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Nectandrin B, a lignan isolated from nutmeg, inhibits liver X receptor- α -induced hepatic lipogenesis through AMP-activated protein kinase activation

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Nonalcoholic fatty liver disease is recognized as the most commonly occurring chronic liver disease. Liver X receptor α (LXR α) and sterol regulatory element-binding protein (SREBP)-1c play a central role in *de novo* fatty acid synthesis. This study investigated pharmacological effects of nectandrin B, a lignan isolated from nutmeg extract, on hepatic lipogenesis stimulated by LXR α -SREBP-1c-mediated pathway and the possible molecular basis. The reporter gene assay revealed that nectandrin B completely represses LXR α activity enhanced by a synthetic LXR α ligand (T0901317) in HepG2 cells. The inhibitory effect was further supported by the suppression of mRNA expression of LXR α target genes, SREBP-1c and LXR α itself. Nectandrin B also inhibited the increase in SREBP-1c expression promoted by insulin plus high glucose, major contributors to hepatic lipid accumulation. LXR α -SREBP-1c-mediated induction of acetyl-CoA carboxylase 1 and fatty acid synthase, major genes for *de novo* lipogenesis, was suppressed by nectandrin B. Moreover, Oil Red O staining showed that nectandrin B notably attenuates LXR α -induced lipid accumulation. AMP-activated protein kinase (AMPK) inhibits the activities of LXR α and SREBP-1c. Nectandrin B strongly activated AMPK signaling in HepG2 cells. Taken together, the suppressive effects of nectandrin B on lipogenic gene expression and lipid accumulation in hepatocytes may be due to its inhibitory effect on the LXR α -SREBP-1c pathway presumably *via* AMPK activation. These results suggest the potential of nectandrin B as a therapeutic candidate for fatty liver disease.

1. Introduction

Nonalcoholic fatty liver disease (NAFLD) is recognized as the most commonly occurring chronic liver disease (Asrih and Jornayvaz 2015). Dysregulation of hepatic fatty acid metabolism can cause excess lipid accumulation in hepatocytes, a major characteristic of NAFLD (Fabbrini et al. 2010). A previous study in patients with NAFLD demonstrated that about 26% of fatty acids in triglycerides produced in liver originate from *de novo* hepatic fatty acid synthesis, suggesting the significant contribution of *de novo* lipogenesis pathway to hepatic triglyceride accumulation (Donnelly et al. 2005). Furthermore, hepatic expression of major lipogenic genes, such as acetyl-CoA carboxylase (ACC) 1 and fatty acid synthase (FAS) is enhanced in parallel with the increase in their key transcription factors, liver X receptor α (LXR α) and sterol regulatory element binding protein (SREBP)-1c in NAFLD patients (Higuchi et al. 2008). LXR α is a nuclear receptor activated by endogenous oxysterols and serves as a crucial regulator of cholesterol, fatty acid, and glucose metabolism (Calkin and Tontonoz 2012; Jakobsson et al. 2012). LXR α plays an essential role in the regulation of hepatic *de novo* fatty acid synthesis through the transacti-

vation of SREBP-1c, a master transcription factor for the major genes involved in *de novo* lipogenesis (Schultz et al. 2000; Repa et al. 2000; Shimomura et al. 1998). Moreover, LXR α increases FAS gene expression by directly binding to its promoter region (Joseph et al. 2002). Interestingly, LXR α is required for insulin-stimulated SREBP-1c gene transcription and fatty acid synthesis (Chen et al. 2004).

AMP-activated protein kinase (AMPK) as a critical regulator of glucose and lipid metabolism, reduces the synthesis of fatty acid and cholesterol in the liver, but promotes fatty acid oxidation in skeletal muscle (Lage et al. 2008). ACC1, a rate-limiting enzyme in hepatic fatty acid synthesis pathway is directly inactivated *via* AMPK-mediated phosphorylation (Lage et al. 2008). Furthermore, AMPK suppresses the expression of lipogenic genes by regulating the transcriptional activities of LXR α and SREBP-1c through direct phosphorylation (Hwahng et al. 2009; Li et al. 2011).

