



## Ginsenoside Rb1 Induces Beta 3 Adrenergic Receptor–Dependent Lipolysis and Thermogenesis in 3T3-L1 Adipocytes and *db/db* Mice

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Lim S, Park J and Um J-Y (2019) Ginsenoside Rb1 Induces Beta 3 Adrenergic Receptor–Dependent Lipolysis and Thermogenesis in 3T3-L1 Adipocytes and db/db Mice. Front. Pharmacol. 10:1154. doi: 10.3389/fphar.2019.01154 Obesity is constantly rising into a major health threat worldwide. Activation of brown-like transdifferentiation of white adipocytes (browning) has been proposed as a promising molecular target for obesity treatment. In this study, we investigated the effect of ginsenoside Rb1 (Rb1), a saponin derived from Panax ginseng Meyer, on browning. We used 3T3-L1 murine adipocytes and leptin receptor mutated db/db mice. The lipid accumulation, AMP-activated protein kinase alpha (AMPKa)-related pathways, lipolytic and thermogenic factors were measured after Rb treatment in 3T3-L1 adipocytes. Body weight change and lipolysis-thermogenesis factors were investigated in Rb1-treated db/db mice. Beta 3 adrenergic receptor activation (B3AR) changes were measured in Rb1-treated 3T3-L1 cells with or without β3AR inhibitor L748337 co-treatment. As a result, Rb1 treatment decreased lipid droplet size in 3T3-L1 adipocytes. Rb1 also induced phosphorylations of AMPKa pathway and sirtuins. Moreover, lipases and thermogenic factors such as uncoupling protein 1 were increased by Rb1 treatment. Through these results, we could expect that the non-shivering thermogenesis program can be induced by Rb1. In db/db mice, 6-week injection of Rb1 resulted in decreased inguinal white adipose tissue (iWAT) weight associated with shrunken lipid droplets and increased lipolysis and thermogenesis. The thermogenic effect of Rb1 was possibly due to β3AR, as L748337 pre-treatment abolished the effect of Rb1. In conclusion, we suggest Rb1 as a potential lipolytic and thermogenic therapeutic agent which can be used for obesity treatment.

Keywords: ginsenoside Rb1, obesity, non-shivering thermogenesis, lipolysis, beta 3 adrenergic receptor

## INTRODUCTION

Obesity is rising as a major health issue worldwide, especially in developed countries. As it is a risk factor for type 2 diabetes mellitus (T2DM) and other chronic metabolic disorders, the interest in managing obesity is constantly growing (Hossain et al., 2007). There are currently five different drugs approved by the United States Food and Drug Administration: orlistat, lorcaserin, phentermine/topiramate, naltrexone/bupropion, and liraglutide (Daneschvar et al., 2016). These

drugs indeed help people lose weight, but serious side effects, such as steatorrhea, headache, nausea, vomiting, etc., encourage the search for alternative treatments (Azhar et al., 2016).

Obesity is a condition defined where excess/abnormal fat accumulation impacts health. As they are the key players in energy homeostasis, the importance of adipose tissues cannot be neglected in the strategy for obesity treatment. Adipose tissues can be categorized into two subsets based on function and morphology: white adipose tissue (WAT) and brown adipose tissue (BAT). The function of WAT is to store the energy in the form of lipids (Luo et al., 2016), while BAT dissipates energy as heat by a process called non-shivering thermogenesis (Yao et al., 2011). Non-shivering thermogenesis is a defense mechanism to fight against cold through enhanced mitochondrial content and uncoupling protein 1 (UCP1). Since its identification in the 16th century, intense research on BAT revealed its role in thermogenesis during hibernation (Smith, 1961), and it is now appreciated as the main organ for non-shivering thermogenesis (Virtanen, 2016).

Recently, apart from classical BAT, recruitable brown fat has risen into another potential strategy for obesity management (Harms and Seale, 2013). These brown-like white adipocytes, also called "beige" or "brite" adipocytes, mimic the role of brown adipocytes by expressing abundant mitochondria associated with high expression levels of UCP1 (Carobbio et al., 2019). Beige and brown adipocytes share numerous characteristics, from the UCP1-mediated thermogenesis to several differentiation factors such as peroxisome proliferatoractivated receptor gamma (PPARy) coactivator 1 alpha (PGC1a) and RD1-BF1-RIZ1 homologous domain containing 16 (PRDM16) (Castro et al., 2017). As various stimuli such as cold exposure, endogenous signals, as well as dietary factors and pharmacological agents can induce the transdifferentiation of white adipocytes into beige adipocytes (Azhar et al., 2016), the search for relatively safe natural products which can induce "browning" of white adipocytes is a promising, attractive target for obesity treatment.

