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The effects of naringenin and naringin on the glucose uptake and AMPK phosphorylation in high glucose treated HepG2 cells

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ABSTRACT

Background: Naringin and its aglycone naringenin are citrus-derived flavonoids with several pharmacological effects. On the other hand, the mechanism for the anti-diabetic effects of naringenin and naringin are controversial and remain to be clarified further.

Objective: This study examined the relationship between glucose uptake and AMP-activated protein kinase (AMPK) phosphorylation by naringenin and naringin in high glucose-treated HepG2 cells.

Methods: Glucose uptake was measured using the 2-NBDG fluorescent D-glucose analog. The phosphorylation levels of AMPK and GSK3 β (Glycogen synthase kinase 3 beta) were observed by Western blotting. Molecular docking analysis was performed to evaluate the binding affinity of naringenin and naringin to the γ -subunit of AMPK.

Results: The treatment with naringenin and naringin stimulated glucose uptake regardless of insulin stimulation in high glucose-treated HepG2 cells. Both flavonoids increased glucose uptake by promoting the phosphorylation of AMPK at Thr172 and increased the phosphorylation of GSK3 β . Molecular docking analysis showed that both naringenin and naringin bind to the γ -subunit of AMPK with high binding affinities. In particular, naringin showed higher binding affinity than the true modulator, AMP with all three CBS domains (CBS1, 3, and 4) in the γ -subunit of AMPK. Therefore, both naringenin and naringin could be positive modulators of AMPK activation, which enhance glucose uptake regardless of insulin stimulation in high glucose-treated HepG2 cells.

Conclusions: The increased phosphorylation of AMPK at Thr172 by naringenin and naringin might enhance glucose uptake regardless of insulin stimulation in high glucose treated HepG2 cells.

Keywords: Naringenin; naringin; glucose uptake; AMPK phosphorylation; molecular docking

INTRODUCTION

AMPK activation is involved in non-insulin mediated glucose uptake, suggesting it is a potential strategy to enhance glucose uptake in an insulin-resistant state, such as Type 2 diabetes [1]. AMPK is activated by a low energy status (increased AMP/ADP:ATP) and regulates the metabolic process and energy homeostasis by triggering ATP-generating

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Conflict of Interest

The authors declare no conflicts of interest.

Authors Contributions

Conceptualization: Han CH, Dayarathne LA; Data curation: Ranaweera SS; Funding acquisition: Han CH; Investigation: Dayarathne LA, Han CH; Resources: Han CH; Software: Premkumar N, Rajan P; Supervision: Han CH; Visualization: Dayarathne LA, Han CH; Writing - original draft: Dayarathne LA; Writing - review & editing: Premkumar N, Rajan P.

processes, including glucose uptake and fatty acid oxidation [2]. The activation of AMPK regulates insulin-independent glucose uptake via stimulating GLUT2 translocation and regulating the systemic glucose homeostasis [3]. Once activated, AMPK phosphorylates GSK3, which is responsible for maintaining glucose homeostasis [4,5]. The phosphorylation and inhibition of GSK3 activate glycogen synthase (GS), which is a target for glycogen synthesis [6]. Therefore, AMPK is an important pharmaceutical target for the treatment of diabetes [1].

AMPK is inactive unless phosphorylated on the α -subunit activation loop at Thr172. On the other hand, AMP binding to the γ -subunit enhances Thr172 phosphorylation by upstream kinases, liver kinase B1 (LKB1), and Ca^{2+} /calmodulin-dependent protein kinase kinase (CaMKK) [7]. A previous study reported that flavonoids and other polyphenolic compounds are possible activators of AMPK, which have similar functions to adenosine nucleotides [8,9].

Naringin is a flavanone glycoside found in grapes and citrus fruits. Naringenin is formed by the attachment of two rhamnose units to the 7-carbon of naringin (**Fig. 1A**) [10]. For naringenin and naringin, diverse biological activities of therapeutic interests have been described, including anti-diabetic effects [11,12]. On the other hand, the mechanisms through which these flavonoids promote glucose uptake in the insulin-resistant state remains to be clarified.

