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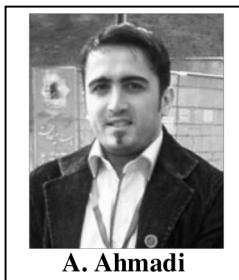
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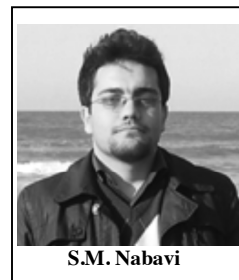
The Role of Hesperidin in Cell Signal Transduction Pathway for the Prevention or Treatment of Cancer

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Abstract: During past two decades, plant-derived bioactive compounds have been reported as novel therapeutic agents for prevention and/or mitigation of different human diseases such as cancer, inflammation, cardiovascular and neurodegenerative diseases. Hesperidin is known as one of the most common and bioactive constituents of *Citrus* (C) species which possesses multiple health-promotion effects. A plethora of scientific literature reported that hesperidin possesses *in-vitro* and *in-vivo* anticancer activities. In addition, there are numerous scientific evidences regarding the molecular mechanisms of anticancer activities of hesperidin and its aglycone, hesperetin. However, in this case, the number of comprehensive reviews on molecular mechanisms underlying the anticancer effects of hesperidin is sparse. Therefore, in this work we present a critical review of the available literature regarding the molecular mechanisms of the anticancer effects of hesperidin and its aglycone, hesperetin.

Keywords: Aglycone, anticancer, citrus, flavonoid, hesperetin, hesperidin.

1. INTRODUCTION

Signal transduction is the process of conversion of external signals produced by outside stimuli, including some hormones, growth factors, neurotransmitters, inflammatory cytokines and chemokines, to a biochemical response leading to a cellular response [1]. As a result of these responses, changes in cellular metabolism, expression of different genes, cell division as well as cell death may occur. It has been reported that in most diseases, these signaling dysfunctions have occurred. Cancer is a complex process, comprised of different stages such as initiation, promotion, and progression. Among them, the promotion stage has advantages over the others in therapeutics intervention because

during this stage the single initiated cancerous cell proliferated and became resistant. Therefore, comprehending the molecular mechanisms of promotion and progression is critical for the discovery and development of chemopreventive compounds with high efficacy [2].

During past two decades, much attention has been paid to the discovery of novel therapeutic agents from nature [3-7]. Because of the availability, efficacy, and safety of herbal compounds, recent research has been focused on the beneficial effects of plant-derived bioactive compounds [8-14]. Among them, flavonoids are known as one of the most common groups of bioactive phytochemicals that possess multiple pharmacological actions such as antioxidant, anticancer, anti-inflammatory, etc [15-18].

Hesperidin is known as one of the most common flavonone glycosides, which is found in different *Citrus* (C) species. It has been reported that hesperidin possesses different pharmacological effects such as antioxidant, anti-inflammatory, and anticancer effects [19]. A growing body of evidence has shown potent antican-

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cer effects of both hesperidin and its aglycone form [20, 21]. Furthermore, it can be hypothesized that anti-inflammatory and anti-oxidant effects of hesperidin and its aglycone play crucial roles in the anticancer effects of these compounds [22]. In the chemical structure of hesperidin, the presence of hydroxyl moieties is responsible for these pharmacological effects.

Chemically, hesperidin consists of an aglycone (the form lacking the sugar moieties), hesperetin, and the sugar rutinose: hesperetin-7-rutinoside. Hesperidin is the main flavonoid in *C* fruits, and can be found in large concentrations in *C* rinds. Until now, it has been isolated from rinds of different *C* species such as *C. aurantium* L., *C. sinensis* L., and *C. unshiu* Marcov. [23].

Hesperidin is absorbed from the intestine intact as a glycoside. Its aglycone hesperetin appears in the plasma 3 h after ingestion and reaching a peak concentration between 5 and 7 h. It has also been reported that glucuronides and sulfoglucuronides are known as circulating forms of hesperetin [24]. In the present review, we have focused on anticancer effects of hesperidin in addition to its chemistry and natural sources.