Ginsenoside Rb1 (C54H92O23, Rb1) (chemical structure shown in Figure 1A) is a triterpenoid saponin derived from the highly valued herb, Panax ginseng Meyer (P. ginseng). This herb has been used as a powerful tonic for qi and blood for over 50 centuries in traditional Chinese and Korean medicine and is also appreciated by the western countries as well (Zhou et al., 2019). Among the known saponins of P. ginseng, Rb1 is considered as one of the most abundant ingredients which may be responsible for the various biological functions of P. ginseng (Zhu et al., 2019). Several studies report the anti-adipogenic effect of Rb1 (Park et al., 2008; Xiong et al., 2010; Shen et al., 2013; Lin et al., 2014; Yu et al., 2015). In contrast, an early work by Shang et al. reported that Rb1 promotes adipogenesis by enhancing two major adipogenic factors, CCAAT/enhancer binding protein alpha (C/EBPa) and PPARy (Shang et al., 2007). Furthermore, this was later supported by Mu's study, as it showed that Rb1-induced increase of PPARy and C/EBPa could have been a result of its browning effect in adipocytes. Mu's team reported that Rb1 significantly increased the levels of UCP1, PGC1a, and PRDM16, thus leading to increased thermogenic capacity of 3T3-L1 adipocytes (Mu et al., 2015). However, although the browning effect of Rb1 has already been reported, its detailed mechanism still remains unknown to date. We hereby show that Rb1 treatment indeed resulted in browning of 3T3-L1 adipocytes, and this effect was due to regulation of beta 3 adrenergic receptor (B3AR)-mediated lipolysis induced by Rb1.

### MATERIALS AND METHODS

#### **Chemical Reagents and Antibodies**

Rb1 (>98%, ab142646) was purchased from Abcam (Cambridge, UK). 3-Isobutyl-1-methylxanthine (IBMX), dexamethasone (Dex), insulin, and Oil Red O powder were purchased from Sigma (St. Louis, MO, United States). L748337 was from Tocris Bioscience (Bristol, UK). Dulbecco's Modified Eagle's Medium (DMEM) and fetal bovine serum (FBS) were



**FIGURE 1** Chemical structure of Rb1 and cytotoxicity of Rb1 in 3T3-L1 adipocytes. (A) Chemical structure of Rb1 is shown. (B) An MTS assay was performed in order to evaluate the cytotoxicity of Rb1 on 3T3-L1 adipocytes. Data are expressed as mean  $\pm$  standard error of the mean (S.E.M.) of three or more experiments. \*p < 0.05 vs. untreated cells. Rb1, ginsenoside Rb1.

purchased from Corning (NY, United States). Antibodies for liver kinase B1 (LKB1) (3047S), pLKB1 (Ser428) (3482S), AMP-activated protein kinase alpha (AMPKa) (2532S), pAMPKa (Thr172) (2535S), acetyl-CoA carboxylase (ACC) (3676S), pACC (Ser79) (3661S), silent information regulator T1 (SIRT1) (8469S), SIRT3 (5490S), phospho-hormone sensitive lipase (pHSL) (Ser563) (4139S), phospho-PKA substrate (9624S), UCP1 (14670S), and β-actin (3700S) were purchased from Cell Signaling Technology (Beverly, MA, United States); the antibody for glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (sc-32233) antibody was purchased from Santa Cruz Biotechnology (Santa Cruz, CA, United States); antibodies for PGC1a (ab54481), comparative gene identification 58 (CGI58) (ab59488), adipose triglyceride lipase (ATGL) (ab207799), HSL (ab45422), and β3AR (ab94506) were purchased from Abcam (Cambridge, UK); the antibody for PPARa (PA1-822A) was purchased from Thermo Fisher Scientific (Waltham, MA, USA).