The present study hypothesized that naringenin and naringin increase glucose uptake by activating the AMPK pathway in high glucose-treated HepG2 cells. For this, this study examined the effects of naringenin and naringin on the glucose uptake and the phosphorylation of AMPK and GSK3 β in high glucose-treated HepG2 cells. In addition,

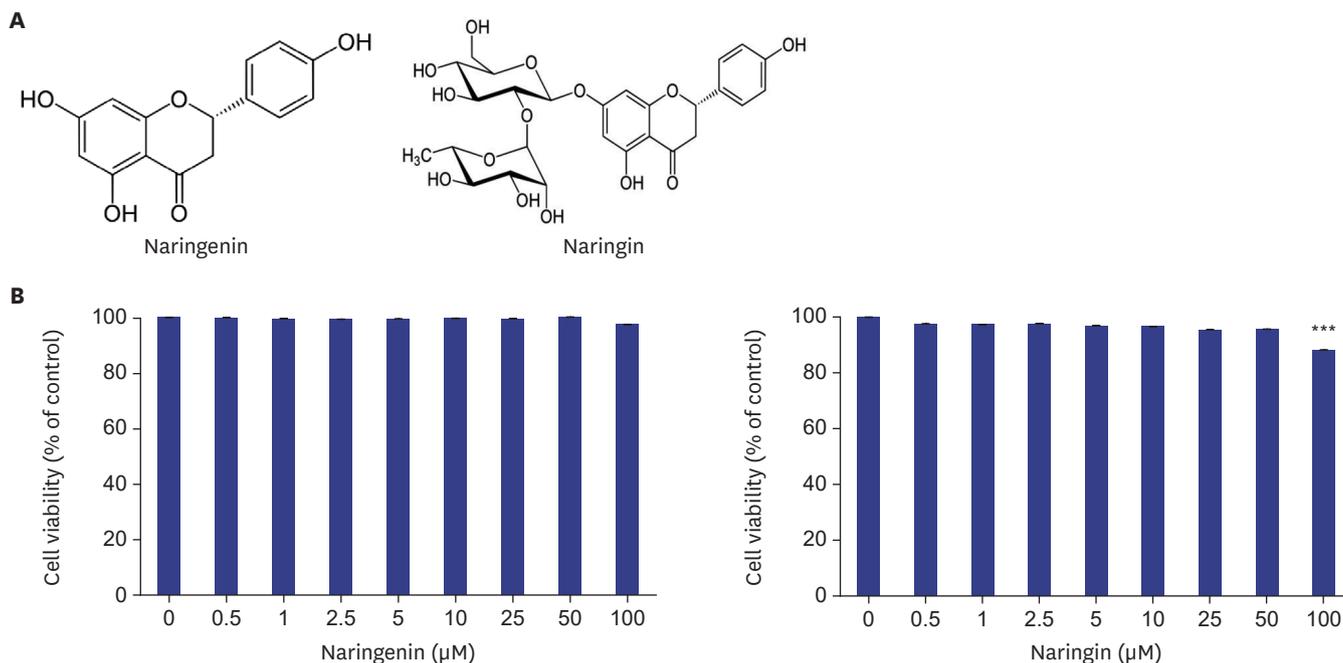


Fig. 1. Effect of naringenin and naringin on the viability of HepG2 cells. Chemical structures of naringenin and naringin (A). Effect of naringenin and naringin on the HepG2 cell viability. HepG2 cells were cultured at a density of 1×10^5 in a 96 well plate. After reaching confluence, the cells were treated with 0–100 μM of flavonoids for 24 h, and the cell viability was measured using a MTT assay (B). The values are the mean \pm SE.

*** $p < 0.0005$ vs. control.

molecular docking analysis was conducted to observe the binding affinity of naringenin and naringin to the γ -subunit of AMPK.

MATERIALS AND METHODS

Cell culture and treatment

HepG2 cells (KCLB 42707) were obtained from the Korean Cell Line Bank (KCLB) (Seoul, Korea) and cultured routinely in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% Fetal Bovine Serum (FBS), 100 U/mL penicillin, and 100 μ g/mL streptomycin (Gibco, USA) in a humidified atmosphere containing 5% CO₂ at 37°C. After reaching confluence, the cells were then seeded in culture plates for further experiments.