Nutmeg, the dried semen of *Myristica fragrans* Houtt. (Myristicaceae) is a popularly used spice and has also been used as a traditional medicine for the treatment of diarrhea (Nguyen et al. 2010; Van Gils and Cox 1994; Grover et al. 2002). Several pharmacological activities of nutmeg extracts, such as

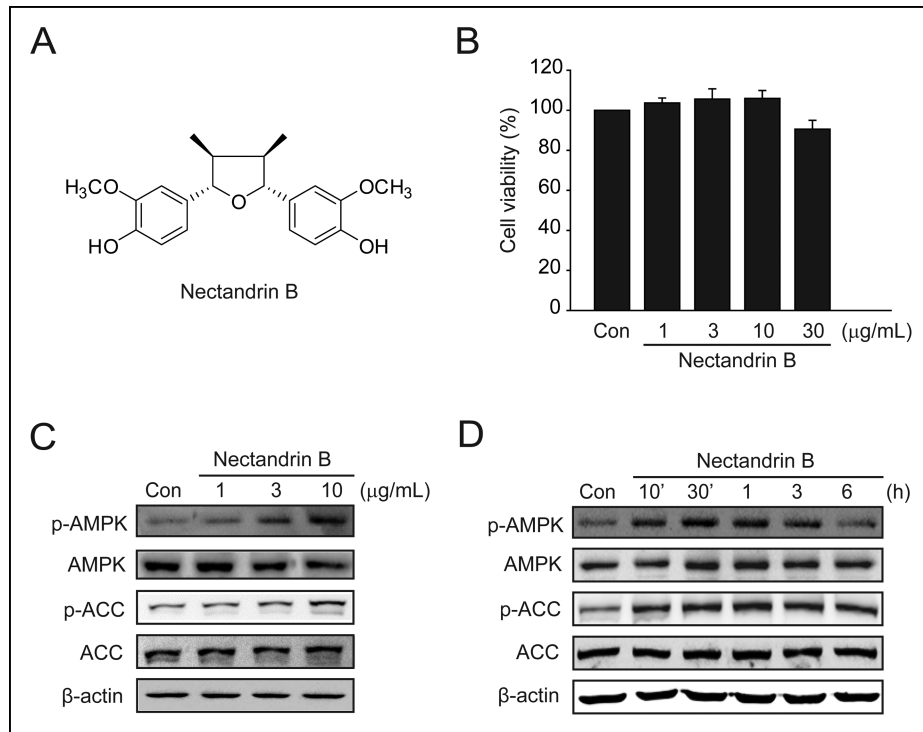


Fig. 1: Effect of nectandrin B on cell viability and AMPK signaling in HepG2 cells. (A) Chemical structure of nectandrin B. (B) Cells were incubated with the indicated concentrations of nectandrin B for 24 h, and then the cell viability was assessed by MTT assay. Data represent the mean \pm S.E. (n = 4). (C-D) Cells were exposed to the indicated concentrations of nectandrin B for 1 h or treated with nectandrin B (10 μ g/mL) for the indicated times. Cell lysates were obtained and each protein level was determined by Western blot analysis.

anti-inflammatory, anti-hyperlipidemic, and anti-atherosclerotic effects have been reported (Olajide et al. 1999; Ram et al. 1996; Sharma et al. 1995). In a previous study, a total extract of nutmeg was shown to activate AMPK and seven tetrahydrofuran-type lignans were isolated from this extract as active constituents (Nguyen et al. 2010). Among those lignans, nectandrin B exhibits a strong activating effect on AMPK signaling in differentiated C2C12 cells (Nguyen et al. 2010).

In the present study, we investigated the pharmacological effects of nectandrin B on hepatic lipogenesis. We determined if nectandrin B activates AMPK signaling pathway in hepatocytes. We also evaluated its ability to inhibit the transcriptional activity of LXR α and down-regulate LXR α -mediated lipogenic gene expression. Furthermore, we report that nectandrin B attenuates lipid accumulation stimulated by an LXR α ligand in hepatocytes.

2. Investigations and results

2.1. Effect of nectandrin B on HepG2 cell viability

An MTT colorimetric assay was used to examine the cytotoxicity of nectandrin B in HepG2 cells. Cell viability tended to slightly decrease at a concentration of 30 μ g/mL (Fig. 1B). Thus, 1-10 μ g/mL of nectandrin B was treated for subsequent experiments.