2. CHEMISTRY OF HESPERIDIN

Hesperidin is the glycoside form of hesperetin (methyl eriodictyol) (Fig. 1), which is bonded to rutinose [25, 26]. It has been reported that the glycoside moiety in the chemical skeleton of hesperetin is a disaccharide generated from rhamnose and glucose and widely found as rutinose and/or neohesperidose [26].

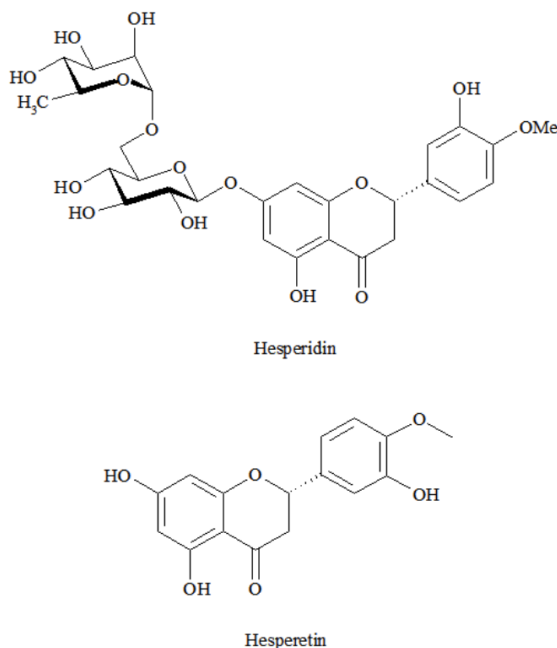


Fig. (1). Chemical structures of hesperetin and hesperidin.

It is well known that rutinose (6-*O*-(α -L-rhamnopyranosyl)-D-glucopyranose and/or 6-*O*-(α -L-rhamnosyl)-D-glucose) is widely found in nature [26-28]. Another isomeric form, neohesperidose (2-*O*- α -L-rhamnopyranosyl-D-glucopyranose), has been identified in different *C* species as hesperetin 7-*O*-neohesperidoside [29, 30]. It is well known that the presence and structures of the glycoside moieties are responsible for *C* bitterness [26]. It has also been reported that the presence of rutinoides and neohesperidosides is responsible for tastelessness and bitterness of *C* species [26]. A plethora of evidence has shown that hesperidin is most frequently identified as the rutinoides form, which is non-bitter [26, 31, 32]. However, the neohesperidoside form of hesperidin has been identified in grapefruit [26, 32, 33]. The aglycone form of hesperidin (hesperetin, (*S*)-2,3-dihydro-5,7-dihydroxy-2-(3-hydroxy-4-methoxyphenyl)-4*H*-1-benzopyran-4-one) is a bioflavonoid with a heterocyclic structure [25, 26, 34]. It has been reported that the presence and number of hydroxyl moieties in both the aromatic and heterocyclic rings of bioflavonoids are responsible for their pharmacological effects [26]. For example, it has been reported that the presence and number of hydroxyl moieties in the structure of hesperetin closely correlates with its antioxidant activity [26].

3. NATURAL SOURCES

A number of sources have shown that hesperetin has been identified in different parts of *C* such as the epicarp, mesocarp, and endocarp [26, 35]. Hesperetin has also found in the *C* juice and seeds [36]. A growing body of literature has shown that hesperidin is found predominantly in the rinds of *C. aurantium*, *C. sinensis*, and *C. unshiu* in large scale amounts [37-40]. It has been found that the total amount of hesperetin in albedo, membranes and pith of *C* is higher than juice vesicles and seeds. Peppermint is another important and common source of hesperidin [41].