The insulin-resistant HepG2 cell model was established using the reported method with slight modifications [13]. Briefly, after seeding in 96 well plates, the cells were serum-starved for 12 h. and incubated in serum-free DMEM (Gibco, USA) containing either normal concentrations of glucose (5.5 mM D-glucose) or high concentrations of glucose (30 mM D-glucose) with or without the flavonoids (Sigma-Aldrich, USA) (10 and 50 μ M) and metformin (Sigma-Aldrich, USA) (2 mM) for an additional 24 h. The high glucose-treated cells were used as the insulin-resistant model. The cells were stimulated with or without 100 nM insulin (Sigma-Aldrich, USA) for 30 min before harvesting.

Cell viability assay

The HepG2 cells were seeded in 96-well plates at a cell density of 1×10^5 cells/well and cultured overnight using routine culture media. After reaching confluence, the cells were treated with different concentrations (0–100 μ M) of the flavonoids for 24 hrs. The cell culture medium was removed, and fresh medium containing 10% Ez-cytox (DogenBio, Korea) was added to each well, according to the manufacturer's instructions. The plates were incubated for 3 h at 37°C and 5% CO₂. The cell viability indicated by formazan production was measured using an ELISA microplate reader (TECAN, Austria) at 450 nm.

2-NBDG glucose uptake

HepG2 cells were cultured in 96-well plates. After reaching confluence, the cells were serum-starved for 12 h and incubated in a serum-free medium containing either normal or high glucose concentrations with or without the samples (flavonoids 10 μ M and 50 μ M) for an additional 24 h. The cells were incubated with 40 μ M 2-NBDG [2-N-(7-Nitrobenz-2-oxa-1,3-diazol-4-yl)amino-2-deoxy-d-glucose] (Invitrogen, Carlsbad, USA) with or without 100 nM insulin (Sigma-Aldrich, USA) for 30 min at 37°C. The cells were washed three times with PBS, and the fluorescent images were taken from IncuCyte ZOOM Fluorescent Microscope at 20 \times magnification (Essen BioScience, Inc. USA). IncuCyte ZOOM Fluorescent Processing Software was used to measure the total fluorescent intensities of each well.

Western blotting

The cells were washed with PBS (Gibco, USA) and lysed with ice-cold RIPA buffer (Tech and Innovation, Korea) containing protease inhibitor mixture. The whole-cell lysates were centrifuged at 12,000 rpm for 10 min. The supernatant was separated, and the amount of protein was assessed using a Bradford assay (Bio-Rad Laboratories, USA). Equal amounts of protein from the cell homogenates were subjected to SDS PAGE and transferred to PVDF membranes. The membranes were probed with the primary antibodies, GSK3 β , p-GSK3 β ,

AMPK, and p-AMPK (Cell signaling technology, USA), and detected with peroxidase-conjugated secondary antibodies for 1hr at room temperature. β -actin (ThermoFisher, USA) was used as a loading control. A chemiluminescence bioimaging instrument (NeoScience Co., Ltd., Korea) was used to detect the proteins of interest. Densitometry analysis was performed using ImageJ analysis software.

Molecular docking

The crystal structure of AMPK in complexes with AMP (PDB ID; 2V8Q) was obtained from Protein Data Bank (PDB). The sitemap tool (Schrodinger Software, Germany) was used to identify the four CBS domains (CBS1, CBS2, CBS3, and CBS4) in the γ -subunit of AMPK. All the AMP molecules in the γ -subunit of AMPK were removed for docking of the flavonoids. The 3D structures of AMP and flavonoids (naringenin and naringin) were downloaded from the PubChem database (<https://pubchem.ncbi.nlm.nih.gov/>) and minimized energetically using PyRx software (Python Prescription 0.8, The Scripps Research Institute). The grid box used for focused docking was set to $26 \times 44 \times 46 \text{ \AA}$ to ensure the structure of the γ -subunit of AMPK. The docking experiments were carried out using the AutoDock Vina module (Molecular Graphics Lab, The Scripps Research Institute, USA). The best-docked pose was selected based on binding energy, and 3D images were generated using the PyMOL (The PyMOL Molecular Graphics System, Ver.2.5.0, Schrodinger, LLC, USA). The docked complex of AMPK was optimized further, validated, and explored using the Discovery Studio visualizer (Ver.21.1.0.20298). The hydrogen bond and hydrophobic interactions between the receptor and ligand were analyzed using the Ligplot program [14].

Statistical analysis

The values are presented as the mean \pm SE for each group. Statistical analysis was performed using IBM SPSS Statistics (Ver.17.0; USA) and one-way analysis of variance with Tukey's post hoc tests for multi-group comparisons. The p values < 0.05 were considered significant.