2.2. Activation of AMPK signaling by nectandrin B in HepG2 cells

Many studies have demonstrated the critical roles of AMPK in the regulation of lipid metabolism, particularly the inhibitory effects on LXR α and SREBP-1c activities (Lage et al. 2008; Hwang et al. 2009; Li et al. 2011). AMPK was notably activated in response to the treatment of nectandrin B in HepG2 cells as evidenced by the increase in the phosphorylation of

AMPK at Thr172 (Fig. 1C and 1D). This result was also confirmed by the increased phosphorylation of ACC, a downstream substrate of AMPK. Nectandrin B enhanced the phosphorylation of AMPK and ACC in a concentration-dependent manner (Fig. 1C). The concentration of 10 μ g/mL of nectandrin B capable of strongly activating AMPK was applied to the subsequent experiments unless specified otherwise. The levels of phosphorylated forms of AMPK and ACC began to increase from 10 min after treatment of 10 μ g/mL of nectandrin B and the effects were maintained up to at least 3 h. These results demonstrate that nectandrin B can activate AMPK in hepatocytes, implying the possibility of its inhibitory action on hepatic lipogenesis.

2.3. Preventive effect of nectandrin B on hepatic SREBP-1c induction

SREBP-1c serves as a common transcription factor for the induction of major lipogenic enzymes, thus leading to hepatic steatosis (Shimomura et al. 1998; Moon et al. 2012). LXR α plays a central role in the transcriptional regulation of SREBP-1c (Repa et al. 2000). We therefore used T0901317 (a synthetic LXR α ligand) to induce SREBP-1c expression and hepatic lipogenesis in the present study.

In order to investigate the ability of nectandrin B to prevent hepatic lipogenesis, we first determined the effect of nectandrin B on the SREBP-1c induction stimulated by T0901317. Nectandrin B abrogated T0901317-induced SREBP-1c up-regulation at the concentration of 10 μ g/mL (Fig. 2A). The inhibitory effect of nectandrin B on the LXR α -dependent SREBP-1c induction was confirmed by the study using GW3965, a more specific LXR α ligand (Fig. 2B). Moreover, we observed the ability of nectandrin B to attenuate the notable increase in SREBP-1c protein expression after exposure to insulin plus high glucose, major contributors to the pathogenesis of hepatic lipid accumulation in metabolic syndromes (Fig. 2B). Our results suggest that nectan-

