phytochemicals in mitigating UVR-induced skin damage via regulation of Nrf2mediated antioxidant response to offer an insight into ethnopharmacological potential for development of novel and effective anti-photodamaging agents.

The Role of UVR-Induced Oxidative Stress in Skin Photodamage

The skin is a primary target of oxidative stress because it is constantly exposed to environment including UVR, which induces reactive oxygen species (ROS) generation in the skin. It has been well accepted that both UVA and UVB rays play a significant role in the premature aging and photodamage of the skin through various mechanisms involving oxidative stress (Lephart, 2016; Gegotek et al., 2017). While UVB has biological impact on the skin primarily by causing direct damage to DNA and inflammation (Halliday and Lyons, 2008), UVA accounts for skin photodamage by generating various types of ROS, such as superoxide anion radical (O2[•]), singlet oxygen $({}^{1}\Delta gO_{2})$ and hydrogen peroxide (H₂O₂). ROS can interact with biomolecules and interfere with cell signaling, affecting cell survival and function of the skin cells (Dunaway et al., 2018). Considerable studies have reported that UVA exposure significantly led to ROS accumulation responsible for oxidative damage to biomolecules including DNA (O'Donovan et al., 2005), lipid (Dissemond et al., 2003) and protein in the skin cells including fibroblasts, keratinocytes (Brem et al., 2017) and melanocytes. Oxidative damage mediated by both UVA and UVB is associated with apoptosis and necrosis of the skin cells associated with sunburn reaction and photoaging process (Didier et al., 2001; Suschek et al., 2001; Kawachi et al., 2008; Parrado et al., 2016).

Skin Photoaging

Photoaging is characterized by epidermal thickness, termed hyperkeratosis, due to increased keratinocyte hyperproliferation as well as degradation or degeneration and disorganization of collagen fibers caused by upregulation of MMPs (Quan et al., 2009; Pittayapruek et al., 2016). In addition, dysregulated proliferation of transformed neoplastic keratinocytes or actinic keratosis is the key event in the progression from photoaged skin to squamous cell carcinoma (Berman and Cockerell, 2013). UVR (both UVA and UVB) is well accepted to play a vital role in photoaging via several mechanisms including DNA damage, oxidative stress, apoptosis, senescence, inflammation, immunomodulation (Rijken and Bruijnzeel-Koomen, 2011; Brand et al., 2017) and degradation and/or remodeling of the extracellular matrix (ECM) (Bosch et al., 2015). Generally, the characteristic hallmarks of photoaged skin are alterations in the ECM including accumulation of disorganized elastin fibers and depletion of collagens, the main structural proteins of the dermal connective tissues. Both UVA and UVB radiation can induce hyperkeratosis and several types of MMPs (including MMP-1 or collagenase) in mouse models of photoaging (Chaiprasongsuk et al., 2017; Misawa et al., 2017). Several in vitro

and in vivo studies have reported that UV irradiation stimulates expression of MMP-1, MMP-3 and MMP-9, which are the major UV-inducible collagenolytic enzymes, regulated at the transcriptional level (Afaq et al., 2009; Quan et al., 2009; Pittayapruek et al., 2016). MMPs are co-expressed in response to various stimuli including oxidative insults, inflammatory cytokines and growth factors (Greenlee et al., 2007; Lee et al., 2021). MMPs are suggested to be downstream targets within signaling pathways of upstream response genes, which encode several signaling proteins that activate different transcription factors capable of binding the promoters of MMP genes. The key transcription-binding sites involved in the regulation of MMP genes include the activator protein-1 (AP-1) site, the nuclear factor kappa B (NF- κ B) site and the signal transducer and activator of transcription (STAT) site (Fanjul-Fernandez et al., 2010). In addition, MMPs can be co-regulated because they share several transcription-binding sites in their promoter sequences. NF-KB and AP-1 are the transcription factors that can bind the promoters of MMP-1, 3 and 9 (Watanabe et al., 2004). The AP-1 transcription complex, a family of dimeric transcription factors composed of members of the Jun and Fos family proteins, is the main transcription factor regulating MMP-1 gene (Angel et al., 2001). In general, the c-Jun and c-Fos genes are activated rapidly and transiently in response to stimuli and are thus considered immediate-early response genes. Binding of heterodimer complexes of c-Jun with c-Fos to the AP-1 site, which is specific DNA sequences (5'-TGAG/CTCA-3'), termed TREs (TPA (tetradecanoylphorbol-12-Acetate)-response elements), is responsible for transactivation of AP-1 that regulates MMP-1, 3 and 9 expressions (Mackay et al., 1992; Watanabe et al., 2004). Both c-Jun and c-Fos are controlled by mitogenactivated protein kinase (MAPK) signaling pathways which are stimulated by extracellular stimuli including growth factors and cytokines as well as environmental stimuli including UVR. Three distinct types of MAPKs, ERK (extracellular signal-regulated kinase), JNK (c-Jun NH2-terminal kinase) and p38 MAPK, differentially affect AP-1 activity in response to various stimuli (Karin, 1995). The ERKs generally are triggered by growth factors and hormones as well as JNK and p38 MAPK are activated by environmental stresses including UVR and proinflammatory mediators, such as tumor necrosis factor (Chang and Karin, 2001; Silvers et al., 2003; Whitmarsh, 2007). UVR is suggested to primarily cause the greatest increases in JNK activity. Upon exposure to some stimuli, phosphorylation of c-Fos in the AP-1 complex at two C-terminal sites (Ser362 and Ser374) by MAPKs, in particular ERK, is required for transactivation at the specific AP-1 site (McBride and Nemer, 1998). In addition, c-Jun is activated and stabilized by JNK- and p38-catalyzed phosphorylation at the NH2-terminal sites (Ser63 and Ser73) located within transactivation domain of c-Jun. p38 MAPK indirectly activates AP-1 by phosphorylating other transcription factors such as AP-1 family proteins ATF2 (the activating transcription factor) forming a heterodimer with c-Jun, which then binds to the promoter elements in the c-Jun gene and regulates its transcription, leading to the subsequent upregulation of c-Jun expression and synthesis (Pramanik et al., 2003).

ROS participates in the photoaging process through several mechanisms including DNA damage, apoptosis, upstream modulation of MAPK/AP-1, NF-KB and JAK (janus kinase)-STAT signaling cascades, activation of cytokine and growth factor receptors and immune reaction of melanocytes and keratinocytes. Upregulation of MAPK/AP-1 signaling results in induction of transcription and production of MMPs (such as collagenase-1 (MMP-1), stromelysin-1 (MMP-3), and gelantinase A (MMP-2), that subsequently degrade ECM including collagen and elastin as well as suppress the collagen synthesis in the dermal fibroblasts. Furthermore, keratinocytes play an indirect role in photoaging through secreting paracrine factors, which stimulate the signaling cascades-mediated upregulation of MMPs in dermal fibroblasts. Several studies including ours suggested that ROS formation is involved in the molecular mechanisms of photoaging via activating MAPK/AP-1 signaling pathway, resulting in both upregulation of MMPs and downregulation of procollagen I production (Li et al., 2019). ROS induced by UVR is implicated in MAPKsdependent activation of AP-1 signaling, leading to upregulation of various MMPs including MMP-1, MMP-3 and MMP-9 in both keratinocytes and fibroblasts (Rittie and Fisher, 2002; Pittayapruek et al., 2016). Our in vitro and in vivo studies demonstrated the role of ROS induced by UVA exposure in upregulation of MMP-1 through activation of MAPK/AP-1 signaling pathway in keratinocytes and mouse skin. In addition, UVB has been shown to trigger MMP-1 and MMP-3 expressions through ROS generation and MAPK/AP-1 activation in irradiated keratinocytes and fibroblasts (Kim et al., 2013; Kim et al., 2015; Lu et al., 2016). In addition, UVB radiation is suggested to induce ROS formation, leading to increased MMP-9 activity and expression in mouse embryonic fibroblasts and HaCaT keratinocytes (Chang et al., 2017; Ma et al., 2018). UVB-induced oxidative stress was observed to activate MAPK signaling in association with increased expression of MMP-9 in UVB-exposed dermal fibroblasts (Gunaseelan et al., 2017). Moreover, ROS was found to be involved in activation of p38 MAPK and induction of MMP-9 expression in UVB-exposed HaCaT keratinocytes (Li et al., 2017b) and mouse dermis (Li et al., 2017b). In addition to collagen and elastin degradation by MMPs, UVB plays a role in a reduction of procollagen type I synthesis through activating AP-1-mediated downregulation of transforming growth factor beta (TGF-β) signaling (Pittayapruek et al., 2016; Gao et al., 2018a).

It should also be taken into account that photoaging of the skin is a complex multifactorial process. Apart from ROS/MAPK/AP-1 signaling cascades, UVR which can activate various cell surface receptors can stimulate downstream signaling pathways that control other different transcription factors regulating expression of many genes involved in photoaging process. Eventually, it is important to inhibit ECM degradation that leads to solar scar, a process taking place with each exposure to UV even at low doses. Since ROS serve as important second messengers, which act upstream of MAPK/AP-1 signalingmediated induction of MMPs and reduction of procollagen implicated in pathogenesis of photoaging, controlling ROS homeostasis could represent promising pharmacological and molecular approaches to impede photoaging.