However, a close correlation between level of hesperetin and stage of fruit maturity has been reported [26, 42]. In addition, seed germination has positive effects on the level of hesperetin, which is due to the effects of light exposure that stimulates hesperetin production [26, 42, 43].

In addition to *C* species, hesperetin has also been found in several species from the families Fabaceae, Betulaceae, Lamiaceae and Papilionaceae [26, 44-46]. Hesperetin has also been found in *Acanthopanax setchuenensis* (Araliaceae), *Zanthoxylum avicennae* and *Zanthoxylum cuspidatum* (Rutaceae) [47-49]. Fur-

thermore, hesperetin has been identified as neohesperidin form in *Cynara* (Asteraceae) species [26, 50].

4. ANTICANCER EFFECTS OF HESPERIDIN: CELL SIGNAL TRANSDUCTION PATHWAY

4.1. Janus Kinase-signal Transducers and Activators of Transcription (JAK-STAT) Pathway

The Janus kinase/signal transducer and activator of transcription, JAK/STAT, is known as one of the most important signaling pathways for different cytokines and growth factors. This pathway affects many cellular functions such as metastasis, proliferation, growth, and immune response [51]. Up until now, 4 mammalian JAKs (JAK1, 2, 3 and Tyk2) and 7 mammalian STATs (STAT1, 2, 3, 4, 5a, 5b and 6) have been recognized. Of the seven STAT family members, STAT1 and STAT3 have been involved in cancer progression but have opposite effects. STAT1 has been reported to enhance anti-proliferative and pro-apoptotic responses in cancerous tissues while STAT3 can significantly enhance proliferation [52]. Constitutive activation of STAT is promoted by immoderate stimulation through cytokines and/or growth factors.

Through the activation of various tyrosine kinases, STATs can undergo phosphorylation, dimerization, and nuclear localization, and these proteins can bind with specific Deoxyribonucleic acid (DNA) elements to activate transcription of targeted genes [53]. It has also been reported there is a close correlation between JAK/STAT activation and an increase in the invasion and metastasis of different types of cancer. Among the target gene of STATs, cyclins D1/D2, Myc, B-cell lymphoma-extra-large (Bcl-xL), and Myeloid cell leukemia 1 (Mcl-1) are known to play crucial roles in cancer metastasis through activation of cell cycle and apoptosis [54]. These cellular events play important roles in regulation of hematopoiesis, immune development, tumor biology and other processes [55]. Many studies have demonstrated that the JAK/STAT pathway is involved in various cancers such as prostate, breast, pancreatic, and colon [56-59]. In addition, there is evidence that various chemopreventive agents are potent inhibitors of the JAK-STAT signaling pathway and cancer metastasis [60].

Furthermore, it has been reported that JAK/STAT signaling can be activated by interleukin 6 (IL6), which leads to an increase in the migration and invasion under in-vitro conditions and metastasis under in-vivo conditions [62]. Li *et al.* reported that 7,3'-dimethoxyhesperetin repressed the serum levels of IL-6 as well as IL-6 mRNA expression in the synovium of

rats with adjuvant arthritis. They found that 7,3'-dimethoxyhesperetin significantly decreased mRNA expression of JAK2 and STAT3 as well as protein expression of p-JAK2 and p-STAT3 in the synovium of rats [61].

Another study showed that 5,7,3'-triacetylhesperetin (TAHP) (66, 132 mg/kg) down-regulated the serum level of IL-6 and also decreased the level of IL-6 in the synovial tissues of adjuvant arthritis rats. In addition, it was shown that administration of 5,7,3'-triacetylhesperetin reduced expression of mRNA of STAT3 and JAK2, as well as the ratio of p-JAK2/JAK2 protein and p-STAT3/STAT3 protein in the synovial tissues of rats [62].