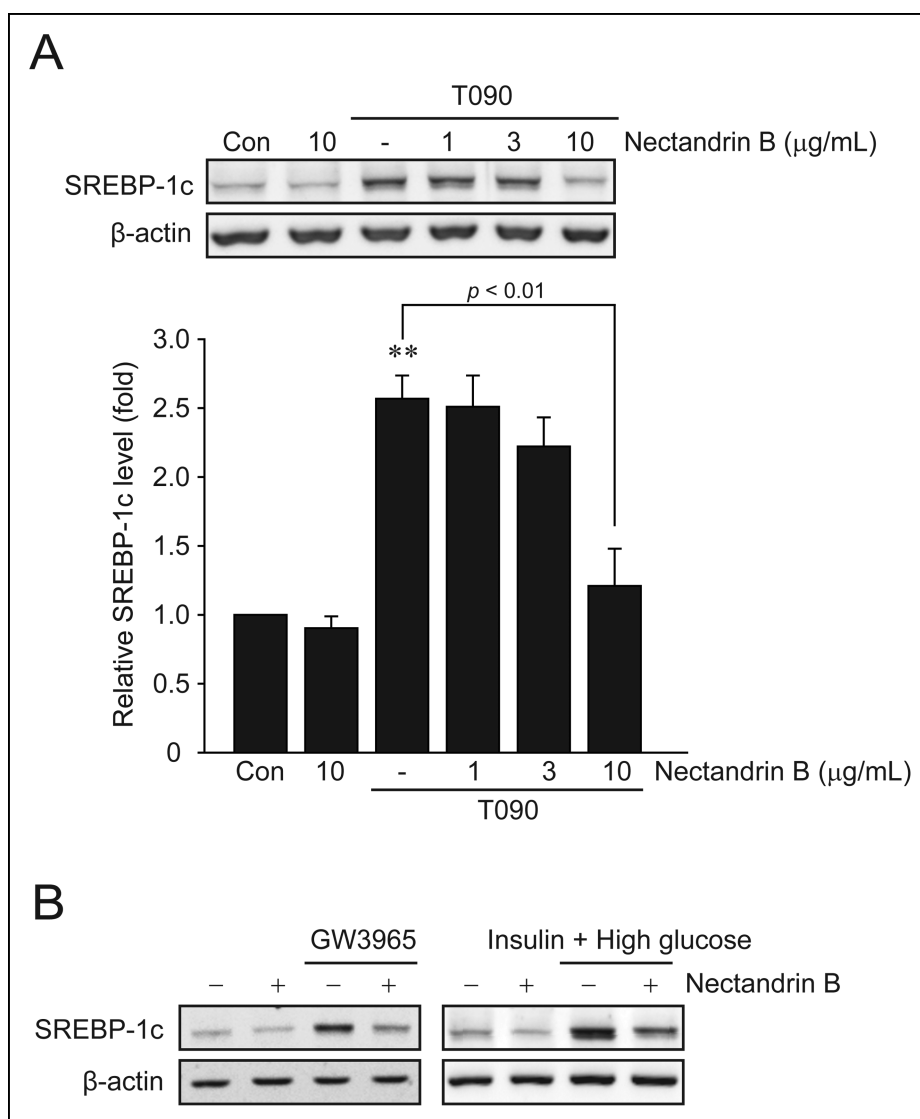


Fig. 2: Effect of nectandrin B on SREBP-1c induction by LXR α agonists or insulin plus high glucose in HepG2 cells. (A) Cells were pretreated with vehicle or nectandrin B (1–10 μ g/mL) for 1 h and subsequently exposed to T0901317 (T090, 10 μ M) for 12 h. The cell lysates were then immunoblotted and band intensity was determined by scanning densitometry. The SREBP-1c levels were normalized to those of β -actin. Data represent the mean \pm S.E. (n = 5). ** $P < 0.01$ (compared with the vehicle-treated control) (B) Cells were pretreated with nectandrin B (10 μ g/mL) for 1 h and then incubated with GW3965 (10 μ M) for 12 h, or insulin (100 nM) plus high glucose (25 mM) for 24 h. The SREBP-1c protein levels were measured in the cell lysates.

drin B can block the LXR α -dependent transcriptional induction of SREBP-1c.

2.4. Inhibitory effect of nectandrin B on LXR α transcriptional activity and LXR α -mediated lipogenic gene induction

We next measured the LXRE luciferase activities to verify the inhibitory effect of nectandrin B on LXR α transactivation. The pTK-CYP7A1-LXRE(x3)-LUC reporter gene construct including three copies of an LXRE sequence in rat *Cyp7a1* gene was used for this study. The reporter gene assay revealed that nectandrin B significantly attenuates the ability of T0901317 to enhance the CYP7A1-LXRE-luciferase activity (Fig. 3A). Multiple LXREs are found in human LXR α gene promoter and LXR α can autoregulate its own promoter (Laffitte et al. 2001). Accordingly, the mRNA expression of LXR α was increased by 12 h exposure of HepG2 cells to T0901317, and pretreatment with nectandrin B completely abrogated this increase (Fig. 3B left). Similarly, nectandrin B repressed the T0901317-induced mRNA expression of SREBP-1c, another LXR α target gene

(Fig. 3B right). Both FAS and ACC1, main enzymes for *de novo* fatty acid synthesis, are regulated by LXR α and SREBP-1c (Joseph et al. 2002; Lopez et al. 1996). As expected, nectandrin B significantly reduced the LXR α -SREBP-1c-mediated induction of FAS and ACC1 (Fig. 3C).

Taken together, these results demonstrate that nectandrin B inhibits the LXR α activity and subsequently suppresses the increase in hepatic lipogenic gene expression, implying its possible beneficial effect on hepatic lipid accumulation.

2.5. Attenuation of LXR α -induced lipid accumulation by nectandrin B in primary mouse hepatocytes

We further determined whether nectandrin B exerts anti-lipogenic effects in primary hepatocytes isolated from mice. In consistence with the results obtained in HepG2 cells, the inhibitory effect of nectandrin B on the T0901317-induced SREBP-1c protein expression was confirmed in primary cultured mouse hepatocytes (Fig. 4A). Next, we examined the effect of nectandrin B on the T0901317-stimulated lipid accumulation. The primary cultured mouse hepatocytes were stained