Skin Inflammation and Skin Barrier Dysfunction

The skin barrier dysfunction and oxidative stress are suggested to play a role in the development of chronic inflammatory skin conditions (e.g., dermatitis) and alteration of wound healing process (Wikramanayake et al., 2014). The accumulation and alteration of external stimuli exposures result in a compromised barrier function of the skin through cutaneous inflammation and the imbalance of skin homeostasis (Egawa and Kabashima, 2018). Previous studies have demonstrated that UVB irradiation has a negative impact on epidermal morphology and barrier function by increasing stratum corneum (SC) thickness, causing changes in SC lipids and stimulating transepidermal water loss (Biniek et al., 2012). ROS generation induced by UVR plays a role in epidermal barrier dysfunction through oxidative damage to proteins and lipids, leading to alteration of tissue structure (Rinnerthaler et al., 2015). Regulation of Nrf2 activity has also been proposed to offer a potential strategy to improve skin barrier integrity by mitigating UVR-induced damage of keratinocytes and modulating inflammatory responses of the skin. Downregulation of Nrf2 signaling was shown to be involved in UVB-induced upregulation of pro-inflammatory mediators such as tumor necrosis factor alpha (TNF-a), cyclooxygenase-2 (COX-2), interleukin-6 (IL-6), interleukin-1 beta (IL-1β), and interleukin-8 (IL-8) in keratinocyte HaCaT cells (Park et al., 2021). A previous study using a mouse model of UVB-induced photodamage revealed that basal activity of Nrf2 in keratinocytes of normal skin is vital for improvement of skin barrier integrity and for prevention of skin carcinogenesis. UVB-mediated apoptosis of epidermal cells was involved in impaired skin integrity and activation of Nrf2 was observed to protect against UVB-induced apoptosis of basal keratinocytes in a paracrine, glutathione (GSH)/cysteine-dependent manner. Furthermore, enhanced levels of Nrf2-dependent genes in all layers of epidermis in response to UVB exposure were involved in the suppression of apoptosis in vivo (Schafer et al., 2010). The connection between the Nrf2 and antioxidant response element (ARE) system proved the protective pathways of skin inflammation via the regulation of the inflammatory factors (Saha et al., 2020). The NFE2L2 gene encoding for Nrf2 contains ARE-like sequences, providing a positive feedback mechanism to amplify antioxidant and anti-inflammatory signaling such as glutathione S-transferase (GST), NAD(P)H quinone oxidoreductase-1 (NQO-1), heme oxygenase-1 (HO-1) (Nguyen et al., 2009; Luo et al., 2018). Nrf2 has been suggested to play a role in modulating several signaling pathways involved in the inflammatory responses include NF-KB, MAPK, and JAK-STAT. Previous studies have reported the crosstalk between Nrf2 and NF-kB pathway. Nrf2 negatively regulated the NF-kB signaling pathway and proinflammatory cytokine production by inhibiting oxidative stress-induced NF-KB and preventing the IκB-α (NF-κB inhibitor) proteasomal degradation (Ma,

2013; Saha et al., 2020). In addition, several proinflammatory cytokines (e.g., IL-6, TNF-a and IL-1β), growth factors (e.g., epidermal growth factor (EGF), fibroblast growth factor, keratinocyte growth factor (KGF) and vascular endothelial growth factor (VEGF) and MMPs (e.g., MMP-2 and MMP-9) play a role in wound repair consisting of a series of multiple stages including inflammation, proliferation and remodeling (Shao et al., 2019; Suntar et al., 2021). Activation of Nrf2 in response to ROS production in inflamed tissues is thus suggested to play a role in promoting wound healing and regulating repair-related inflammation. Moreover, Nrf2 transcripts several genes encoding skin barrier structural and functional components including the keratins (KRT), the cornified envelope family members, small proline rich proteins, secretory leukocyte protease inhibitor, and the EGF family member epigen (Rojo de la Vega et al., 2017). In addition to Nrf2, STAT3 (in cell proliferation and differentiation), Smad proteins (in collagen production) and Forkhead box protein N1 (FOXN1) (in re-epithelization) are important transcriptional regulators involved in the process of wound repair. Furthermore, without involvement of inflammatory cells, upregulation of Nrf2 activity and its target antioxidant NQO-1 or HO-1 was demonstrated to promote the migration of corneal epithelial cells during wound repair in vitro and in vivo (Hayashi et al., 2013). In response to ROS produced in the early phase of wound repair, upregulation of Nrf2 as a target of KGF in keratinocytes is involved in the healing process in association with modulating proinflammatory cytokine IL-1, IL-6, and TNF- α and TGF- β 1 and VEGF *in vitro* and *in vivo* (Braun et al., 2002). Nevertheless, the regulatory role of Nrf2 in epidermal homeostasis is complex and needs further clarification as prolonged activation of Nrf2 in keratinocytes could interfere skin homeostasis. Previous in vitro and in vivo studies demonstrated that increased activity of Nrf2 in keratinocytes resulted in epithelial abnormalities, altered epidermal barrier and development of hyperkeratosis (Kypriotou et al., 2012; Schafer et al., 2012).

Skin Hyperpigmentation

While melanin plays a crucial role in protecting the skin against harmful effects of UVR, excessive production of melanin could be detrimental because melanin precursors and intermediate metabolites produced during melanogenesis in response to UVR exert phototoxic properties (Schmitz et al., 1995). Whereas hyperpigmentation mediated by UV irradiation could reflect a sign of defensive response of the skin to stress, alteration in melanin synthesis may be implicated in skin damage, particularly in individuals with fair skin. UVR-dependent elevated melanogenesis has been suggested to be biologically harmful, genotoxic and contributed to development of melanoma skin cancer, especially in lightly pigmented individuals. The incidence of skin cancer has dramatically risen in particular among fair-skinned populations, primarily due to lifestyle changes and increased recreational exposure to UVR including outdoor activities and sunbathing for cosmetic purposes (Narayanan et al., 2010; D'Orazio et al., 2013; Watson et al., 2016). Furthermore, the growth of skin fairness products is dramatic particularly in Asia and Africa, although the use of skin

bleaching products is associated with adverse side effects (Shroff et al., 2017). Thus, there is a need to develop effective and safe strategies for improvement of skin dyspigmentation or uneven complexion. Melanogenesis in melanocytes is a complex biosynthetic process involving the tyrosinase-catalyzed oxidation of tyrosine. Two main types of melanin, pheomelanin and eumelanin, are found in human skin and hair. Eumelanin is the brown/black insoluble pigment, characterizing dark phenotypes, and pheomelanin is the red/ yellow, sulfur-containing pigment, predominating in red-haired individuals (Slominski et al., 2004). Eumelanin functions as a UV absorbent and subsequently has photoprotective action. Pheomelanin is photolabile and can produce ROS as byproducts that lead to further DNA damage and is thus suggested to be carcinogenic following UVR (Brenner and Hearing, 2008).

Tyrosinase, a copper-containing membrane-bound located in melanosomes, catalyzes hydroxylation of L-tyrosine to L-DOPA, which is the first and the rate-limiting step of melanogenesis for both eumelanin and pheomelanin. In addition to tyrosinase, crucial enzymes involved in eumelanin synthesis include tyrosinase related proteins (TRP-1) and dopachrome tautomerase (DCT or TRP-2). Pheomelanin is produced via benzothiazine intermediates deriving from the oxidative polymerization of cysteinyl dopa derivatives generated through the condensation of the cysteine or GSH with the dopaquinone (Lu et al., 2021). Environmental stimuli (e.g., UVR and drugs), endogenous factors (e.g., hormone and mediators) and genetic factors can influence melanogenesis regulated by tyrosinase via various signal pathways, primarily the melanocortin 1 receptor (MC1R)-microphthalmia-associated transcription factor (MITF) signaling. MC1R is a G protein-coupled receptor that controls the quantity and quality of melanin synthesized in melanocytes. Important agonists of MC1-R acting as the main intrinsic regulator of pigmentation are peptide hormones and neuropeptides including stimulating hormone (a-MSH), endothelin-1 (ET-1) and adrenocorticotropic hormone (ACTH), which are cleavage products of proopiomelanocortin (POMC) (Lin and Fisher, 2007). The major signal transduction pathways that mediate the regulation of melanogenesis involve the binding of agonists to MC1R that trigger events inside melanocytes through raising intracellular cyclic 3'-5'-cyclic adenosine monophosphate (cAMP) and activating the adenylate cyclase enzyme, protein kinase A (PKA), leading to phosphorylation of the cAMP responsive binding element (CREB), which promotes the activation of MITF, which is the master transcription factor that regulates expression of several melanogenic genes including tyrosinase, TYRP1 and TYRP2. Moreover, upon activation of MC1R, enhanced levels of cAMP and subsequent activation of PKA were observed to activate the MAPK signaling cascades including p38, leading to activation of MITF (Smalley and Eisen, 2000). However, inhibition of MC1R in normal melanocytes and melanoma cells was observed to trigger PI3K/Akt and MAPK/ERK pathways, leading to inhibition of MITF and subsequent suppression of melanogenesis (Chae et al., 2017; Wu et al., 2018). Mechanisms underlying the role of phytochemicals in

regulating pigmentation involve the direct suppression of tyrosinase activity and/or gene expression, direct scavenging of ROS, promotion of Nrf2-regulated antioxidant defense and inhibition of signaling pathways involved in inflammatory responses (Vomund et al., 2017; Boo, 2019). Nrf2 is suggested to play a role in modulating crucial signaling pathways including MAPK, PI3K/Akt and MC1R-MITF signaling cascades involved in regulation of melanin synthesis (Shin et al., 2014; Chaiprasongsuk et al., 2016). Moreover, exposure of the skin to UVR can stimulate keratinocytes to secrete hormones including ACTH, ET-1, α-MSH that bind to MC1R, activating and upregulating melanogenesis-related proteins. MITF Activation of Nrf2 has been observed to suppress the paracrine factors (such as α-MSH) derived from keratinocytes that results in downregulation of signaling pathways (including the cAMP/CREB/MITF pathway) involved in melanogenesis in melanocytes (Hseu et al., 2020; Chen SJ. et al., 2021). Therefore, application of compounds having abilities to activate Nrf2 might represent a promising approach to prevent and treat hyperpigmentation disorders.