## **Cell Culture and Differentiation**

3T3-L1 adipocytes from mouse embryo fibroblasts cell lines were obtained from the American Type Culture Collection (Rockville, MD, USA), cultured, and differentiated into mature white adipocytes as previously described (Jung et al., 2018). Briefly, cells were grown in DMEM containing 10% FBS and 100 units/ml of penicillin streptomycin solution at 37°C, in 5% CO<sub>2</sub>, at 95% humidity until confluence. After 2 days from full confluence (day 0), the cells were differentiated by a 48 h incubation in differentiation medium (DMEM plus 10% FBS containing 0.5 mM IBMX, 1  $\mu$ M Dex, and 1  $\mu$ g/ml insulin). At day 2, the cells were cultured in the maintenance medium (DMEM plus 10% FBS supplemented with 1  $\mu$ g/ml insulin) and various concentrations of Rb1 (1, 5, 10 and 20  $\mu$ M) for another 48 h followed by fresh maintenance medium.

Stromal vascular fraction (SVF) cells were isolated, cultured, and differentiated into mature beige adipocytes as previously described (Aune et al., 2013). Briefly, inguinal white adipose tissue (iWAT) was dissected from male 6-week-old C57BL/6J mice and digested in collagenase D and dispase II in 37°C with constant agitation at 150 rpm for 40-50 min. The cells were filtered using a cell strainer (50-70 µm diameter) and plated in 10 cm plates, and the red blood cells, immune cells, and other contaminants were removed by washing with PBS. Cells were grown in complete medium (DMEM containing 10% FBS and 100 units/ml of penicillin streptomycin solution) at 37°C, in 5% CO<sub>2</sub>, at 95% humidity until 95–97% confluence (day 0). At day 0, the cells were differentiated by a 48 h incubation in differentiation medium consisting of maintenance medium (complete medium containing 5 µg/ml insulin and 1 nM T3) including 0.5 mM IBMX, 2 µg/ml Dex, and 125 µM indomethacin. After 48 h (day 2), the cells were cultured in maintenance medium with 0.5 µM rosiglitazone and 20 µM Rb1. After an additional 48 h (day 4), the medium was changed to fresh maintenance medium with 1 µM rosiglitazone for additional 2 days. At 6 days, cells are fully differentiated to mature beige adipocytes.

Human adipose tissue–derived mesenchymal stem cells (hAMSCs) (Cell Engineering for Origin, Seoul, Korea) were grown in DMEM containing 10% FBS and 100 units/ml of penicillin streptomycin solution at 37°C, in 5% CO<sub>2</sub>, at 95% humidity until confluence. After 2 days from full confluence (day 0), the cells were differentiated by a 72 h incubation in differentiation medium (DMEM plus 10% FBS containing 0.5 mM IBMX, 1  $\mu$ M Dex 1  $\mu$ M insulin, and 100  $\mu$ M indomethacin). After 72 h (day 3), the cells were placed in fresh differentiation medium for an additional 3 days. At 6 days, the cells were cultured in the maintenance medium (consisting of DMEM with 10% FBS supplemented with 1  $\mu$ M insulin) and 20  $\mu$ M Rb1 for 72 h. From days 9 to 15, the maintenance medium was changed every 3 days.

Brown preadipocytes were obtained as described previously (Klein et al., 1999). Briefly, interscapular BAT was dissected from newborn FVB mice (age, post-natal day 1) and subjected to collagenase digestion for 30 min. The cells were filtered through a 100 µm strainer. The collected cells were centrifuged at 200 g for 5 min. Cells were seeded on 10 cm plates and grown in DMEM containing 10% FBS and 100 units/ml of penicillin streptomycin solution at 37°C, in 5% CO<sub>2</sub>, at 95% humidity. On day 2 after confluence (day 0), the cells were differentiated by a 48 h incubation in differentiation medium (DMEM plus 10% FBS containing 0.5 mM IBMX, 0.5 µM Dex, 20 nM insulin, 125 mM indomethacin, and 1 nM T3). At day 2, the cells were cultured in the maintenance medium (DMEM plus 10% FBS supplemented with 20 nM insulin and 1 nM T3) and 20 µM Rb1 for another 48 h followed by fresh maintenance medium.

## **Cell Cytotoxicity Assay**

Cell viability was measured with a Cell Proliferation MTS kit (Promega Co., Madison, WI, USA) as previously described (Jung et al., 2018). Briefly, cells were seeded ( $2 \times 10^4$  cells per well) on 96-well plates and incubated for 24 h, followed by incubation with various concentrations (1, 5, 10, 20, 40, and 100  $\mu$ M in 3T3-L1 cells) of Rb1 in culture medium for an additional 48 h. The absorbance was measured at 490 nm in a VERSAmax microplate reader (Molecular Devices, Sunnyvale, CA, USA).