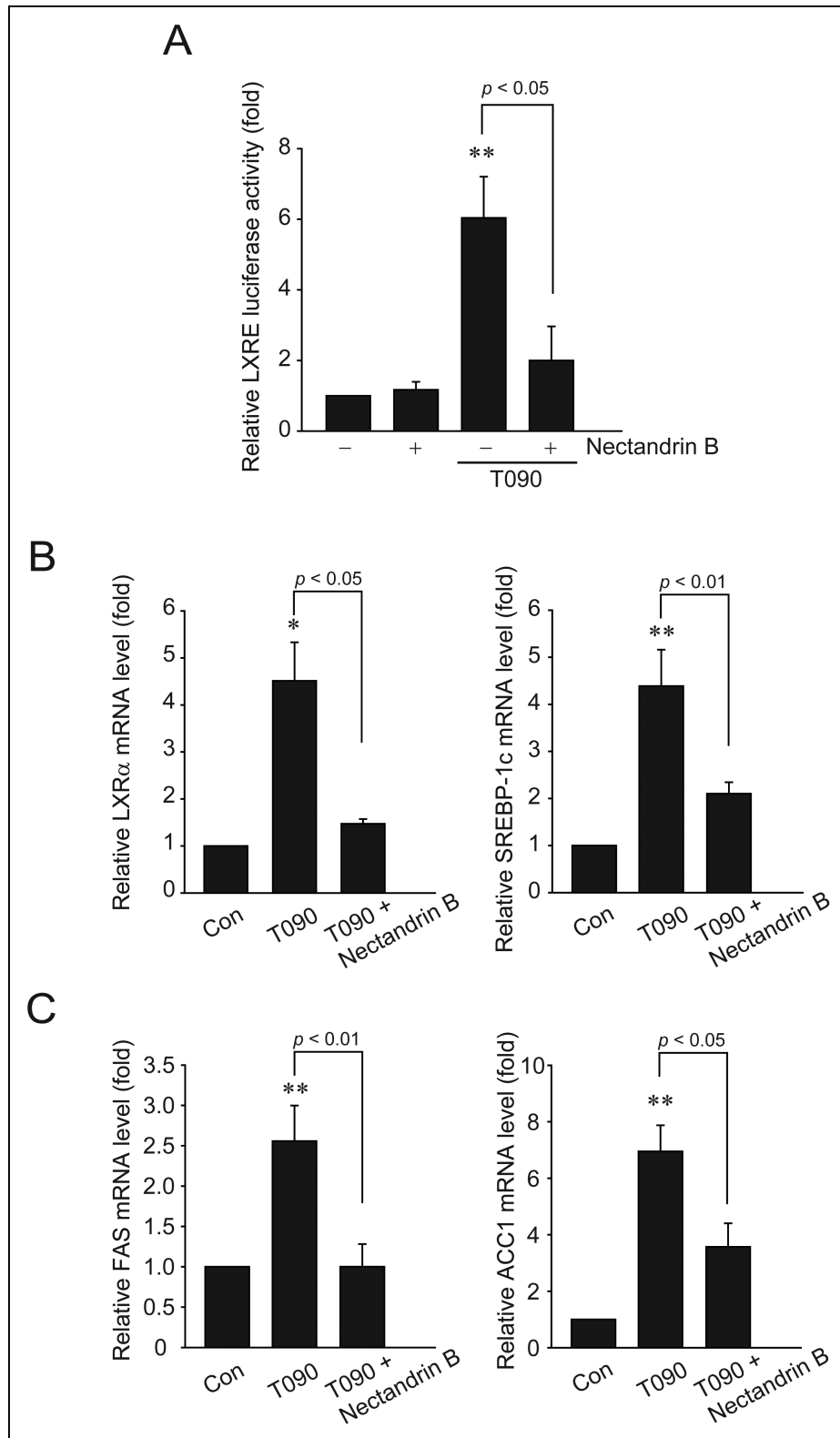


Fig. 3: Effect of nectandrin B on LXR α transcriptional activity and LXR α -mediated lipogenic gene induction in HepG2 cells. (A) Cells were transfected with an LXRE-luciferase construct and then exposed to T0901317 (T090, 10 μ M) for 12 h after 1 h pretreatment with vehicle or nectandrin B (10 μ g/mL). The relative LXRE luciferase activity was measured in the cell lysates. Data represent the mean \pm S.E. (n=4). ** P <0.01 (compared with the vehicle-treated control) (B-C) Cells were treated with vehicle or 10 μ g/mL nectandrin B for 1 h and subsequently exposed to T0901317 (T090, 10 μ M) for 12 h. Each mRNA level was analyzed by using real-time PCR assay and normalized to that of GAPDH. Data represent the mean \pm S.E. (n=3-6). * P <0.05, ** P <0.01 (compared with the vehicle-treated control).

with Oil Red O to visualize intracellular lipid droplets. Oil Red O staining revealed that LXR α -induced accumulation of lipid droplets was notably attenuated by the pretreatment of nectandrin B (Fig. 4B). These results indicate that nectandrin B has anti-steatotic effects through the inhibition of LXR α -mediated lipogenic gene induction in hepatocytes.