RESULTS

Cell viability

The MTT assay was performed to assess the cellular toxicity of naringenin and naringin on HepG2 cells. No significant cellular toxicity was observed in HepG2 cells at flavonoid concentrations up to $50 \mu\text{M}$ (**Fig. 1B**). Accordingly, the experiments were conducted using the non-toxic concentrations of naringenin and naringin (10 and $50 \mu\text{M}$).

Glucose uptake

The effects of naringenin and naringin on the glucose uptake in high glucose-treated HepG2 cells were determined using a 2-NBDG uptake assay (**Fig. 2**). The glucose uptake was reduced in the high glucose-treated HepG2 cells compared to low glucose-treated cells. In contrast, the treatment of metformin (positive control) resulted in a significant increase ($p < 0.005$) the 2-NBDG uptake compared to the high glucose-treated HepG2 cells (**Fig. 2A and B**). In addition, the naringenin and naringin treatment increased the 2-NBDG glucose uptake in high glucose treated cells, regardless of insulin stimulation (**Fig. 2A and B**). In particular, the naringin treatment showed higher glucose uptake than the naringenin treatment in high glucose treated cells. The results suggest that naringenin and naringin enhanced the glucose uptake in high glucose-treated HepG2 cells regardless of insulin stimulation.

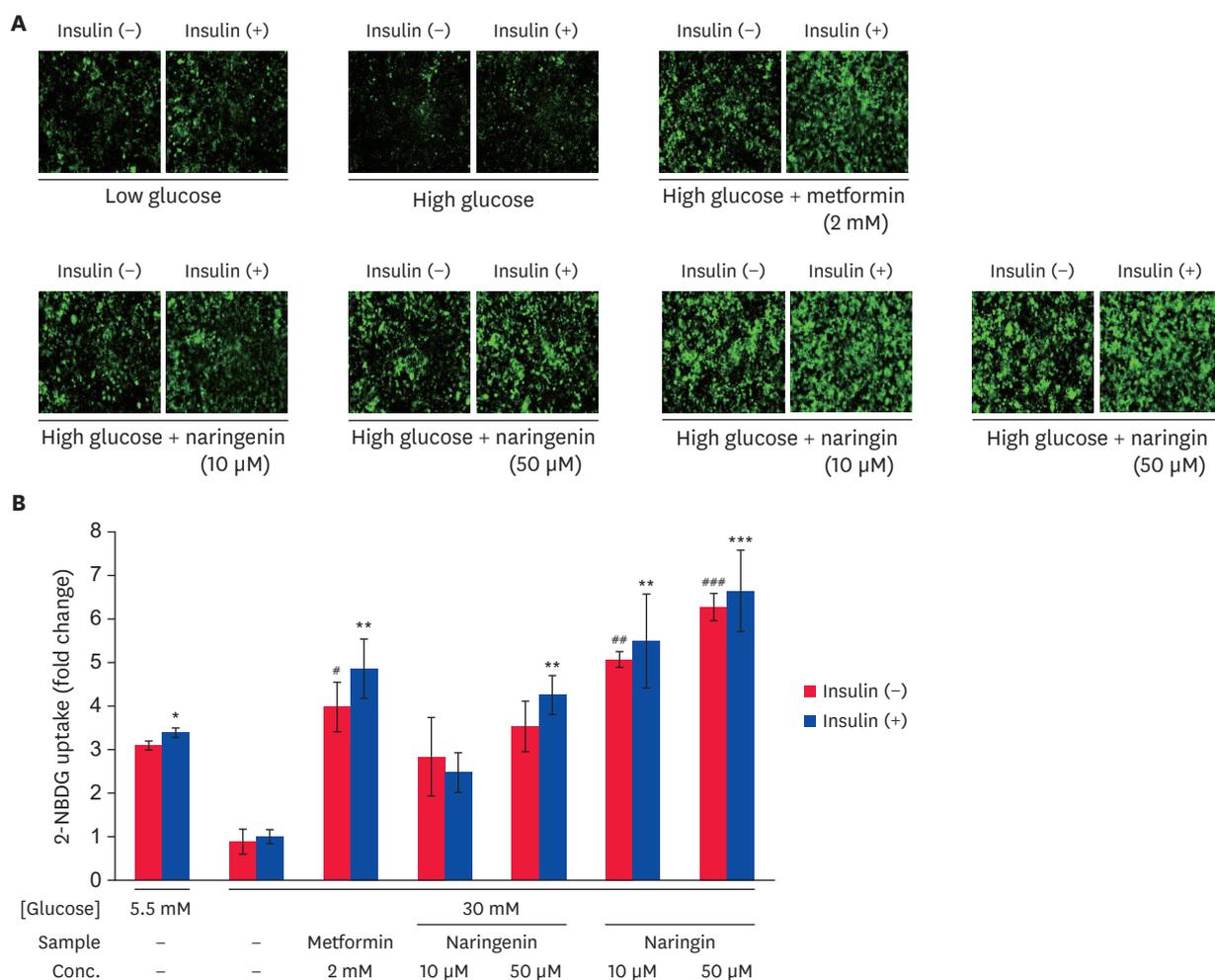