## **Oil Red O Staining**

Intracellular triglyceride (TG) accumulation was measured using the Oil Red O staining method as previously described (Jung et al., 2018). Photomicrograph images were obtained from a regular light microscope (Olympus, Tokyo, Japan), and absorbance was measured at 500 nm using a VERSAmax microplate reader (Molecular Devices, Sunnyvale, CA, USA).

## Protein Extraction and Western Blot Analysis

Western blot analyses were performed as previously described (Jung et al., 2018). In brief, homogenized tissues and harvested cells were lysed in lysis buffer (Cell Signaling Technology, Beverly, MA, United States), and protein concentration was determined using a protein assay reagent (Bio-Rad Laboratories, Hercules, CA, USA). Equal amounts of total protein were resolved by 8–12% sodium dodecyl sulfate–polyacrylamide gel electrophoresis and transferred to a polyvinylidene difluoride membrane. The membranes were incubated with the primary antibody at 4°C overnight and then incubated with a 1:10,000 dilution of the proper horseradish peroxidase (HRP)– conjugated secondary antibody (Jackson Immuno Research, West Grove, PA, USA) for 1 h at RT.

#### Immunofluorescence Assay

Immunofluorescence (IF) analyses were performed as previously described (Jung et al., 2018). BODIPY 558/568 C12 (D3835) for staining lipid droplets was purchased from Thermo Fisher Scientific (Waltham, MA, USA). Images were acquired using a fluorescence microscope (Logos Biosystems, Anyang, Korea).

#### Oxygen Consumption Rate Assay

Oxygen consumption rate (OCR) was measured by a Mito-ID Extracellular  $O_2$  sensor kit (OCR; ENZ-51045, ENZO Life Sciences, Farmingdale, NY, USA) according to the manufacturer's instructions. Briefly, cells were prepared in 96-well black plates, and an  $O_2$  sensor probe (10 µl) was added into each well. After sealing the wells by adding two drops of provided oil, the absorbance was analyzed with a fluorescence plate reader.

#### Free Fatty Acid Assay

Cellular free fatty acid (FFA) was measured by an EZ-Free Fatty Acid Assay Kit (DG-FFA100, Dogenbio, Seoul, Korea) according to the manufacturer's instruction. Briefly, the FFA buffer was added to the prepared cells and reacted with the acyl-CoA synthesis for 30 min. After reaction, the reaction buffer was added and reacted at 37°C without light for 30 min. Then, the absorbance was read with a microplate reader.

#### Cyclic Amp Assay

Cyclic AMP (cAMP) was measured using a cAMP ELISA Kit (STA-500, Cell Biolabs, Inc., San Diego, CA, USA) according to the manufacturer's instruction.

## **RNA Isolation and Real-Time Reverse Transcription Polymerase Chain Reaction**

RNA extraction was using a GeneAll RiboEx Total RNA extraction kit (GeneAll Biotechnology, Seoul, Korea) according to the manufacturer's instruction. cDNA synthesis was using a Maxime RT PreMix Kit (iNtRON Biotechnology, Seoul, Korea). Realtime reverse transcription polymerase chain reaction (RT-PCR) was performed with SYBR Green Power Master Mix (Applied Biosystems, Foster City, CA, USA) and the Step One Real-Time PCR System (Applied Biosystems, Foster City, CA, USA) according to the manufacturer's instruction provided. The primers used in this study are provided in **Table 1**.

#### **Animal Experiment**

Male 6-week-old db/db mice and age-matched wild-type (WT) C57BL6/J mice were purchased from Daehan Biolink Co. (Eumsung, Korea) and maintained for 1 week prior to the experiments, provided with a laboratory diet and water ad libitum. Healthy mice were randomly divided into three groups as follows (n = 5 per group): a WT group, a db/db group, and a db/db group administered with Rb1. The control groups (WT group, db/db group) were administered distilled water (i.p.), while the experiment group was administered Rb1 prepared in distilled water (10 mg/kg of body weight, i.p.) five times per week for 6 weeks. Body weight and food intake were measured three times per week. At the end of the experiment, the animals were anesthetized; tissues were collected and placed in a tube and then stored at  $-80^{\circ}$ C.