Moreover, in response to UVR, melanogenesis acts as a shield against the harmful effect of UVR on the skin and thus approaches promoting melanin production can mitigate UVRinduced melanocyte damage. We previously demonstrated the role of Nrf2 in regulating the release of paracrine factor α -MSH by keratinocytes that influenced UVB-mediated melanocyte responses including DNA damage, oxidative stress, apoptosis and inflammation (Jeaveng et al., 2017). Several natural compounds such as flavonoids and coumarins having abilities to induce melanogenesis and restore melanocyte viability might thus be useful in the prevention and treatment of hypopigmentation disorders such as vitiligo (Niu and Aisa, 2017). Therefore, phytochemicals have been proposed to exert beneficial effects against abnormal melanogenesis via improving hyperpigmentation or hypopigmentation caused by disruption of melanocyte homeostasis and/or loss of functional melanocytes. While this review highlights the studies demonstrating the roles phytochemicals improving UVR-induced of in hyperpigmentation via Nrf2-dependent mechanisms, it should be taken into account that melanocyte biology is complex and the role of phytochemicals in regulating melanogenesis involved in maintaining the skin homeostasis needs further clarification.

The Role of Nrf2-Regulated Antioxidant Defense Against Cutaneous Photodamage

The primary endogenous antioxidant defenses include antioxidant and detoxification enzymes such as catalase, glutamate cysteine ligase (GCL) (composed of a catalytic subunit GCLC and a modifier subunit GCLM), the ratelimiting enzyme in GSH synthesis, glutathione peroxidase (GPx), GST, HO-1, NQO-1 and superoxide dismutase (SOD) regulated by various transcription factors including Nrf2 (Surh, 2003; Hseu et al., 2012). Nrf2 is a master regulator of antioxidant and cytoprotective genes involved in the human skin adaption to the environmental insults including UVR and thus plays a beneficial role in maintenance of skin homeostasis. Activity of Nrf2 is tightly regulated by proteins including Kelch-ECH associated protein 1 (Keap 1) and proteasome degradation system and thus regulation of Nrf2-mediated antioxidant response pathway is complicated. Under homeostatic conditions, two molecules of Keap1 bound to Nrf2 is responsible for the continuous ubiquitylation and degradation of Nrf2. In response to Nrf2 activating stimuli or oxidative stress, Keap1 is oxidized at critical cysteine residues, especially Cys151, leading to dissociation of Keap1-Nrf2 that allows Nrf2 to escape from Keap1-mediated ubiquitination. Nrf2 is then translocated into the nucleus and binds to the ARE promoter, a *cis*-acting enhancer sequence located in the 5'-flanking regions of genes encoding phase II and antioxidant cytoprotective enzymes including GST, NQO-1 and GCL (Schafer et al., 2010; Liu et al., 2016; Boo, 2020b).

Nrf2 plays a vital role in maintaining redox homeostasis and cellular metabolism in skin cells involved in the skin's structural integrity and function (Ikehata and Yamamoto, 2018). Oxidative insults, such as UVR and H₂O₂, and electrophilic chemicals, such as butylated hydroxyanisole and its active de-methylated metabolite *tert*-butyl hydroquinone (tBHQ); phenolic flavonoids [e.g., green tea polyphenols and epigallocatechin-3gallate (EGCG)]; and the naturally occurring isothiocyanates including sulforaphane (SFN) and curcumin, can stimulate Nrf2 activity via modification of Keap1 cysteine residues, suggested as the stress sensors for Nrf2 activator (Baird and Yamamoto, 2020). The cysteine modifications result in a conformational change in the associated motif of Keap1-Nrf2 that facilitates the dissociation of Nrf2 from Keap1 and subsequently Nrf2 nuclear translocation (Kong et al., 2001). Generally, various environmental stressors including UVR lead to post-translational activation of Nrf2 through Keap1 inactivation. The upregulation of Nrf2-mediated antioxidant defense system was demonstrated in vitro and in vivo to protect the human skin from harmful effects of UVR. UVA-1mediated lipid oxidation induces expression of antioxidant response genes, which is dependent on the redox-regulated transcription factor Nrf2 in dermal fibroblasts (Gruber et al., 2010). Exposure of keratinocytes (including primary human epidermal keratinocytes and HaCaT keratinocyte cell lines) to UVA (20 J/cm²) increased Nrf2 activity via enhancing Keap1 expression. UVA exposure led to stimulation of Nrf2 activity and its target proteins (HO-1, NQO-1, GST) in HaCaT keratinocytes and dermal fibroblasts, although Nrf2 activity was minimally affected in UVA-irradiated primary keratinocytes (Rysava et al., 2020). In fact, the regulatory role of Nrf2 in skin cell survival and function affected by UVR is complex because UVR can either upregulate or downregulate Nrf2-mediated antioxidant defense in various skin cell types. Changes in the Nrf2 activity are dynamic and dependent on types of UV ray, UVR's intensity and time following the exposure (Chaiprasongsuk et al., 2016; Rysava et al., 2020; Rysava et al., 2021). Previous observations indicate that both UVA and UVB downregulate Nrf2 antioxidant signaling pathway in skin keratinocytes, fibroblasts and melanocytes in vitro and in skin tissues in vivo. UVB exposure led to reduced expressions of Nrf2 and its target antioxidant HO-1 proteins in HaCaT keratinocyte cells and mouse skin in vivo



number of phenolic rings they contain and of the structural elements binding the rings to one another. They are generally divided into four classes including phenolic acids, flavonoids, stilbenes and lignans. Phenolic acids are further classified as hydroxyl benzoic and hydroxyl cinnamic acid derivatives (Johnsson, 2004; Nishiumi et al., 2011).

(Rodriguez-Luna et al., 2018; Rodriguez-Luna et al., 2019). UVR was observed to downregulate antioxidant and detoxifying enzymes including GST, NQO-1 and y-GCS (νglutamylcysteine synthetase) in the skin cells through modulating activity of Nrf2 (Kannan and Jaiswal, 2006; Lohakul et al., 2021b). The DNA damage or modulation of signaling cascades (including MAPKs) that take place rapidly in response to UVR exposure is suggested to mediate the downregulation of Nrf2 antioxidant response pathway (Lopez-Camarillo et al., 2012). The p38 was suggested to reduce Nrf2 nuclear translocation and its transcriptional activity (Boo, 2020b). In addition, activation of Nrf2 signaling has been suggested to protect against UVR-mediated skin damage via several mechanisms including promotion of antioxidant and cytoprotective defense, DNA repair, anti-inflammatory signaling. Upregulation of Nrf2/HO-1 signaling accompanied with increased activities and protein levels of catalase, GPx and SOD was observed to suppress apoptosis induced by UVR [UVA (3 J/cm²)+UVB (90 mJ/cm²)] via activating the PI3K/Akt signaling pathway in the 3D skin model (Xian et al., 2019). Thus, understanding the role of Nrf2 in the pathogenesis of skin photodamage could give an insight into development of potential compounds having Nrf2 inducing activity for prevention and treatment of skin photodamage.

Phytochemicals Targeting Nrf2 in Skin Photodamage: Development of Botanicals and Phytochemicals as Promising Photoprotective Agents

Several reports have highlighted the potential role of bioactive phytochemicals of plant-based diets and botanical drugs that have been used in ethnomedicine or reported in



ethnopharmacological studies. The bioactive phytochemicals are naturally occurring compounds in botanical products including plants and botanical drugs which exert biological activities providing medical and nutritional benefit. These compounds exert antioxidant effects by directly scavenging ROS or by promoting the antioxidant defense system through the Nrf2dependent pathway (L Suraweera et al., 2020). Phytochemicals include polyphenols and the non-phenolic phytochemicals. Phenolic compounds are classified into different groups on the basis of the number of phenolic rings they contain and of the structural elements binding the rings to one another. They are generally divided into four classes including phenolic acids, flavonoids, stilbenes and lignans (Figure 1). Phenolic acids are further classified as hydroxyl benzoic and hydroxyl cinnamic acid derivatives (Pandey and Rizvi, 2009). The natural sources of polyphenols include fruits (e.g., apple, berries, cherries, grapes, strawberries and pomegranate), vegetables, soybeans, cereals, tea, cocoa, soy and Phlebodium aureum (L.) J.Sm. The common flavonoids include catechins, quercetin, genistein, epicatechin,

catechin and anthocyanins (Bosch et al., 2015). The common phenolic acids are hydroxycinnamic acids including caffeic acid and ferulic acid as well as gallic acid (also known as 3,4,5trihydroxybenzoic acid). Gallic acid is the most abundant phenolic acid found in plant-based diets (Hano and Tungmunnithum, 2020). For non-flavonoid phenolics, the most widely studied stilbene is resveratrol. The commonly studied non-phenolic phytochemicals include carotenoids, caffeine and sulforaphane (SFN) (Bosch et al., 2015).