3. Discussion

The liver is a primary organ in regulating carbohydrate and lipid metabolism to maintain whole-body energy homeostasis (Fabbrini et al. 2010). Nonalcoholic simple fatty liver can progress to nonalcoholic steatohepatitis (NASH), a more severe condi-

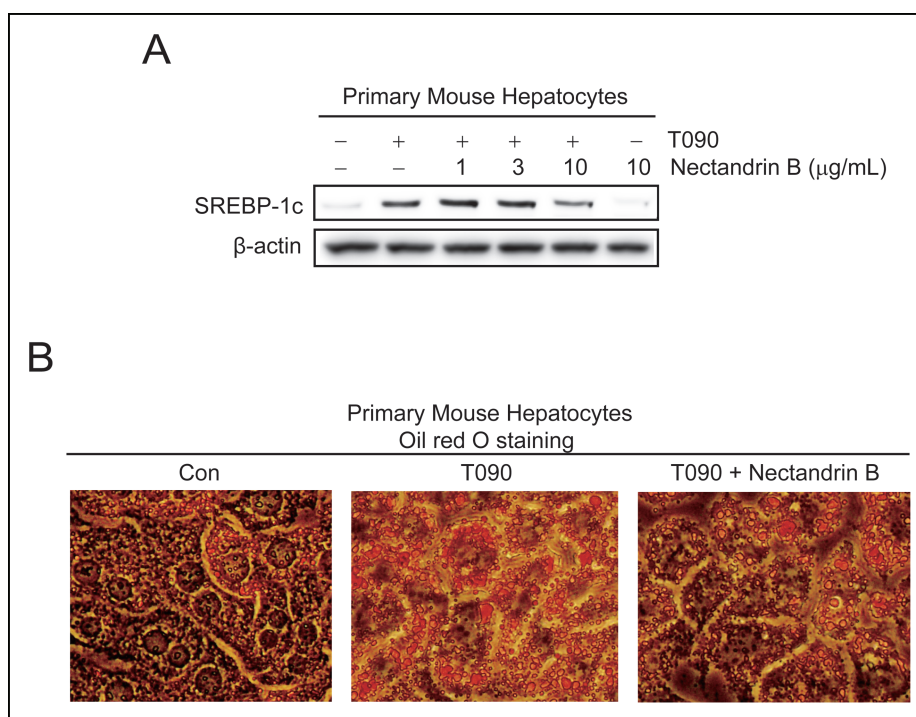


Fig. 4: Effect of nectandrin B on LXR α -induced SREBP-1c expression and lipid accumulation in primary mouse hepatocytes. (A) Cells were treated as described in the legend to Fig. 2A. (B) Cells were incubated with T0901317 (T090, 10 μ M) for 24 h in the presence or absence of 10 μ g/mL nectandrin B. Lipid droplets were detected by staining with Oil Red O. Representative images are shown.

tion with inflammation and fibrosis that elevates the risk of liver cirrhosis, an irreversible liver disease (Contos et al. 2004). Furthermore, NAFLD is referred to as hepatic manifestation of metabolic syndrome which is defined as the cluster of metabolic abnormalities including abdominal obesity, hyperglycemia, dyslipidemia, and hypertension (Asrih and Jornayvaz 2015; Alberti et al. 2005). Phytochemicals in plants are receiving growing attention because of their therapeutic potential for metabolic diseases (Soory 2009).

Nectandrin B is a tetrahydrofuran-type lignan isolated from nutmeg extract (Nguyen et al. 2010). In previous studies, nectandrin B has been shown to strongly activate AMPK, thus leading to vasorelaxation due to phosphorylation of endothelial nitric-oxide synthase, inhibition of neointima formation, and suppression of monocyte adhesion to endothelial cells (Hien et al. 2011; Ki et al. 2013; Hien et al. 2014). AMPK has been recognized as an effective therapeutic target for metabolic disorders including fatty liver, insulin resistance, and type 2 diabetes (Hardie 2011). It has been reported that the mixture of seven tetrahydrofuran-type lignans of nutmeg extract containing nectandrin B prevents the increases in body weight, epididymal fat mass, and the levels of blood glucose and LDL cholesterol in mice fed with high-fat diet (Nguyen et al. 2010). However, the molecular mechanisms for these anti-obesity and hypolipidemic effects have not been elucidated. We report here, for the first time, the protective effects of nectandrin B on the enhanced *de novo* fatty acid synthesis pathway and resultant lipid accumulation in hepatocytes.