The inhibition of STAT3 phosphorylation has led to a reduction of STAT3 expression targets such as cyclin D1, survivin, and Bcl-xL, which are responsible for the survival of cancerous and cancer stem cells (CSCs). In addition, oxidized low-density lipoprotein (LDL) caused the activation of JAK2, STAT1, and STAT3 via oxidative stress in the cells. It has been reported that hesperetin significantly reduced the oxidized LDL-induced 2',7'-dichlorofluorescein staining, which is due to the inhibitory effects of hesperetin against oxidized LDL-triggered reactive oxygen species and apoptosis. Choi *et al.* indicated that hesperetin decreased p38MAPK phosphorylation and its downstream c-myc and signal STAT1, which was caused by oxidized LDL. Furthermore, hesperetin suppressed oxidized LDL-activated JAK2/STAT3-dependent signaling pathways [63] (Fig. 2).

4.2. The PI 3-Kinase/Akt and mTOR Pathways

The phosphatidylinositol-3-kinase (PI3K)/Akt and the mammalian target of rapamycin (mTOR) pathway control various biological processes and play an important role in different aspects of cell growth and survival in physiological as well as in pathological conditions [64].

It has been reported that activation of several growth factor receptor protein tyrosine kinases results in autophosphorylation on tyrosine residues. Thereafter, PI3K is recruited to the membrane and due to its activation, second messenger phosphatidylinositol-3,4,5-triphosphate (PI3,4,5-P3) from the substrate phosphatidylinositol-4,4-bisphosphate (PI-4,5-P2) is produced. Subsequently phosphatidylinositol 3-phosphate (PIP3) recruits 3-phosphoinositide dependent protein kinase-1 (PDK1) and Akt/protein kinase B (PKB), signaling proteins with pleckstrin homology (PH) domains, to the membrane [60].

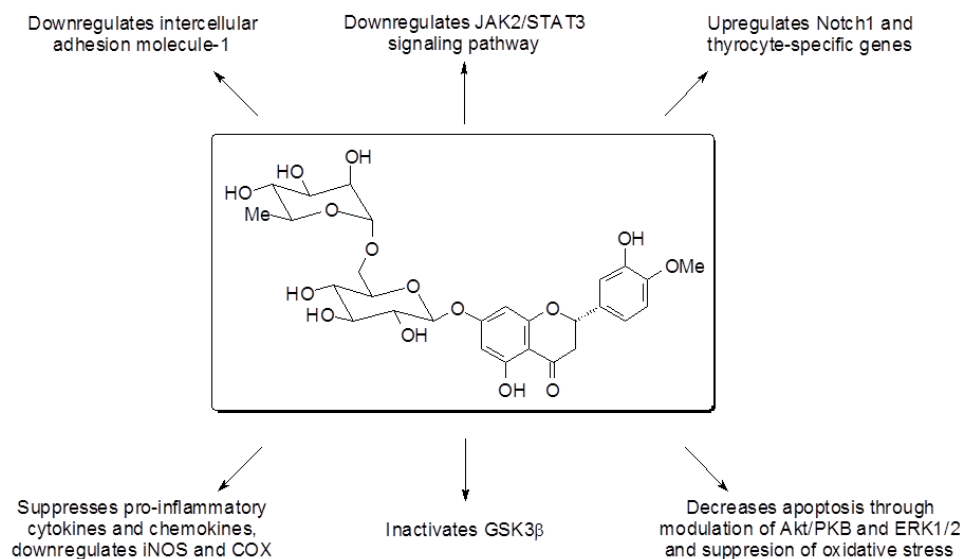


Fig. (2). Molecular mechanisms underlying anticancer effects of hesperidin.

Full activation of Akt also requires phosphorylation at a second site by a distinct protein kinase (mTORC2), which is also stimulated by growth factors. It has been reported that once activated, Akt phosphorylates some of the target proteins such as proteins that are known as direct regulators of cell survival transcription factors, as well as other protein kinases [65].