Phytochemicals play a crucial role in photoprotection against UVR-induced skin photodamage *via* UV-absorbing, antioxidant, melanin-modulating, anti-inflammatory properties Antioxidant phytochemicals have been demonstrated to mitigate skin photodamage *in vitro* and *in vivo via* directly scavenging ROS, promoting antioxidant defense capacity, modulating various signaling pathways involved in inflammation, controlling DNA repair, cellular viability and function of the skin (Boo, 2020a; Garg et al., 2020). This review focuses the photoprotective role of phytochemicals in UVR-mediated photoaging,

hyperpigmentation and inflammation affecting skin barrier integrity via Nrf2-dependent pathway (Figure 2). The phytochemicals as electrophiles can promote cytoprotective proteins and antioxidant defenses via upregulating Nrf2 signaling. Keap1 and Cul3 comprise a unique ubiquitin E3 ligase responsible for degradation of Nrf2. Keap1 is a homodimeric protein belonging to the BTB (Broad complex, Tramtrack, Bric-á-brac)-Kelch family of proteins, which are named Kelch-like 1 to 42 (KLHL1-42). The BTB domain of Keap1 is necessary for Keap1 homodimerization and for mediating interactions with cul3/Rbx1 E3 ubiquitin ligase system. The BTB domain contains reactive cysteine residue responsible for interaction with electrophiles and thus plays a crucial role in sensing environmental electrophiles. Posttranslational modifications of the highly reactive Cys 151 in Keap1 result in dimerization of Keap1, resulting in loss of Nrf2 ubiquitination, which stabilize the Nrf2 proteasomal degradation (Shin et al., 2020) and subsequent accumulation of Nrf2 and activation of the Nrf2-driven cytoprotective gene machinery (Cleasby et al., 2014). Phytochemicals, which are thiolreactive electrophiles, covalently bind to the cysteine residue(s) in the dimerization domain of Keap1. Then, the activated ligase complex fails to degrade Nrf2, allowing the transcriptional activation of Nrf2 target genes (Yamamoto et al., 2008). Wellknown Nrf2 activators including the isothiocyanate SFN, the alkylating agent iodoacetamide, tBHQ and diethylmaleate were demonstrated to modify C151 in Keap1, that mediates proteasomal degradation, leading to Nrf2 stabilization and enhancing its nuclear accumulation (Deshmukh et al., 2017; Dayalan Naidu et al., 2018; Dayalan Naidu and Dinkova-Kostova, 2020; Taguchi and Yamamoto, 2020).

In addition to the role of phytochemicals as indirect antioxidants upregulating antioxidant by genes, phytochemicals can act as direct antioxidants by their ROS scavenging activities (Dinkova-Kostova and Talalay, 2008). Direct antioxidants, such as (-)-epicatechin-3-gallate and carotenoids (i.e., β-carotene and lycopene) can protect skin cells from ROS-induced damage to the skin cells. Their protective effects are short-lived and involve their abilities to neutralize ROS. ROS is involved in regulating the activity of Nrf2 via several mechanisms. Keap1 is considered as a cysteinebased sensor for a variety of endogenous and exogenous stressors including electrophiles and oxidants. ROS is involved in modulate Nrf2 activity via Keap-1-dependent and independent mechanisms. For Keap1-dependent mechanism, ROS (e.g., H₂O₂) has been demonstrated to promote Nrf2 activity via oxidative modification of Keap1 cysteines, leading to the release of Nrf2 and allowing its nuclear translocation (Espinosa-Diez et al., 2015; Suzuki et al., 2019). For Keap1independent regulation of Nrf2 activity, MAPKs and glycogen synthase kinase-3 (GSK-3) are suggested to play a role in posttranslational modifications of Nrf2 via phosphorylation accountable for the alterations in its binding to the proteins involved in controlling Nrf2 stability and subsequent transcriptional activity (He et al., 2020). For instance, GSK-3 activation can stimulate Nrf2 nuclear export as well as ubiquitination and degradation, leading to downregulation of the Nrf2/ARE signaling pathway of brain ischemia and reperfusion injury (Chen et al., 2016). In addition, suppression of GSK-3 by activation of upstream phosphoinositide 3-kinase-protein kinase B/Akt (PI3K-PKB/Akt) results in Nrf2 stabilization. Activation of ERK leads to Nrf2 downregulation in diabetic hearts in response to oxidative stress (Tan et al., 2011). Thus, it is possible that ROS which can act as second messengers in protein kinase cascades also have a regulatory role in Nrf2 activity *via* Keap1-independent manner. Phytochemical polyphenols that act as both pro-oxidants through autoxidation to generate ROS (Babich et al., 2011) and direct antioxidants that can increase and decrease cellar ROS levels can affect Nrf2 activity *via* both Keap1-dependent and -independent manners.

In fact, skin is the largest body organ that is continuously exposed to environmental stressors including UVR. Oxidative stress induced by UVR plays a role in the stress responses of keratinocytes, melanocytes and fibroblasts responsible for photodamaged skin. Hence, development of phytochemicals that can activate the Nrf2 transcription factor is considered a promising pharmacological strategy to prevent and treat UVRinduced skin damage. The phytochemical derivatives that are effective for these photoprotective strategies include polyphenols, flavonoids, non-flavonoids and non-phenolic derivatives. The phytochemicals having ROS scavenging properties could suppress UVB-induced MMP-1 expression in HaCaT cells and human dermal fibroblasts and promote type I procollagen production in human dermal fibroblasts via downregulation of MAPK/AP-1 signaling cascades in association with upregulation of Nrf2 signaling (Kim et al., 2015). Moreover, botanical extracts including extracts of sunflower (Helianthus annuus L.) (Hwang et al., 2019), cherry blossoms (Li et al., 2018) and Foeniculum vulgare Mill. (Sun et al., 2016), which are rich sources of antioxidant phytochemicals, were observed to exert the protective effect against UVB-induced ROS formation, MMPs (MMP-1 and MMP-3) production and procollagen type I depletion via downregulation of MAPK signaling in association with upregulation of Nrf2 activity in human dermal fibroblasts. The anti-photoaging actions of the sunflower extract were also related to suppression of UVBinduced inflammatory cytokines including IL-6, COX-2, iNOS (inducible nitric oxide synthase), and TNF-a production (Hwang et al., 2019). In addition, the phytochemicals acting as direct or indirect Nrf2 inducers were demonstrated to exert the antiphotoaging effects via downregulation of MMPs including MMP-1 via MAPK/AP-1 signaling pathways in mouse skin (Sun et al., 2016; Li et al., 2018). Thus, indirect or direct targeting of Nrf2-dependent antioxidant response could offer a promising pharmacological strategy for prevention and inhibition of skin photodamage. The crosstalk between Nrf2 and other signaling pathways (e.g., MAPK/AP-1 pathway) involved in the mechanisms underlying the protective effects of phytochemicals on photodamaged skin were shown in Figure 3 (Kim et al., 2015; Sun et al., 2016; Li et al., 2018; Hwang et al., 2019; Garg et al., 2020). Several in vitro, in vivo and clinical studies showing the protective roles of botanicals and phytochemicals against photoaging, inflammation, skin barrier



FIGURE 3 The regulatory mechanism of Nrf2-dependent antioxidant and cytoprotective actions against photoaging, inflammation and hyperpigmentation. Phytochemicals play a photoprotective role against skin photodamage directly *via* scavenging reactive oxygen species (ROS) and indirectly via activation of nuclear factor erythroid 2-related factor 2 (Nrf2) signaling, leading to upregulation of antioxidant and cytoprotective genes. In response to ultraviolet ray (UVR)-induced oxidative stress, ROS, produced in keratinocytes, melanocytes and fibroblasts, can modulate several related signaling pathways involved in photoaging, inflammation and melanogenesis. Exposure of human skin to UVR causes cells to produce ROS, which can modulate the signaling pathways involved in upregulation of matrix metalloproteinases (MMPs) which includes MMP-1/3/9, leading to collagen degradation, inflammation (nuclear factor kappa B also known as NF-xB and its downstream signaling) and the upregulation of melanogenesis-related genes including the microphthalmia-associated transcription factor (MITF), tyrosinase (TYR), and tyrosinase related proteins (TRP-1). Dietary phytochemicals as natural sources of antioxidants play the protective roles against UVR-induced ROS by the inhibition of ROS formation and the activation of Nrf2 signaling. In response to oxidative insults, Nrf2 is activated by the phosphorylation and disassociation of Nrf2 from Kelch-ECH associated antioxidant and phase II detoxifying enzymes including glutamate cysteine ligase (GCL), glutathione S-transferase (GST), NAD(P)H quinone oxidoreductase-1 (NQO-1), heme oxygenase-1 (HO-1). The activation of Nrf2 improves oxidative status of the cells and promotes cytoprotection against skin oxidative damage and inflammation.

dysfunction and hyperpigmentation *via* Nrf2-dependent mechanisms are described below and summarized in **Table 1**.

Isothiocyanates

Isothiocynates are sulfur-containing compounds having the general formula R–N=C=S and are commonly found in cruciferous vegetables from the *Brassica* genus including broccoli, brussels sprouts, cabbage and cauliflower. Members of isothiocynates widely known as Nrf2 activators including SFN and phenethyl isothiocyanate (PEITC) protected human *ex vivo* full skin against UVR-induced sunburn cells, apoptosis and the decreased activity of the antioxidant enzyme catalase in correlation to upregulation of Nrf2 activity and its target genes (γ -GCS, HO-1, NQO-1) in HaCaT keratinocytes (Kleszczynski et al., 2013). SFN (0.6 μ M/cm²) was observed to exert antiphotoaging effects on mouse skin via inhibition of MAPK/AP-1 signaling in UVA-irradiated mouse skin (Chaiprasongsuk et al., 2017). Treatment of keratinocyte cell line NCTC2544 with SFN (10 μ M) combined with the Fernblock[®] XP (1 and 2 mg/ml),

obtained from the tropical fern *Phlebodium aureum* (L.) J.Sm., substantially suppressed the production of MMP-1, MMP-3 and IL-1 in association with a decrease in ROS production. The SFN (5 and 10 μ M) and Fernblock[®] XP (1 mg/ml) combination also showed inhibitory effects on melanoma cell growth and migration *in vitro* in association with the ability to inhibit the inflammatory microenvironment and neo-angiogenesis (Serini et al., 2020).