In view of the important role of LXR α -SREBP-1c pathway in *de novo* lipogenesis and pathogenesis of fatty liver diseases, targeting this pathway would be therapeutically attractive for the treatment of NAFLD. In the present study, we demonstrated that nectandrin B completely represses the transcriptional activity of LXR α enhanced by a synthetic LXR α agonist. This inhibitory effect of nectandrin B was further supported by the suppression of mRNA expression of LXR α target genes, such as SREBP-1c and LXR α itself. Furthermore, we showed that

nectandrin B also inhibits the increase in SREBP-1c expression promoted by insulin plus high glucose, two key endogenous stimuli for hepatic fatty acid synthesis. Several mechanisms have been identified to explain the activation of lipogenic pathway by either insulin or glucose, but commonly involve LXR α and SREBP-1c. Both insulin and high glucose activate SREBP-1c by stimulating its proteolytic cleavage and also increase the protein level of the precursor form of SREBP-1c (Hegarty et al. 2005; Guillet-Deniau et al. 2004). Moreover, LXR α is necessary for insulin-stimulated SREBP-1c induction (Chen et al. 2004) and glucose can act as an endogenous ligand for LXR α (Mitro et al. 2007). Collectively, we demonstrate here that nectandrin B has the ability to repress the induction of SREBP-1c precursor protein, presumably via the inhibition of LXR α activity.

The LXR α transcriptional activity is regulated by several kinases including protein kinase A, AMPK, p70 ribosomal S6 kinase-1, and c-Jun N-terminal kinase 1 (Yamamoto et al. 2007; Hwahng et al. 2009; Kim et al. 2010). AMPK suppresses LXR α activity by direct threonine-phosphorylation as demonstrated *in vitro* and in cell kinase assays (Hwahng et al. 2009). Several other mechanisms also contribute to the anti-lipogenic effects of AMPK (Lage et al. 2008). AMPK activation inactivates ACC1 by direct phosphorylation (Lage et al. 2008). Additionally, a recent paper suggested that AMPK directly phosphorylates SREBP-1c at Ser372, thus blocking the transcriptional activation of SREBP-1c target genes through the inhibition of SREBP-1c cleavage (Li et al. 2011). Taken together, the repression of LXR α -SREBP-1c pathway by nectandrin B can be explained at least in part by its stimulatory effect on AMPK activation. Finally, the suppressive effects of nectandrin B on the expression of major lipogenic genes, FAS and ACC1, and resultant fat accumulation in hepatocytes may be due to its inhibitory effect on the LXR α -SREBP-1c pathway.

In conclusion, we demonstrate that nectandrin B shows a potent stimulating effect on the AMPK activity in hepatocytes and has the ability to prevent hepatic lipogenesis and fat accumulation promoted by LXR α -SREBP-1c-mediated pathway. These

results may suggest the potential of nectandrin B as a promising therapeutic candidate for fatty liver disease.

4. Experimental

4.1. Materials

Antibodies against AMPK, phospho-AMPK, ACC, and phospho-ACC were obtained from Cell Signaling Technology (Beverly, MA). Anti-SREBP-1c antibody was purchased from Santa Cruz Biotechnology (Santa Cruz, CA). T0901317 was supplied by Calbiochem (San Diego, CA). Anti- β -actin antibody, GW3965, insulin, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) and other reagents were provided by Sigma-Aldrich Co. (St. Louis, MO). Nectandrin B was kindly provided by Korea Bioactive Natural Material Bank (KBNMB, Republic of Korea) at Seoul National University. It was isolated from nutmeg as described in the previous study (Nguyen et al. 2010) and its purity was confirmed to be >95% by HPLC.

4.2. Cell culture

HepG2 cells were purchased from ATCC (Manassas, VA) and cultured 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 in a humidified atmosphere of 5% CO₂ at 37 °C.