Akt/PKB and its substrates such as Bad, GSK-3, pro-caspase9, I κ B kinase (IKK), forkhead transcription factors and Yes-associated protein (YAP) have anti-apoptotic role [66]. mTOR is a serine/threonine protein kinase, and its activity is due to two different complexes, mTORC1 and mTORC2, which regulate metastasis cascades including cell growth, proliferation, motility, survival, and cell invasion and migration. In addition, the canonical pathway of mTOR activation depends on mitogen-driven signaling by the PI3K/Akt pathway, although alternative non-Akt-dependent activation by the Ras/MEK/ERK pathway has also been identified.

Taken together, activation of this pathway leads to increased protein synthesis. These include several that have been implicated in the pathogenesis of various tumors, e.g., cyclin D1, which allow progression of cells through the cell cycle, and hypoxia-inducible factor (HIF), which drive the expression of pro-angiogenic growth factors such as vascular endothelial growth factor (VEGF) [64]. Several phytochemicals have been identified that suppress the activation of Akt/PI3K/mTOR pathway. Saiprasad *et al.* indicated that hesperidin caused apoptosis and also triggers autophagic factors via suppression of Aurora-A-mediated PI3K/

Akt/mTOR and glycogen synthase kinase 3 beta (GSK-3 β) signaling cascades in experimental colon carcinogenesis. Their finding showed that Akt and PI3K phosphorylation and activation increased in azoxymethane (AOM) induced animals. Phosphatase and tensin homolog (PTEN), a tumor suppressor, negatively regulates the PI3K and Akt activation suppressing tumorigenesis. Hesperidin treatment restored the decreased PTEN activity seen in AOM-induced animals and this may likely be a reason for hesperidin's inhibitory role against PI3K/Akt signaling cascade.

Meanwhile, hesperidin significantly reduced mTOR expressions associated with an increase in autophagic proteins Beclin-1 and microtubule-associated protein 1A/1B-light chain 3-phosphatidylethanolamine conjugate (LC3-II). GSK-3 β is another downstream effector of the PI3K/Akt pathway [53]. Moreover, GSK-3 β initiates the degradation and inactivation of several oncogenic transcription factors (c-myc and c-jun) and proto-oncoproteins (β -catenin) via phosphorylation. These results have demonstrated that hesperidin administration significantly inhibits phosphorylation of GSK-3 β , leading to the reduction of oncogenic β -catenin, c-myc and c-jun. Indeed, this study revealed that hesperidin was able to actively inhibit the PI3K/Akt/GSK-3 β and mTOR signaling cascades [67].

In nervous tissues, hesperetin (100 nmol/L) and its metabolite 5-nitro-hesperetin efficiently prevent apoptosis via Akt/PKB activation/phosphorylation and also via an activation of extracellular signal-regulated kinase 1/2 (ERK1/2) [68] (Fig. 2).

4.3. Notch Pathway

The Notch pathway plays an important role in the cell fate determination, proliferation, differentiation and survival at different stages including development, neurogenesis and homeostasis. It has also been involved in cell–cell communications. Direct interaction of Notch family receptors (Notch 1–4) with ligands (JAG1, JAG2, Dll-1, Dll-3, and Dll-4), which are expressed on adjacent to cells, leads to initiation of Notch pathways. It has also been reported that gamma-secretase stimulates the Notch intracellular domain releasing from the membrane. Thereafter, the Notch intracellular domain translocates into the cell nucleus and interacts with DNA-binding transcription factors and in this way leads to Notch target genes. Impairment of this signaling pathway is linked to several diseases including cancer. The aberrant regulation of Notch signaling contributes to tumor development [69]. Notch genes possess oncogenic and also tumor suppression activities in different malignancies for example, inducing keratinocyte growth arrest in the skin [70]. It has also been reported that Notch inhibitors or inducers can serve as growth inhibitors. Notch1 signaling activity is downregulated in many cancers such as thyroid cancer, pancreatic cancer, liver cancer, brain tumor, etc.