Benzyl isothiocyanate and 6-(Methylsulfinyl)hexyl isothiocyanate derived from Wasabi, have been reported to suppress a transcriptional levels of COX-2, an enzyme synthesizing the pro-inflammatory mediators (Lee et al., 2009; Uto et al., 2012). The disturbance of phosphorylated MAPKs signaling, ERK, p38 kinase, and JNK, was observed in the ITCs treatment, resulting in the downregulation of the transcription of inflammatory genes such as COX-2, iNOS, TNF- α , IL-1 β , and IL-6 (Lee et al., 2009; Latronico et al., 2021). The effects of allylisothiocyanate and SFN on the Nrf2 nuclear translocation were associated with the downregulation of p65 protein, a subunit of

Phytochemicals	Active compounds and sources	Effects	Treatment and study model	Mechanism of action	References
Isothiocyanates	Sulforaphane (SFN), broccoli, brussels sprouts, cabbage and cauliflower	Anti-photoaging	Topical administration of SFN (0.6 µM/cm ²) for 2 weeks	J: MMP-1, 8-OHdG DNA damage, MAPKs signaling, c-Jun, and c-Fos	Chaiprasongsuk et al. (2017)
			In vivo: BALB/c mice	↑: Nrf2 translocation, Nrf2-target	
	SFN, broccoli, brussels sprouts, wasabi	Anti-inflammation and improvement of skin barrier function	Topical administration of SFN (2.5–10 mg/kg)	genes, collagen J: Janus kinase 1/STAT3 signaling, skin thickness and eosinophil accumulation in atopic dermatitis	Wu et al. (2019)
			In vivo: BALB/c mice	mouse skin lesions [†] : Nrf2 and Nrf2-dependent antioxidant enzymes (HO-1)	
	SFN	Hyper-pigmentation	B16F10 cells treated with 10 μM of SFN for 6 h	1: Tyrosinase activity, melanin content, ROS and 8-OHdG ↑: Nrf2 and Nrf2-dependent antioxidant enzymes (GCL, GST, NQO-1)	Chaiprasongsuk et al. (2016)
Flavonoids	Flavanones (Hesperetin; HSP)	Anti-photoaging	Topical administration of HSP (0.3, 1, and 3 mg/cm ²) for 2 weeks	↓: MMP-1, 8-OHdG DNA damage, MAPKs signaling, c-Jun, and c-Fos	Lohakul et al. (2021a)
			In vivo: BALB/c mice	↑: Nrf2 translocation, Nrf2-target	
	Fisetin	Anti-inflammation and improvement of skin barrier function	Topical administration of Fisetin (25–100 μΜ) for 10 weeks	genes, collagen J: pro-inflammatory mediators (COX-2, IL-6, and NF-κB), aquaporin and filaggrin (the protein	Wu et al. (2017)
				markers of skin barrier function)	
	Caffeic acid, ferulic acid, quercetin and rutin	Hyper-pigmentation	In vivo: BALB/c mice Treatment of cells with of caffeic acid, ferulic acid, quercetin and rutin prior to	î: Nrf2 ↓: Melanogenesis	Chaiprasongsuk et al. (2016)
			UVA irradiation In vitro: B16F10 melanoma cells	↑: Nrf2 and Nrf2 targeted genes	
	Ellagic acid		Treatment with ellagic acid (20-80 µM) for 24-72 h	J: protein levels of the paracrine factors, proopiomelanocortin (POMC), α-MSH, and AKT/JNK/ ERK signaling	Yang et al. (2021)
			<i>In vitro</i> : keratinocyte HaCaT cells	↑: Nrf2 nuclear protein	
Carotenoids	Rosemary extract (carnosic acid)	Anti-photoaging	Pre-treatment with rosemary extract containing carnosic acid (2.5–10 µM) for 6–9 h	↓: matrix metalloproteinases (MMPs)	Calniquer et al. (2021)
			In vitro: HaCaT keratinocytes and KERTr keratinocytes	↑: Nrf2/ARE systems	
	Crystalline lycopene preparations purified from	Anti-inflammation and improvement of skin barrier function	Pre-treatment with the different compounds at a	\downarrow : NF-κB activity and IL-6	Calniquer et al. (2021)
	tomato extract (>97%), carotenoid-rich Tomato Nutrient Complex (TNC), rosemary extract		concentration of 5 μM <i>In vitro</i> : HaCaT keratinocytes	↑: ARE/Nrf2 activity	
	Fucoxanthin		topical application of cream containing the fucoxanthin (0.2% w/w) to mouse skin	↓: melanin index and skin edema, COX and IL-6	Rodriguez-Luna et al. (2018)
			In vivo: Female Swiss CD-1 mice	↑: Nrf2-dependent antioxidant enzymes; heme oxygenase-1 (HO-1)	

TABLE 1 | The protective roles of botanicals and phytochemicals against photoaging, inflammation, skin barrier dysfunction and hyperpigmentation via Nrf2 regulation.

(Continued on following page)

TABLE 1 (Continued) The protective roles of botanicals and phytochemicals against photoaging, inflammation, skin barrier dysfunction and hyperpigmentation via Nrf2 regulation.

Phytochemicals	Active compounds and sources	Effects	Treatment and study model	Mechanism of action	References
Resveratrol and stilbenoid	Grape peel extract, dried heartwood of <i>Pterocarpus</i> <i>marsupium</i> Roxb.	Anti-photoaging	Oral administration of either 2 g GPE or 2 mg resveratrol per kg body weight in mice <i>In vivo</i> : mice	 J: skin wrinkle formation î: Nrf2-dependent antioxidant enzymes; heme oxygenase-1 	Kim et al. (2019)
	Resveratrol	Hyper-pigmentation	Treatment of UVB- irradiated skin with	(HO-1) ↓: MITF and its target proteins including TYR, TRP1, TRP2	Lee et al. (2014)
	Pterostilbene extracted from the dried heartwood of <i>Pterocarpus marsupium</i> Roxb.		resveratrol In vivo: Guinea Pig Skin Treatment with pterostilbene In vitro: B16F10 mouse melanoma cells	 ↑: Nrf2/HO-1 proteins ↓: melanogenesis and tyrosinase activity ↑: Nrf2-mediated HO-1, γ-GCLC, and NQO-1 protein expressions 	Li et al. (2017a), Majeed et al. (2020)
	Pterostilbene (Pter)	Anti-inflammation and improvement of skin barrier function	Keratinocyte HaCaT cells Pre-treatment with Pter (5 and 10 µM) for 24 h prior to UVB irradiation (300 mJ/cm ²) <i>In vitro</i> : HaCaT keratinocytes	 J: ROS generation 1: Nuclear translocation and phosphorylation of Nrf2, expression of Nrf2-dependent antioxidant enzymes, DNA repair activity, phosphatidylinositol-3-kinase (PI3K) phosphorylated kinase, Akt 	Li et al. (2017a)
Sesquiterpene lactones	Santamarine isolated from Asteraceae scoparia, artichoke (<i>Cynara</i> <i>scolymus</i> L)	Anti-photoaging	Pre-treatment with Santamarine (1–10 µM) for 1 h <i>In vitro</i> : human dermal fibroblasts	J: ROS levels, MAPKs/AP-1, and MMP-1/3/9, p-c-Fos and p-c-Jun f: collagen I, TGF-β/Smad signaling, Nrf2-dependent intracellular antioxidant mechanism (SOD and	Oh et al. (2021)
	Cynaropicrin (Cyn)	Anti-inflammation and improvement of skin barrier function	Pre-treatment with Cyn (up to 100 µM) prior to UVB irradiation (50 mJ/cm ²) <i>In vitro</i> : Normal human epidermal keratinocyte (NHEKs)	HO-1) ↓: ROS generation, TNF-α, BaP ↑: Nrf2, Nrf2-dependent antioxidant enzymes (NQO-1)	Takei et al. (2015)
Tannins	Red raspberries (<i>Rubus idaeus</i> L.) extracts	Anti-photoaging	Pre-treatment with <i>Rubus idaeus</i> L. (1–100 µM) for 1 h <i>In vitro</i> : normal human	J: MAPK/AP-1, NF-κβ and TGF-β/ Smad î: type I procollagen and Nrf2	Gao et al. (2018b)
	<i>Alchemilla mollis</i> (Buser) Rothm. (AM) extract		dermal fibroblasts (NHDFs) Treatment with AM (1, 10 and 100 μ g/ml) for 4 h after UVB irradiation (144 mJ/cm ²)	nuclear transfer]: ROS production, TGF-β1, MMP- 1, IL-6, and nucleus NFATc1 dephosphorylation, wrinkle formation, skin thickening, water loss, and erythema	Hwang et al. (2018)
			<i>In vitro:</i> NHDFs <i>In vivo:</i> hairless mice	↑: type I procollagen and elastin expression, Nrf2-dependent antioxidant enzymes (NQO-1 and HO-1)	
	Red raspberries extract (RBE)	Anti-inflammation and improvement of skin barrier function	Pre-treatment with RBE (62.5–1,000 µg/ml) for 48 h prior to UVB irradiation (100 mJ/cm ²)]: Cell viability, epidermal thickness	Wang et al. (2019)
			Topical treatment with RBE (750 µg/ml) <i>In vitro</i> : HaCaT cells <i>In vivo</i> : nude mice (ICR-	↑: Nrf2, Nrf2-dependent antioxidant enzymes (catalase, SOD, NQO-1, and HO-1)	

TABLE 1 (Continued) The protective roles of botanicals and phytochemicals against photoaging, inflammation, skin barrier dysfunction and hyperpigmentation via Nrf2
regulation.