Fig. 2. Effect of naringenin and naringin on glucose uptake. The glucose uptake assay was done using the fluorescent D-glucose analog 2-NBDG. The HepG2 cells were serum-starved for 12 h and incubated in serum-free medium containing either normal (5.5 mM) or high glucose (30 mM) concentrations for an additional 24 h in the presence and absence of samples. The cells were then stimulated with or without insulin (100 nM) and 2-NBDG (50 μM) for 30 min. 2-NBDG uptake by cells was detected by IncuZyte Zoom at 20× magnification. Representative images for each sample (A). Quantitative 2NBDG uptake from each sample (B). The values are mean ± SE.

* $p < 0.05$, ** $p < 0.005$, and *** $p < 0.0005$ vs. insulin-stimulated high glucose control and * $p < 0.05$, ** $p < 0.005$, and *** $p < 0.0005$ vs. without insulin-stimulated high glucose control.

Phosphorylation of AMPK

Western blot analysis was conducted to observe the effects of naringenin and naringin on AMPK phosphorylation (Fig. 3). The levels of AMPK phosphorylation (Thr172) in high glucose-treated HepG2 cells were decreased significantly ($p < 0.05$) compared to low glucose-treated cells. In contrast, the level of AMPK phosphorylation (Thr172) was increased significantly by the metformin, naringenin, or naringin treatments in high glucose-treated HepG2 cells (Fig. 3A and B). The effects of naringin on the levels of AMPK phosphorylation (Thr172) were comparatively higher than the effect of naringenin. The results suggest that both naringenin and naringin increased the level of AMPK phosphorylation in high glucose-treated HepG2 cells.

Phosphorylation of GSK3β

Western blotting was performed to determine the effects of naringenin and naringin on the level of GSK3β phosphorylation (Ser9) (Fig. 4). The phosphorylation levels of GSK3β (Ser9)