Previous studies have suggested that Notch signaling is reduced in neuroendocrine tumors, including carcinoids, and it has been shown that increased signaling of this pathway reduces carcinoid cell proliferation as well as neuroendocrine tumor marker expression among the tumors [71]. Zarebczan and colleagues demonstrated that hesperetin causes expression of Notch1 among the carcinoid cells (BON), with subsequent suppression of cell proliferation and hormone generation. Indeed, hesperetin reduced the expression of neuroendocrine tumor markers such as ASCL1 and CgA [72]. In anaplastic thyroid cancer (ATC), Notch1 acts as tumor suppressor. Therefore restoration of functional Notch1 signaling is a therapeutic approach in ATC. Recently, Patel *et al.* showed that administration of hesperetin induces expression of Notch1 and its downstream effectors hairy and enhancer of split 1 (Hes1), and Hes1 related with YRPW motif was observed in hesperetin-treated ATC cells. Hesperetin exhibited antiproliferative effect on ATC cell proliferation. Inhibition of cell growth was mainly due to apoptosis as demonstrated by an increase in the ratio of cleaved poly ADP ribose polymerase (PARP) and cleaved caspase-3 as well as diminished survivin. Moreover, these workers found that hesperetin administration upregulates thyrocyte-specific genes including

thyroid transcription factor 1 (TTF1), TTF2, paired box gene 8, thyroid stimulating hormone receptor, and sodium/iodide symporter [73] (Fig. 2).

4.4. MAPK-ERK Pathway

Mitogen-activated protein (MAP) kinases are known to be regulators of a wide range of cellular processes including cell growth and differentiation, gene expression, mitosis, cell motility, metabolism, survival as well as apoptosis by phosphorylation of target protein substrates. The classic MAP kinases are classified into three sub-families including extracellular signal-regulated kinases (ERK; ERK1 and ERK2), c-Jun N-terminal kinases (JNK; JNK1, JNK2, and JNK3), as well as p38 mitogen-activated protein kinase (p38-MAP kinases) (a, b, d, and g) [74, 75].

Phytochemicals such as hesperidin suppressed ROS-induced inflammations that are involved in most diseases including cancer. Moon and Kim demonstrated the anti-inflammatory effects of HES in HaCaT cells. Their result indicated that HES inhibited the hydrogen-peroxide-induced interleukin-8 and tumor necrosis factor- α generation and also p38 phosphorylation as well as activation of COX-2. [76]. In addition, ERK activation has been reported in response to oxidative damage and served as an important factor in the cell survival. Peroxynitrite is also known as an important contributor in the initiation and progression of different diseases such as cancer, cardiovascular and neurodegenerative diseases. Hesperetin exposure to fibroblasts is shown to upregulate ERK1/2 phosphorylation and to modify peroxynitrite-mediated reduction in ERK1/2 phosphorylation [77]. A study by Pollard *et al.* has shown that modulation of fibroblast signaling, as well as scavenging of intracellular peroxynitrite, plays important role in the promising effects of hesperetin on fibroblasts [77].

In addition, treatment with hesperetin modified the kidney nuclear factor-kappa B (NF- κ B)-mediated gene expression through regulation of ERK, Jun N-terminal kinase (JNK) and p38 in rats [78]. Rainey *et al.* concluded that hesperetin significantly increased the level of ERK1/2 phosphorylation when used at concentrations of 100–300 nM, although, at this concentration, hesperetin did not increase CREB phosphorylation. Treatment with hesperetin (300 nM concentration) reduced staurosporine-caused cell death among primary neurons [79]. Indeed, it was shown that hesperetin and 5-nitrohesperetin, when used at the concentration of 100 nM/L, were effective in preventing neuronal apoptosis via two mechanisms: activation/phosphorylation of both ERK1/2 and Akt/PKB [68, 80] (Fig. 2).