Phytochemicals	Active compounds and sources	Effects	Treatment and study model	Mechanism of action	References
Terpenoids: diterpene, triterpene and sesquiterpene	Ginsenosides compound Mx (C-Mx) from Notoginseng stem-leaf	Anti-photoaging	Pre-treatment with the Ginsenosides C-Mx (1–20 uM) for 3–72 h	↓: MMP-1 and 3	Liu et al. (2018)
			In vitro: NHDFs were obtained via skin biopsy from a healthy young male donor	↑: Nrf2, Nrf2-dependent antioxidant enzymes (NQO-1 and HO-1), procollagen	
	Ginsenoside Rg1	Anti-inflammation and improvement of	Pre-treatment with Rg1 (50 μM) for 1 h	↓: IL6 and 8	Li et al. (2016)
		skin barrier function	In vitro: HaCaT cells	↑: Nrf2, Nrf2-dependent antioxidant enzymes (GCLC, GCLM, and HO-1)	
	Ginsenoside C-Y, a Ginsenoside Rb2 Metabolite from American Ginseng	Hyper-pigmentation	Pre-treatment with C-Y (10, 20, 30 IM) for 72 h <i>In vitro</i> : NHDFs	 t: melanin content and tyrosinase activity t: Nrf2, Nrf2-dependent antioxidant enzymes (NQO-1 and HO-1) 	Liu et al. (2019)

the transcription factor NF- κ B (Wagner et al., 2012). A previous study also suggested a connection between activation of Nrf2 and expression of keratin 16, a key intermediate cytoskeletal protein responsible for maintaining the skin barrier integrity in response to injury or inflammation. While genetic deletion of Nrf2 caused an early onset of hyperkeratotic lesions in *Krt16* null mice which developed palmoplantar keratoma, topical treatment with SFN prevented the development of the skin lesions in footpad skin in association with restoring redox balance (Kerns et al., 2016).

Isothiocyanates including SFN and 7-methylsulfonylheptyl isothiocyanate (7-MSI), the sulfur-rich phytochemicals found in cruciferous vegetables, have been observed to exert antimelanogenic effects via downregulation of MAPKs, the main regulatory pathways of melanogenesis (Shirasugi et al., 2010). 7-MSI treatment significantly reduced melanogenesis in B16F1 melanoma cells *via* activation of ERK signaling, leading to activation of autophagy and downregulation of MITF, tyrosinase and TRP-1 (Kim et al., 2021). Additionally, SFN exerted protective effects on particulate matter-induced melanogenesis *via* decreasing the release of paracrine factors by keratinocytes (Ko et al., 2020). Our previous study also revealed that the mechanisms underlying the anti-melanogenic effects of SFN involved the activation of Nrf2-mediated antioxidant response (Chaiprasongsuk et al., 2016).

Flavonoids

Flavonoids are most abundant polyphenols found in fruits, vegetables, grains, chocolate and beverages including tea and wine. This group has a common basic structure consisting of two aromatic rings bound together by three carbon atoms forming an oxygenated heterocycle. The flavonoids include flavanols (e.g., catechin, epicatechin), flavonols (e.g., quercetin, kaempferol, rutin), flavanones (e.g., hesperetin), flavones (e.g., apigenin, luteolin, hispidulin), isoflavones (e.g., daidzein, genistein) and anthocyanins (e.g., cyanidin) (Panche et al., 2016).

Rutin (Q-3-O-rutinoside) is a flavonol glycoside abundantly found in plants such as tea, onions, wine, apples and berries.

Previous studies reported the protective role of rutin in aging on human dermal fibroblasts (HDFs) via upregulation of collagen type 1 and downregulation of MMP-1 mRNA in HDF. Application of the rutin-containing cream also improved skin elasticity as well as length and area of crow's feet (Choi et al., 2016). Furthermore, the analysis of proteome profiles revealed that rutin treatment caused an induction of proteins involved in the antioxidant defenses and a reduction of proteins involved in the degradation of Nrf2 in UVB-irradiated dermal fibroblasts (Gegotek et al., 2018). Hesperidin, (hesperetin-7-rutinoside), and its aglycone, hesperetin, mostly found in citrus fruits and botanical drugs, have been demonstrated to provide in vitro and in vivo anti-photoaging effects on the skin via stimulating collagen synthesis in association with the antioxidant properties (Lohakul et al., 2021a). Citrus sinensis peel extract containing hesperidin loaded lipid nanoparticles showed photoprotective effects on UVR-mediated induction of MMP-1 via JNK signaling, reduction of collagen accompanied by decreased SOD protein production as well as stimulated inflammatory markers (COXprostaglandin E2) and lipid oxidation product, 2, malondialdehyde (MDA), in mouse skin (Amer et al., 2021). It was also suggested that the protective effect of a mixture of methylated derivatives of hesperidin on UVBinduced skin damage might involve the abilities to promote Nrf2 nuclear translocation and mRNA levels of its target gene GCLC and HO-1 in keratinocytes (Kuwano et al., 2015). In addition, a clinical trial showed that a 12-week topical application of a serum containing 0.1% hesperetin significantly promoted skin hydration and elasticity via enhancing the synthesis of hyaluronic acid (Sheen et al., 2021). Moreover, hesperetin and polyherbal formula extracts containing hesperetin topically applied to mouse skin before UVA exposure three times per week for 2 weeks (a total dose of 60 J/cm²) significantly attenuated MMP-1 upregulation and collagen depletion concomitant with promoting Nrf2 activity and the level of its target proteins (GST and NQO-1) as well as reducing 8-hydroxy-2'deoxyguanosine (8-OHdG), a product of oxidatively damaged DNA damage, in irradiated mouse skin (Lohakul et al., 2021a). Grape fruit stem extract from Muscat Bailey A containing catechin, epicatechin and trans-resveratrol showed protective effects on UVB-induced destruction of collagen fiber through reduction of MMP-1 expression in association with a decrease in lipid peroxidation and restoration of GSH levels in mouse skin (Cho et al., 2018).

Furthermore, flavonoids consequently affect immune mechanisms that are essential in the development of the inflammatory processes (Maleki et al., 2019). Treatment of human epidermal keratinocytes (HaCaT cells) with 6-shogaol, an active ingredient of ginger, resulted in suppressing the UVB (180 mJ/cm²)-induced iNOS, COX-2 and TNF-a, which are the key mediators of inflammatory response, through modulating Nrf2 signaling (Wu et al., 2011; Du et al., 2018; Chen et al., 2019). In addition, the flavonoids have been suggested to exert antiinflammatory actions in association with Nrf2 activation in vitro and *in vivo*. The mechanisms underlying the anti-inflammatory effects of flavonoids involved inhibition of production of proinflammatory cytokines including IL-33, TNF-a, IL-1β, IL-6 and downregulation of NF-KB activity (Staurengo-Ferrari et al., 2018). Gallocatechin-silver nanoparticle was observed to improve wound healing in diabetic rats via inhibiting TLR4/NF-KB inflammatory signaling pathway and modulating Nrf2/HO-1 pathway (Ni et al., 2015). The flavonol Galangin, obtained from Alpinia officinarum Hance and propolis extracts, was able to mitigate imiquimodinduced psoriasis-like skin inflammation in BALB/c mice via inhibiting pro-inflammatory mediators of COX-2, iNOS, NF-KB pathway and pro-inflammatory cytokines IL-17, IL-23, IL-16 in the skin as well as IL-6, TNF-α in both skin and serum. The antiinflammatory effects of galangin were also associated with its ability to induce Nrf2 activity (Sangaraju et al., 2021). Recent evidence has revealed that the flavone luteolin improved impaired healing and promoted re-epithelization of skin wound in streptozotocin-induced diabetic rats via suppressing expressions of inflammatory proteins including MMP-9, TNF-a, IL-6, IL-1β and downregulating NF-KB in association with activation of Nrf2dependent upregulation of antioxidant enzymes (Chen LY. et al., 2021). Topical application of a flavonoid fisetin (50 and 200 µM) for 10 weeks to mouse skin after UVB exposure was demonstrated to mitigate skin photodamage by inhibiting MMP-1 and MMP-2 protein expressions and collagen degradation as well as by improving skin barrier dysfunction. The fisetin treatment resulted in restoring skin hydration and barrier function in UVB-irradiated mouse skin through promoting contents of filaggrin, a structural protein in the stratum corneum, and aquaporins, integral epidermal cell membrane proteins, responsible for epidermal hydration and barrier function. The anti-photodamaging effects of fisetin are suggested to involve upregulation of Nrf2 and downregulation of proinflammatory mediators (COX-2, IL-6, and NF-KB) (Wu et al., 2017).

We previously observed that caffeic acid, ferulic acid, quercetin and rutin provided anti-melanogenic effects *via* enhancing Nrf2-mediated antioxidant defense responses in UVA-irradiated B16F10 cells (Chaiprasongsuk et al., 2016). Ellagic acid was shown to suppress α -MSH-induced melanin synthesis and tyrosinase activity by downregulating cAMPmediated CREB and MITF signaling in B16F10 cells. Ellagic acid also had ability to suppress protein levels of the paracrine factors, POMC and α -MSH, through Nrf2 activation in keratinocyte HaCaT cells (Yang et al., 2021).

Licorice root extracts have traditionally been used for skin problems and are suggested as one the top cosmeceutical ingredients for hyperpigmentation (Searle et al., 2020). The root and rhizome extracts of licorice and several flavonoids identified as its bioactive ingredients have been suggested to provide beneficial effects on the skin through tyrosinase inhibitory activity, ROS scavenging activity, anti-inflammatory activity and Nrf2 inducing activity (Ciganovic et al., 2019). Glycyrrhiza flavonoids and licochalcone A, a major component of the licorice root extracts, showed the inhibitory effects on melanogenesis via activation of ERK and the subsequent downregulation of MITF/tyrosinase pathway in B16F10 cells. Isoliquiritigenin, a flavonoid component from the hydrolysis products of licorice root, was observed to exert antimelanogenic effects on α-MSH-, ACTH- and UV-induced melanin synthesis and on melanocyte dendricity and melanosome transport through downregulation of melanogenic proteins including tyrosinase, TRP-1, DCT, Rab27a and Cdc42 in melanocytes (Lv et al., 2020).