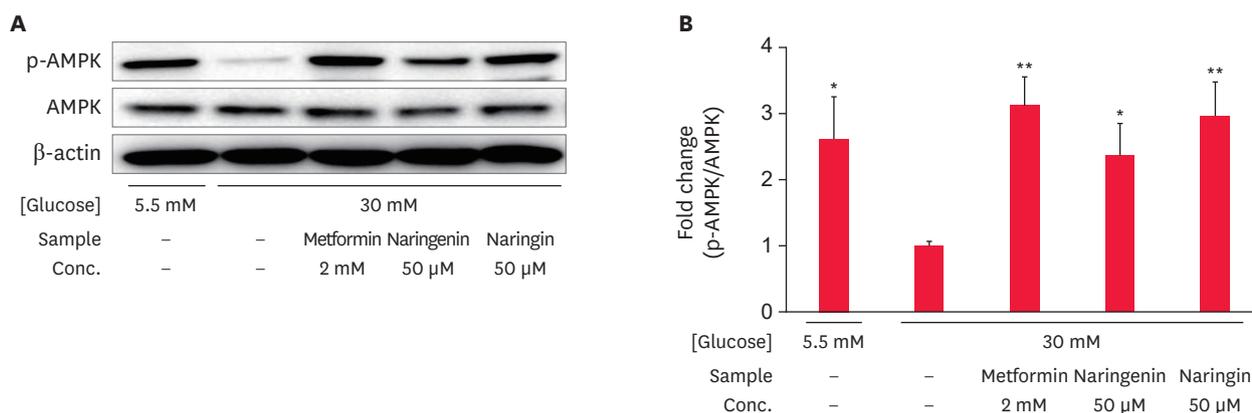


Fig. 3. Effect of naringenin and naringin on the phosphorylation of AMPK. HepG2 cells were starved in serum-free medium for 12 h and incubated in serum-free medium containing either normal (5.5 mM) or high (30 mM) glucose concentrations with or without different samples for an additional 24 h. The total cell extract was subjected to Western blot to observe the phosphorylation of AMPK. Immunoblot bands of AMPK (A), and the result of quantitative analysis of the ratio of p-AMPK to AMPK (B). Values are the mean \pm SE.

AMPK, AMP-activated protein kinase.

* $p < 0.05$ and ** $p < 0.05$ compared to high glucose control.

in high glucose treated HepG2 cells were decreased significantly ($p < 0.0005$) compared to the low glucose-treated cells. In contrast, the levels of GSK3 β phosphorylation were increased by the metformin, naringenin, or naringin treatment compared to high glucose-treated cells in either the presence or absence of insulin. In particular, the treatment of naringin showed significantly higher ($p < 0.0005$) levels of GSK3 β phosphorylation compared to naringenin treatment (Fig. 4A and B). These results suggest that both naringenin and naringin increased the levels of GSK3 β phosphorylation (Ser9) in high glucose-treated HepG2 cells regardless of insulin stimulation.

Molecular docking analysis

Molecular docking analysis was performed to simulate the binding affinity between the flavonoids and γ -subunit of AMPK (Fig. 5). As shown in Fig. 5A, the γ -subunit of AMPK contains four cystathionine β -synthase (CBS) domains. Among the four CBS domains, however, only three CBS (CBS1, CBS3, and CBS4) domains can bind to their modulator,

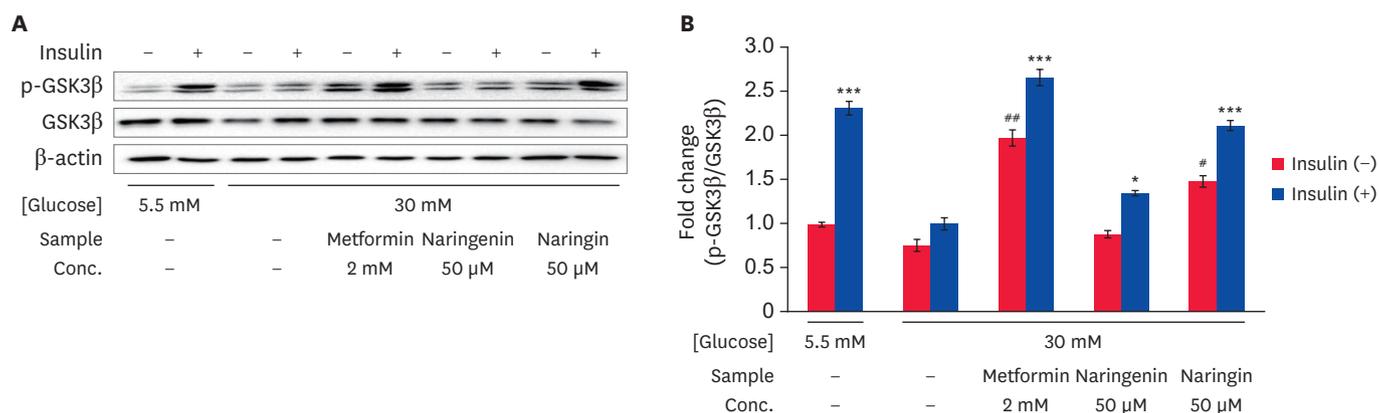


Fig. 4. Effect of naringenin and naringin on the phosphorylation of GSK3 β . HepG2 cells were starved in serum-free medium for 12 h and incubated in serum-free medium containing either normal (5.5 mM) or high (30 mM) glucose concentrations with or without different samples for an additional 24 h. Before harvesting, the cells were stimulated with or without 100 nM insulin for 30 min. The total cell extract was subjected to Western blot to observe the phosphorylation of GSK3 β . Immunoblot band pattern of GSK3 β (A), and the results of quantitative analysis of the ratio of p-GSK3 β to GSK3 β (B). The values are the mean \pm SE. * $p < 0.05$, *** $p < 0.0005$ compared with high glucose control and # $p < 0.05$ and ## $p < 0.005$ vs. without insulin-stimulated high glucose control.