4.5. Wingless and INT-1 (Wnt) Pathway

Members of the Wnt family are secreted growth factors that bind to a complex of receptors of the Frizzled families (canonical pathway). These proteins also activate non-canonical pathways by stimulating cytoskeletal reorganization and regulating planar cell polarity. Stabilization of β -catenin occurs due to Frizzled signaling, which acts as a transcriptional activator in the Wnt pathway [81].

Wnt/ β -catenin pathway was reported to regulate the process of cell proliferation, migration, apoptosis, differentiation, epithelial–mesenchymal interactions, as well as stem cell self-renewal. It has also been reported that β -catenin is phosphorylated through GSK-3 β in complex with the proteins axin and APC in the absence of Wnt signaling. β -Catenin phosphorylation leads to its ubiquitination and degradation. Due to Wnt binding, Disheveled phosphorylation occurs (a cytoplasmic protein that interacts with Frizzled), which then increases phosphorylation of LRP. This process provides binding sites for axin, leading to association of axin with the receptor, followed by disruption of the connection complex of axin, casein kinase-1 and GSK-3 β with β -catenin. Disruption of the axin complex averts β -catenin degradation [65].

Many studies indicated that Wnt/ β -catenin activation is implicated in most human cancers including gastric, leukemia, endometrial, thyroid, melanoma, ovarian, breast, colon, liver, lung and head and neck. Wnt signaling increases accumulation of β -catenin within the cell nucleus, leading to the consequent transcriptional activation of some target genes such as c-myc, cyclin D, c-jun, gastrin, CD44, surviving, VEGF, endothelin-1, interleukin β -8, nanog and snail [60].

Melanogenesis is known to be a complex process that regulates skin pigmentation to protect against photodamage. Huang and colleagues evaluated hesperetin activities on melanin synthesis among murine B16-F10 melanoma cells. Their results showed that hesperetin increased mitogen-activated protein kinases activation, cAMP-responsive element binding protein phosphorylation and glycogen synthase kinase-3 β , and led to induced β -catenin accumulation. GSK3 β was inactivated by hesperetin through phosphorylation of the Serine 9 site of this protein kinase. Thus, β -catenin degradation is reduced by hesperetin. The Wnt/ β -catenin pathway plays an important role in melanocyte differentiation [82].

Another study conducted by Kim *et al.* indicated that hesperetin diminished suppression actions of high

glucose during the differentiation of osteoblasts in periodontal ligament stem cells. They also found that hesperetin up-regulated the expression of total β -catenin in a dose-dependent manner and that hesperetin also caused nuclear translocation of β -catenin. In support of this they showed that suppression of Akt and/or β -catenin decreased the ameliorative activity of hesperetin against high glucose-suppressed osteogenic differentiation [83] (Fig. 2).

4.6. NF- κ B and Cyclooxygenase (COX)-2 Pathways

The NF- κ B family includes different transcription factors such as RELA (p65), c-BEL, RELB, NF- κ B1 (p50/p105), as well as NF- κ B2 (p52/p100) which plays key roles in the immune system and in inflammation as well as in regulation of proliferation and survival. A variety of stimuli, including reactive oxygen species, inflammatory agents, cytokines, carcinogenic materials, tumor inducers, bacterial and viral infections, γ -radiation, UV light, and X-rays, can activate members of this transcription factor family [84].

It has also been reported that tumor necrosis factor (TNF), interleukins, chemokines, COX-2, 5-lipoxygenase (LOX), as well as matrix metalloproteinase (MMP)-9 are restored via NF- κ B. In unstimulated cells, NF- κ B proteins are connected to inhibitory I κ B proteins, which preserve NF- κ B in an inactive form in the cytosol. TNF and Toll-like receptor activation leads to recruitment of adaptor proteins that activate the inhibitor of NF- κ B (I κ B) kinase, which phosphorylates I κ B. Following phosphorylation, I κ B is targeted by the proteasome for ubiquitination and degradation. Released NF- κ B translocates to the cell nucleus and results in expression of its target genes including cyclin D1, Bcl-2, Bcl-xL, MMP, as well as VEGF [60, 65].