Tannins

There are three major classes of tannins: condensed tannins (e.g., proanthocyanidins, flavonol-based compounds); hydrolyzable tannins (gallotannins and ellagitannins) and phlorotannins. Gallic acid, a chemical constituent of tannins, and its derivatives are found in almost all organ of a plant including bark, wood, leaf (in particular tea leaves), fruit, root and seed.

Red raspberries (Rubus idaeus L.) extracts containing high levels of anthocyanins and ellagitannins including Sanguiin H-6 and lambertianin C showed the protective effect on UVBinduced photoaging in normal human dermal fibroblasts (NHDFs). Treatment with the red raspberry extracts resulted in a significant reduction of MMPs secretion and production of pro-inflammatory mediator IL-6 possibly via downregulating MAPK, NF-KB and AP-1 as well as increased procollagen type I production *via* activating the TGF-β/Smad pathway. The anti-photoaging effects of the tannin-rich botanical drugs involved promotion of Nrf2 activity and its target antioxidants including HO-1 and NQO-1 (Gao et al., 2018b). Alchemilla mollis (Buser) Rothm. ethanolic extract possessing gallic acid showed protective effects on UVBinduced photoaging in NHDFs and in mouse skin in vivo. Treatment with Alchemilla mollis (Buser) Rothm. ethanolic extract led to a substantial reduction in ROS formation as well as MMP-1 and IL-6 promotion through downregulating AP-1 activity in NHDFs exposed to UVB (144 mJ/cm²) irradiation. Furthermore, treatment with Alchemilla mollis (Buser) Rothm. extract and gallic acid protected against UVBinduced a reduction of type I procollagen levels in association with promotion of TGF-B1 in vitro and in vivo. Oral administration of Alchemilla mollis (Buser) Rothm.

extract and gallic acid also improved UVB-induced wrinkle formation, skin dryness, epidermal thickening and collagen fiber density in hairless mice. The antioxidant mechanism underlying the anti-photoaging effects of Alchemilla mollis (Buser) Rothm. extract also involved the upregulation of Nrf2/HO-1 pathway (Hwang et al., 2018). Black tea (Fuzhuan-brick tea, rich in gallic acid and tea polyphenols) and gallic acid were also demonstrated to provide antiphotoaging effects via upregulation of Nrf2/HO-1 signaling in association with activation of MAPK signaling (p38 and ERK1/2 phosphorylation) in UVB-exposed keratinocyte HaCaT cells (Zhao et al., 2018). Cocoa phytochemicals including procyanidins and the flavanol catechin and epicatechin have been suggested to have several biological activities including antioxidant and anti-inflammatory activities possibly responsible for their beneficial effects on various age-related diseases including skin aging (Kim et al., 2014). Long term consumption of cocoa beverage for 12 weeks protected against UV-induced skin erythema and improved skin conditions (including erythema, skin roughness and scaling) in women (Heinrich et al., 2006). Procyanidins showed abilities to activate Nrf2 signaling in various in vitro and in vivo models (Truong et al., 2014). Therefore, it is possible that mechanisms underlying the anti-photoaging effects of procyanidins involve activation of Nrf2-regulated antioxidant defenses.

Phlorotannins (PTNs) are tannins found primarily in brown algae and play a role in protecting cells against UVR. PTNs applied topically attenuated radiation-induced inflammatory responses by downregulating NF- κ B signaling and its downstream COX-2 and inflammasome activation in a mouse model of radiation dermatitis. PTNs also showed the abilities to promote wound healing process by enhancing aquaporin 3 involved in epidermal hydration and homeostasis. The mechanisms underlying the anti-inflammatory and wound healing-promoting effects of PTNs on irradiated mouse skin also involved upregulation of Nrf2/HO-1 signaling (Yang et al., 2020).

Gallic acid, gallotannin, valonia tannin and extracts of plants (e.g., Ceylon olive leaves and pomegranate peel) containing gallic acid exerted antimelanogenic effects directly by acting as a competitive inhibitor of tyrosinase and indirectly by inhibiting tyrosinase via antioxidant actions that affect melanogenesis pathway in vitro and in vivo (Chen et al., 2009; Panich et al., 2013; Su et al., 2013; Kanlayavattanakul et al., 2020; Huang et al., 2021; Liu et al., 2021). A randomized, double-blind, placebocontrolled clinical trial demonstrated that continuous administration of apple polyphenol rich in procyanidins for 12 weeks improved UV-induced skin pigmentation in heathy women (Shoji et al., 2020). Our previous study demonstrated that gallic acid protected against UVA (8 J/cm²)-induced melanogenesis via modulation of Nrf2 signaling and promotion of antioxidant defenses (including GSH, catalase, GPx and GST) in B16F10 melanoma cells (Onkoksoong et al., 2018).

Resveratrol and its Derivatives

Resveratrol was observed to provide photoprotective effects in UVB-induced photoaging *via* the antioxidant, anti-inflammatory

and antiapoptotic actions in human keratinocyte HaCaT cells and ICR mice *in vivo*. The mechanism underlying its antiphotoaging actions involves upregulation of Nrf2 signaling and the antioxidant defenses (including HO-1, NQO-1, SOD1, GPx-4) in association with suppression of aging markers (MMP-1 and MMP-9) and proinflammatory mediators (IL-6, TNF- α , VEGF-B) by inhibiting ROS-mediated MAPK and COX-2 signaling cascades (Cui et al., 2022). Oral administration of grape peel extract and resveratrol exerted the anti-photoaging effects on UVB-induced skin wrinkle formation *via* promotion of Nrf2/HO-1 signaling cascades (Kim et al., 2019). In addition, a formulation containing 0.4% pterostilbene, the resveratrol analog, extracted from the dried heartwood of *Pterocarpus marsupium* Roxb., showed substantial reduction of aging markers and improvement of wrinkles, skin hydration elasticity in healthy volunteers (Majeed et al., 2020).

Topical treatment with pterostilbene, the resveratrol analog, suppressed an acute UVB radiation-induced skin inflammation and prevented chronic UVB radiation-mediated carcinogenesis in mice. The mechanisms involved in the photoprotective effects of pterostilbene might be attributed its ability to absorb UVB, protect against oxidative damage to DNA, protein and lipid, promote activities of antioxidant enzymes including catalase, SOD and GPx as well as activate Nrf2-dependent antioxidant response (Pastore et al., 2012; Sirerol et al., 2015). Moreover, resveratrol was observed to promote wound healing by restoring cell proliferation and migration, along with increased Nrf2 activity and Mn-SOD expression in the diabetic rat model (Zhou et al., 2021).

Resveratrol acting as a direct antioxidant, Nrf2 activator and tyrosinase inhibitor has been suggested to suppress melanogenesis (Wang et al., 2018; Boo, 2019). Resveratrol treatment led to a substantial reduction of UVB-induced melanogenesis via downregulation of MITF and its target proteins including TYR, TRP1, TRP2 in correspondence to upregulation of Nrf2/HO-1 proteins in melanocytes (Jian et al., 2011). Topical administration of 0.4% resveratryl triacetate and 0.4% resveratryl triglycolate twice daily for up to 8 weeks after the artificial pigmentation was shown to provide whitening efficacy in human subjects (Ryu et al., 2015; Jo et al., 2018). Pterostilbene, a stilbinoid, found in blueberries and grapes was demonstrated to exert anti-melanogenic effects via promotion of autophagy in melanocytes and downregulation of CREB (cAMP response element-binding protein)-MITF-tyrosinase pathway in B16F10 cells treated with HaCaT conditioned medium. The protective effects of pterostilbene on melanogenesis involved suppression of UVA-induced a-MSH expression and upregulation of Nrf2-mediated HO-1, y-GCLC, and NQO-1 protein expressions in HaCaT cells (Hseu et al., 2021). Additionally, the application of a 0.4% formulation of natural pterostilbene for 4 and 8 weeks showed the skin brightening and anti-aging effects, respectively, in healthy volunteers in an open-label, single-arm, monocentric efficacy study (Majeed et al., 2020).

Carotenoids

Carotenoids, which belong to the tetraterpenes family, are liposoluble pigments responsible for the yellow, orange or red

color of fruits, leaves and flowers. The carotenoids are divided into carotenes, xanthophylls and lycopene, and are abundantly present in tomato, carrots, pumpkin, seaweeds and algae (Milani et al., 2017). The main carotenoids including astaxanthin, canthaxanthin, β -carotene, lycopene and lutein have been suggested to exert photoprotective effects against UVRinduced skin photodamage via inhibition of inflammatory responses and photoaging biomarkers as well as promotion of antioxidant defense system in several in vitro, in vivo and clinical studies (Aust et al., 2005; Camera et al., 2009; Cavinato et al., 2017; Cooperstone et al., 2017; Kohandel et al., 2022). Previous reports have suggested that the carotenoids including astaxanthin and lycopene exert pharmacological activities including antioxidant, anti-inflammatory and chemopreventive activities against skin damage via activation of Nrf2 signaling (Wang et al., 2020; Ahn and Kim, 2021; Kohandel et al., 2021).

Dietary carotenoids combined with rosemary extract containing carnosic acid having abilities to activate Nrf2/ARE system exerted an inhibitory effect on UVB (20–60 mJ/cm²)-induced TNF- α and MMP-1 secretion from dermal fibroblasts (Calniquer et al., 2021). The tomato extracts rich in lycopene also protected against H₂O₂-induced photodamage of fibroblasts via promotion of pro-collagen secretion and suppression of apoptotic cell death and ROS formation (Darawsha et al., 2021).