Table 1. Binding energy of AMP and flavonoids at three sites that were identified in the γ -subunit after removing all the co-crystallized ligands

| Molecule | Description | Binding site of γ -subunit of AMPK | | |
|------------|---------------------------|--|--|--|
| | | CBS1 | CBS3 | CBS4 |
| AMP | Binding energy (kcal/mol) | -7.8 | -7.4 | -8.1 |
| | Hydrogen bonds | D89, R151, T88, T86, K148, H150 | A294, R69, R298, R268, D244, H297, K169, S241 | N202, S225, S313, S315, H150, H297 |
| Naringenin | Hydrophobic interactions | M84, I149, I87 | L276, I239, F243, V296 | I203, V224, I311, T199, K148 |
| | Binding energy (kcal/mol) | -7.7 | -8.0 | -7.6 |
| | Hydrogen bonds | R117, H150, V129 | R298 | A226, R298, T199, S225 |
| Naringin | Hydrophobic interactions | D89, I149, L128, G83, T86, T88, M84 | R268, I239, H270, F243, F272, L276, V275, V296 | S315, V224, I311 |
| | Binding energy (kcal/mol) | -8.5 | -8.3 | -9.1 |
| | Hydrogen bonds | R151, K242, K148 | R298, R268, R69, D244, G273, G247, L276 | R298, N202, A201, H297, S313, S225, T99 |
| Naringin | Hydrophobic interactions | R223, R117, D89, R69, S225, T88, T86, W116, M84, H150, I87, I149, L121 | I239, S241, F243, V296, K169, G295, F272, V275 | A226, R223, D316, I311, I203, H150, K148, S315, V224 |

DISCUSSION

This study examined the effects of naringenin and naringin on glucose uptake and AMPK phosphorylation in high glucose-treated HepG2 cells. The naringenin and naringin treatment showed increased glucose uptake with increased AMPK phosphorylation (Thr172) and GSK3 β . Molecular docking simulation showed that both naringenin and naringin bind to the cavity of the γ -subunit of AMPK with high binding affinities, suggesting that the flavonoids could be positive modulators for AMPK activation.

Naringenin and naringin enhanced the glucose uptake significantly regardless of the insulin stimulation in high glucose treated HepG2 cells, as observed previously [15]. Hepatic glucose uptake is largely an insulin-independent process, which contributes to whole-body glucose homeostasis [16]. For example, hepatic glucose uptake can be enhanced by the entry of glucose into the portal vein, which stimulates a portal glucose signal [17]. When hyperglycemia occurs, plasma glucose enters the cytoplasm of the hepatocytes through GLUT2, which is an insulin-independent system [18]. In this process, AMPK suppresses gluconeogenesis in the liver and promotes glucose uptake in peripheral tissues [19].

In the liver, AMPK enhances glucose uptake by up-regulating the expression of GLUT2 [18]. In addition, the activation of AMPK suppresses hepatic gluconeogenesis and promotes glycogen synthesis by the direct phosphorylation of its substrates, including GSK3 β , which will ultimately increase the glucose uptake by the liver [20,21]. The AMPK-induced phosphorylation of GSK3 β inhibited the transcriptional activity of the cAMP response element-binding protein (CREB), a key transcription factor, which regulates the phosphorylation of gluconeogenic enzymes [20].

AMPK is a heterotrimer consisting of catalytic α subunits and regulatory β and γ subunits [22]. The phosphorylation of Thr172 within the activation loop of the α -subunit can cause AMPK activation [7]. AMP is a natural activator of AMPK and binds to the CBS domains of the γ -subunit and indirectly promotes the activity of the catalytic domain in α -subunit [23]. The binding of AMP to the γ -subunit of AMPK stimulates the phosphorylation of Thr172 in the α -subunit by upstream kinases LKB1 and CaMKK β , and prevents the dephosphorylation of Thr172 in α -subunit [24,25]. In addition, a recent study suggested that transforming

growth factor β -activated kinase (TAK1) phosphorylates AMPK on Thr172 through AMP binding [26]. Therefore, AMP binding to CBS causes allosteric activation subsequent to the phosphorylation of Thr172 in the activation loop of the α -subunit, which results in AMPK activation [25].