COX is a prostaglandin H synthase, which is responsible for the production of prostanoids from arachidonic acid (AA). Three isoforms of COX, COX-1, COX-2, and COX-3, have been recognized. COX-1 mainly exists in most tissues; COX-2, however, is primarily present at sites of inflammation. Among these isoforms, COX-2 expression is normalized by mitogens, tumor cytokines, as well as growth factors. COX-2 expression, regulated by NF- κ B, mediates tumorigenesis [85]. It has been reported that COX-2 expression in human tumors can be triggered by different growth factors, cytokines, oncogenes, as well as other factors [86]. The anti-inflammatory effects of hesperidin are accomplished through various mechanisms including:

1. Down-regulation of intercellular adhesion molecule-1 expression which plays a crucial role in the inhibi-

tion of monocyte adhesion to endothelial cells [87, 88].

2. Suppression of expression of different pro-inflammatory cytokines and chemokines [89-91].
3. Increased production of nitric oxide and consequent improvement in endothelial function [87].
4. Inhibition of Janus kinase (JNK), a group of proteins activated by various types of environmental stress and cytokines, and consequent decreased production of metalloproteinases [92].

Moon and Kim have demonstrated the anti-inflammatory activity of hesperidin in HaCaT cells. Their results indicated that hesperidin inhibits interleukin-8 and tumor necrosis factor- α generation and also its mRNA expression, phosphorylation of p38 and COX-2 activation caused by hydrogen peroxide. They suggested that hesperidin suppressed NF- κ B activation, the phosphorylation of I κ B α and the p38 mitogen-activated protein kinase, and also COX-2 activation [76]. In another study, Kim and coworkers reported that hesperetin plays an important role in modulation of aging. They found that hesperetin inhibited the activation of NF- κ B and also related gene expressions. Indeed, another interesting point of this study was that hesperidin inhibited NF- κ B via different signal transduction pathways including NIK/IKK, ERK, p38, as well as JNK [78].

Previous studies have indicated that in colon malignancies, immunoreactivity of NF- κ B abruptly increased. Additionally, it has been demonstrated that NF- κ B expression in AOM-induced colonic tissues is significantly increased. Saiprasad and coworkers reported that hesperidin treatments significantly down-regulated the NF- κ B-dependent inflammatory responses including inducible nitric oxide synthase (iNOS) and COX-2 activation. Hesperidin supplementation inhibited NF- κ B expression at both the initiation and the post-initiation phases. Since increased NF- κ B expression activates downstream inflammatory molecules such as iNOS and COX-2, hesperidin significantly reduced these inflammatory markers [93]. Chronic inflammation is considered a risk factor for colorectal cancer. Hesperetin supplementation showed a suppression of expression of COX-2 mRNA and induction of apoptosis [94] (Fig. 2).

CONCLUSION

Hesperidin is one of the most common bioactive constituents of *C* species which possesses multiple health-promotion effects. This review has presented that hesperidin and its aglycone, hesperetin, have sup-

pressed the growth of various cancer cells, both *in vitro* and *in vivo* conditions, through different cellular pathways. Hesperidin primarily targets the NF- κ B pathway and also activates different apoptotic and pro-apoptotic pathways in different types of cancerous cells. However, our search in the <http://clinicaltrials.gov/> with keywords “hesperidin” and “hesperetin” in March 8, 2014 demonstrated that there are only 8 clinical trials regarding the impacts of hesperidin and its aglycone, hesperetin. In this regard, it is difficult to make a clear decision about its most effective doses for future clinical trials with respect to its anticancer activity. Finally, we conclude that hesperidin and its aglycone act as anticancer compounds by suppressing carcinogenesis through pleiotropic mechanisms.

CONFLICT OF INTEREST

The authors confirm that this article content has no conflict of interest.

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