4.3. Isolation of mouse hepatocytes

Mouse hepatocytes were isolated from C57BL/6 mice as previously described non-recirculating two-step perfusion method with a minor modification (Klaunig et al. 1981). Briefly, liver was perfused with Hank's balanced salt solution (HBSS) without calcium and magnesium at 37 °C for 5 min and then with DMEM containing 0.1% collagenase at a perfusion flow rate of 8 mL/min for 5 min. Next, the liver was excised and minced gently in DMEM containing 0.1% collagenase. The cell suspension was filtered through membrane filter and centrifuged at 50 g for 2 min at 4 °C. The pellet was resuspended with DMEM containing 10% FBS, 100 U/mL penicillin, 100 μ g/mL streptomycin, and 10 nM dexamethasone and centrifuged at 50 g for 2 min at 4 °C. The viability of isolated hepatocytes was estimated to be above 90%. The isolated mouse hepatocytes were seeded at a density of 4×10^5 cells per well in 6-well plates. The culture medium was changed to serum free DMEM 4 h after plating. The next day, cells were used for further experiment.

4.4. Cell viability assay

HepG2 cells were seeded at a density of 5×10^4 cells per well in 48-well plates. The cells were incubated with 1–30 μ g/mL nectandrin B for 24 h after overnight serum starvation and subsequently exposed to 0.5 mg/mL MTT for 1 h at 37 °C. Formazan crystals produced in each well were dissolved by adding 300 μ l of dimethylsulfoxide after removal of the media. Absorbance was determined at 570 nm with an Infinite M200 PRO microplate reader (Tecan, Salzburg, Austria).

4.5. Immunoblot analysis

HepG2 cells were plated in 6-well plates and grown to reach 70%–80% confluency before serum starvation. Immunoblot analyses were conducted according to the procedures previously described (Kim et al. 2011).

4.6. Luciferase assay

Luciferase assay was performed according to the method previously described (Kim et al. 2011). Briefly, HepG2 cells were seeded in 6-well plates at a density of 4×10^5 cells/well. The next day, cells were transiently transfected with a pTK-CYP7A1-LXRE($\times 3$)-LUC (1 μ g) reporter gene construct containing three copies of an LXR response element (LXRE) sequence in rat *Cyp7a1* gene (Hwahng et al. 2009) and pCMV-LacZ using FuGENE (Roche, Nutley, NJ) in Opti-MEM (GIBCO BRL) for 3 h. The transfected cells were then incubated in Eagle's minimal essential medium (MEM) containing 1% FBS for 15 h and exposed to T0901317 in the presence or absence of nectandrin B for an additional 12 h.

4.7. Real-time PCR assay

Total RNA was extracted from HepG2 cells using Trizol (Invitrogen, Carlsbad, CA) in accordance with the manufacturer's instructions. The cDNA was obtained by reverse-transcription of the isolated RNA (2 μ g) with oligo-d(T)₁₆ primers in SureCycler 8800 (Agilent Technologies, Santa Clara, CA). Real-time PCR was conducted with LightCycler®

480 II instrument using LightCycler® 480 SYBR Green I Master solution according to the manufacturer's instructions (Roche Diagnostics, Indianapolis, IN). In addition, a melting curve analysis was performed to verify the accuracy of each amplicon. The mRNA level of interest was normalized to that of glyceraldehyde-3-phosphate dehydrogenase (GAPDH). The following primers were used; human SREBP-1 (sense: 5'-CGACATCGAAGACATGCTTCAG-3', antisense: 5'-GGAAGGCTT-CAAGAGAGGAGC-3'); human LXR α (sense: 5'-GATCGAGGTGATGCTTCTGGAG-3', antisense: 5'-CCCTGCTTTGGCAAAGTCTTC-3'); human FAS (sense: 5'-GACATCGTCCATTTCGTTTGTG-3', antisense: 5'-CGGATCACCTTCTTGAGCTCC-3'); human ACC1 (sense: 5'-GCTGCTCGGATCACTAGTGAA-3', antisense: 5'-TTCTGCTATCAGTCTGTCCAG-3') and human GAPDH (sense: 5'-GAAGGTGAAGGTCGGAGTC-3', antisense: 5'-GAAGATGGTGA-TGGGATTC-3').

4.8. Oil Red O staining

After serum starvation, isolated mouse hepatocytes were exposed to 10 μ M T0901317 for 24 h with 1 h pretreatment of vehicle or 10 μ g/mL nectandrin B. The cells were washed with PBS and then fixed with 4% paraformaldehyde for 1 h. The cells were washed again with PBS, and stained with Oil Red O solution for 2 h at room temperature. The cells were then washed with DW and examined by microscopy (Nikon Eclipse Ti, Japan).

4.9. Statistical analysis

All data are reported as mean \pm S.E. Statistically significant differences were assessed by the Student's *t*-test or one-way ANOVA with Bonferroni's multiple comparison test. A probability value of less than 0.05 was considered significant.

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