The molecular docking results suggest that both naringenin and naringin bind to the cavity of γ -subunit of AMPK with high binding affinities. AMP and flavonoids interacted with the R-groups of amino acids in CBS domains 1, 3, and 4 of the γ -subunit. In particular, naringin showed higher binding affinities than AMP in the CBS domains 1, 3, and 4 of the γ -subunit, where naringin forms a larger number of hydrophobic interactions or hydrogen bonds than true modulator, AMP. The results suggest that naringin could be a potent positive modulator for activating AMPK. In addition, both naringenin and naringin enhanced AMPK phosphorylation at Thr172, which might enhance glucose uptake by stimulating GLUT2 translocation in HepG2 cells. These results suggest that both naringenin and naringin could positively modulate AMPK activation, enhancing glucose uptake.

As reported previously, flavonoid glucuronic acids had a beneficial effect on glucose homeostasis with high binding affinities to AMPK [27]. A recent study reported that *Lippia citriodora*-derived polyphenols act as direct agonists of AMPK by binding to the AMP binding sites of the γ -subunit and the interaction zones between the γ and β -subunits [28]. An *in silico* study on Chinese medicinal compounds eugenyl β -D-glucopyranoside and 6-O-cinnamoyl-D-glucopyranose revealed similar functions as AMP for AMPK activation [8]. Overall, the molecular docking analysis of the present study confirmed that both naringenin and naringin could be positive modulators of AMPK activation.

Naringin has a higher anti-diabetic potential than naringenin. In general, glycosides have higher biological activities than their aglycones because the bound sugar modifies the hydrophilicity of glycosides, enhancing the bioavailability of the compounds [29,30]. On the other hand, the glycoside linkages are unstable under the acidic conditions in the stomach. In addition, the cleavage of sugar moieties of naringin by glycosidase from intestinal bacteria will produce the aglycone, naringenin [31]. Therefore, the intestinal absorption of naringenin should be clarified [32]. In the present study, however, naringin, being a glycoside, had a higher anti-diabetic effect than its aglycone naringenin on HepG2 cells.

Molecular docking results showed that both naringenin and naringin are potential AMPK activators with high binding affinities to the γ -subunit of AMPK. These findings suggest that the glucose uptake by naringenin and naringin depends on the phosphorylation of AMPK at Thr172, which will enhance glucose uptake by stimulating GLUT2 translocation in HepG2 cells (**Fig. 6**). Overall, the results suggest that both naringenin and naringin could be positive modulators of AMPK activation, enhancing glucose uptake regardless of insulin stimulation in high glucose-treated HepG2 cells. Nevertheless, further efforts will be needed to define the precise role of naringenin and naringin on the glucose uptake underlying the AMPK pathway in HepG2 cells.

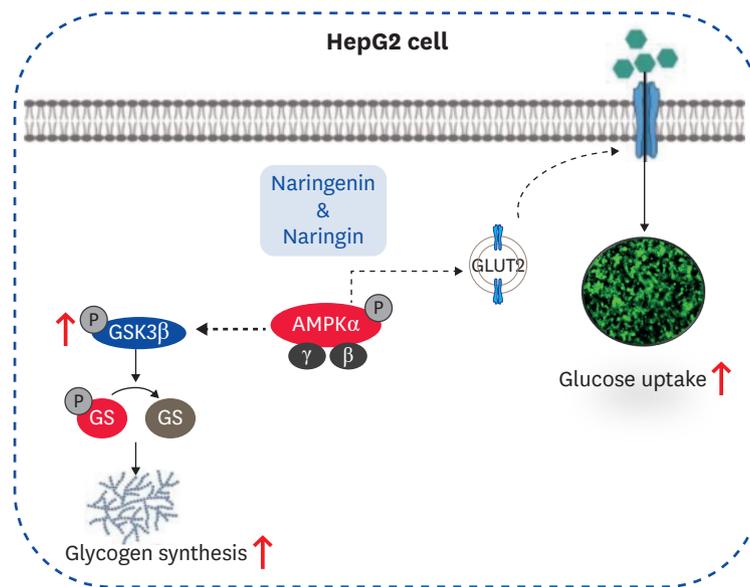


Fig. 6. Proposed model for the enhanced glucose uptake effect of naringenin and naringin on HepG2 cells. AMPK, AMP-activated protein kinase.

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