Lutein, a xanthophyll carotenoid obtained from green leafy vegetables and egg yolk, has been reported to exert the antiinflammatory effects via modulation of oxidant-sensitive inflammatory signaling pathways including NF-KB and STAT3 pathways and suppression of inflammatory cytokines (such as IL-1β, IL-6, TNF-α, COX-2, iNOS) (Ahn and Kim, 2021). Astaxanthin was demonstrated to exert anti-inflammatory effects by suppressing the expression of pro-inflammatory cytokines, for example, COX-2, LOX-1, NF-kB p65, TNF, and IL-1 (Kishimoto et al., 2010). Additionally, combinations of tomato nutrient complex containing lycopene and carotenoids and rosemary extract containing carnosic acid were observed to protect against UVB-induced oxidative stress, photoaging and inflammatory responses by inhibition of NF-KB activity and IL-6 production, along with activation of the ARE/Nrf2 system using HaCaT and KERTr keratinocyte cell lines (Calniquer et al., 2021). Bixin, an apocarotenoid isolated from the Achiote (Bixa Orellana L.), is one of the most consumed food colorants and topical preparations of seed extracts of the achiote have been in ethno-pharmacological use for treatment of wound healing and the pathologies related to epithelial barrier disruption. The mechanisms underlying the pharmacological actions of bioxin in improving skin barrier function were suggested to involve activation of Nrf2mediated antioxidant systems including thioredoxin (TRX)/ thioredoxin reductase (TXNRD1), regulation of peroxisome proliferator-activated receptors (PPARs), responsible for skin homeostasis and epithelial repair and modulation of Toll-like receptor 4 (TLR4)/NF-KB inflammatory signaling pathway (Rojo de la Vega et al., 2017). The marine carotenoid fucoxanthin (found in brown seaweeds, the microalgae and diatoms) showed anti-inflammatory actions through inhibiting proinflammatory cytokines including TNF-a, IL-6

and IL-8 levels as well as suppressing UVB-mediated oxidative stress in keratinocyte HaCaT cells. Moreover, topical application of cream containing the fucoxanthin to mouse skin protected against UVB-induced skin inflammation and hyperplasia *via* downregulation of COX-2 and iNOS, along with upregulation of Nrf2 activity and its target protein HO-1 (Rodriguez-Luna et al., 2018).

Terpenoids: Diterpene, Triterpene and Sesquiterpene

Santamarine, a sesquiterpene lactone, isolated from sunflower family provided anti-photoaging effects via suppression of UVA (8 J/cm²)-induced MAPK/AP-1 pathways involved in upregulation of MMPs and via promotion of TGF-B/Smadmediated collagen production in HDFs. Furthermore, santamarine treatment led to a significant restoration of UVAmediated downregulation of Nrf2-dependent antioxidant defenses including SOD-1 and HO-1 at the mRNA and protein levels (Oh et al., 2021). Zerumbone (ZER), a natural sesquiterpene, from Zingiber zerumbet (L.) Roscoe ex Sm. rhizomes was demonstrated to protect against UVA irradiation (3 J/cm²)-induced ROS formation, MMP-1 activity and collagen III degradation in HDFs. ZER was suggested to exert the anti-photoaging effects via downregulation of AP-1 activity and promotion of Nrf2/ARE pathway (Hseu et al., 2019). Furthermore, treatment of dermal fibroblasts with rosemary extracts rich in the diterpene carnosic acid having abilities to induce ARE/Nrf2 reporter activity protected against TNF-ainduced MMP-1 secretion (Calniquer et al., 2021). Ginsenosides, triterpene saponins, are major bioactive compounds responsible for pharmacological activities of Panax (ginseng), which has traditionally been used to treat and prevent various conditions associated with aging including skin aging. Previous studies demonstrated that the rare minor ginsenosides (C-Mc and Mx), which may act as potential antiphotoaging compounds, suppressed MMP production via regulating MAPK/AP-1/NF-ĸB and promoted pathway collagen production via the TGF-\u03b3/Smad pathway in association with upregulation of Nrf2 signaling in UVB-irradiated human dermal fibroblasts (Liu et al., 2018; Liu et al., 2022).

Cynaropicrin, a sesquiterpene lactone, is the major bioactive phytochemical in the artichoke (Cynara cardunculus L.) that can activate aryl hydrocarbon receptor (AhR), resulting in nuclear translocation of Nrf2. Activation of AhR/Nrf2/NQO-1 pathway by cynaropicrin was involved in its inhibitory effects on UVB-mediated production of proinflammatory cytokines including IL-1 and TNF- α in keratinocytes (Takei et al., 2015). Hemistepsin A, a sesquiterpene lactone isolated from Saussurea lyrata (Bunge) Franch., has been demonstrated to exert pharmacological actions including anti-inflammatory and antioxidant activities. Treatment of keratinocyte HaCaT cells hemistepsin A protected against H₂O₂-induced with cytotoxicity, DNA damage and apoptosis-mediated by mitochondrial dysfunction via upregulation of Nrf2/HO-1 signaling pathway (Park et al., 2020). The ginseng leaf extract rich in ginsenosides (including ginsenoside Rg1) applied topically to mouse skin protected against UVB-induced photoaging and skin barrier dysfunction through suppression of MMP-2, MMP-9 and MMP-13 protein expressions (Hong et al., 2017). Moreover, ginsenoside Rg1 showed anti-inflammatory effects against UVBinduced glucocorticoid resistance in keratinocyte HaCaT cells via promotion of Nrf2 activity (Li et al., 2016).

The sesquiterpene lactone eupalinolide A and B from *Eupatorium lindleyanum* DC. showed inhibitory effects against UVB-induced melanogenesis, skin damage and inflammatory responses *in vitro* and mouse skin *in vivo* (Yamashita et al., 2012). Previous evidence revealed that, apart from the anti-photoaging effect, a minor ginsenoside (C-Y) having ability to induce Nrf2 activity was observed to exert whitening effects by inhibiting melanin production, tyrosinase activity in Melan-a and zebrafish embryos (Liu et al., 2019).

Cannabidinol

Cannabidiol (CBD), the second most prevalent active ingredient in cannabis, is the non-psychoactive phytocannabinoid that has antioxidant and anti-inflammatory effects. CBD has been reported to provide photoprotective effects against UVA and UVB-induced damage to skin cells including NHDFs and keratinocyte HaCaT cells (Vacek et al., 2021). Treatment of 2D and 3D cultured fibroblasts with CBD caused a substantial attenuation in the levels of lipid peroxidation-derived aldehydes (4-hydroxynonenal (HNE), MDA and acrolein-protein adducts) in UVA (20 J/cm²)- and UVB (200 mJ/cm²)-irradiated cells (Gegotek et al., 2019). The formation of aldehyde-protein adducts induced by the highly reactive aldehydes could subsequently change the structure and/or function of several proteins including main ECM elastin in hairless mice exposed to UVA (Larroque-Cardoso et al., 2015). In addition, 4-HNE was suggested to play a role in UVAinduced fibroblast senescence in skin photoaging (Swiader et al., 2021). The compounds having abilities to neutralize aldehydes and inhibit the formation of protein adducts could thus have a beneficial role against skin photoaging.

Moreover, the analytical chemistry revealed that CBD compound could interact with the Nrf2/NF- κ B transcriptional activity (Jastrzab et al., 2019). The biological effects of CBD have been found to maintain membrane integrity by preventing protein and phospholipid modifications (Atalay et al., 2020) and prevent the inflammatory responses (nuclear receptor coactivator-3 and paralemmin-3) (Atalay et al., 2021).

CONCLUSION AND FUTURE CHALLENGES: INSIGHT INTO ETHNOPHARMACOLOGY

Ethnopharmacology is defined as "the interdisciplinary exploration of biologically active agents traditionally employed or observed by man" (Bruhn and Rivier, 2019). Identifying the ingredients and exploring the effects of the ingredients are crucial in the study of traditional medicine. Phytochemicals are bioactive compounds in plant-based products that have been historically used to rejuvenate the skin and alleviate skin disorders. Bioactive compounds of plant origin have thus been considered as invaluable sources of potential preventive or therapeutic agents for dermatological indications due to their pharmacological activities including antioxidant, UV absorption and anti-inflammation. The phytochemicals exert antioxidant effects by directly scavenging ROS or by promoting the antioxidant defense system through activation of Nrf2 signaling. It should also be taken into consideration that while Nrf2 plays a crucial role in maintaining cellular homeostasis under stress and inflammatory conditions, several studies have discussed a detrimental aspect of Nrf2 defined as the "dark side of Nrf2" in the cancer biology as enhanced Nrf2 activity is involved in a pro-carcinogenic effect and therapeutic resistance of cancer cells (Sporn and Liby, 2012). Thus, dietary phytochemicals having the potential to provide either chemopreventive or cancer-promotive properties, depending on the stage of carcinogenesis (L Suraweera et al., 2020). This review discusses the protective role of Nrf2 against UVRinduced skin photodamage and thus application of phytochemicals acting as Nrf2 activators is regarded as a promising strategy to prevent and treat premature aging and photodamage-related skin problems. Targeting Nrf2dependent antioxidant and cytoprotective response has been suggested to represent a promising pharmacological strategy for development of effective and safe anti-photoaging and photoprotective agents. Furthermore, the therapeutic potential of phytochemicals can be limited by their poor bioavailability and thus development of drug delivery systems (such as nano-engineered formulations) is needed to improve efficacy of promising bioactive compounds as effective photoprotective agents.

AUTHOR CONTRIBUTIONS

AC wrote the manuscript and created the graphical figure and tables; UP contributed to the design, the conception of the study and wrote the manuscript. All authors have read and approved the final manuscript.

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