#### Hematoxylin and Eosin Staining

Hematoxylin and eosin (H&E) staining was done as previously described (Jung et al., 2018). Photographs were taken under a microscope, EVOS M7000 (Thermo Fisher Scientific, Waltham, MA, USA).

#### Measurement of Body Fat by Dual-Energy X-Ray Absorptiometry Scan

A body fat scan was performed using dual-energy X-ray absorptiometry (DXA) with an InAlyzer instrument (Medikors, Seongnam, Korea) based on the manufacturer's instructions as in a previous report (Lee et al., 2019).

TABLE 1 | Primer sequences used for real-time reverse transcription polymerase chain reaction (RT-PCR).

Primer, 5'–3'		
Target gene	Sense	Antisense
Atgl	ATATCCCACTTTAGCTCCAAGG	CAAGTTGTCTGAAATGCCGC
Cd137	TCTGTTGCTGGTCCTCAACTT	CTCCTACCTGGTCCTGAAAAC
Cpt1a	GACTCCGCTCGCTCATTCC	GACTGTGAACTGGAAGGCCA
Cpt1b	GCACACCAGGCAGTAGCTTT	CAGGAGTTGATTCCAGACAGGTA
Gapdh	AACTTTGGCATTGTGGAAGG	GGATGCAGGGATGATGTTCT
Prdm16	CCAAGGCAAGGGCGAAGAA	AGTCTGGTGGGATTGGAATGT
Tbx1	ATGATCTCCGCCGTGTCTAG	CGTGGGGAACATTCGTCTGCCTG
Tmem26	GAAATGCACCATGGAACCC	CGGTTCACATACCATGGATAA
Ucp1	AACTGTACAGCGGTCTGCCT	TAAGCCGGCTGAGATCTTGT
Adrb3	GGCCCTCTCTAGTTCCCAG	TAGCCATCAAACCTGTTGAGC

#### **Statistical Analysis**

Data were expressed as the means  $\pm$  standard error of the mean (S.E.M.). Significant differences (p < 0.05) between groups were determined with the ANOVA test. All statistical analyzes were performed using SPSS statistical analysis software version 11.5 (SPSS Inc., Chicago, IL, USA).

### RESULTS

## Rb1 Decreases the Size of Lipid Droplets in 3T3-L1 Adipocytes

To evaluate the cytotoxicity of Rb1, an MTS assay was performed. As in **Figure 1B**, concentrations up to 20  $\mu$ M did not affect cell viability of 3T3-L1 adipocytes. Then, an Oil Red O staining assay was conducted to assess the effect of Rb1 on lipid accumulation (**Figure 2A**). Rb1 treatment did not suppress nor increase TG accumulation in 3T3-L1 adipocytes (**Figure 2B**); however, the size of lipid droplets was significantly reduced in Rb1-treated cells when compared to differentiated adipocytes (**Figures 2C, D**). The portion of large lipid droplets (>160  $\mu$ m<sup>2</sup>) was reduced from 8.4% to 0.4%, while the portion of small droplets (<20  $\mu$ m<sup>2</sup>) was increased from 31.6% to 50.3%. The change in lipid droplet

morphology suggests that Rb1 may have induced browning in 3T3-L1 adipocytes.

### Rb1 Increases Phosphorylation of LKB1– Ampkα–ACC Pathway and Expression of Sirtuin Proteins in 3T3-L1 Adipocytes

AMPKα is a well-known sensor and regulator of energy metabolism (Steinberg and Carling, 2019). As shown in **Figures 3A–D**, Rb1 treatment upregulated the phosphorylation level of AMPKα in a concentration-dependent manner, showing that Rb1 can increase energy metabolism in 3T3-L1 adipocytes. Similar results were shown in the upstream and downstream factors of AMPKα, LKB1, and ACC, respectively.

After observing the effect of Rb1 on the AMPK $\alpha$  pathway, we then performed further experiments to verify whether Rb1 affects the levels of sirtuins. The sirtuins, consisting of seven homologs in mammals, are a family of NAD+-dependent histone/protein deacetylases (Michan and Sinclair, 2007). Among the seven sirtuins, SIRT1 and SIRT3 are closely related to the AMPK $\alpha$  function in lipid homeostasis (Ruderman et al., 2010; Shi et al., 2010). Our results showed that Rb1 also increased the levels of both sirtuins, but without statistical significance in SIRT3 (**Figures 3E–G**).







## Rb1 Induces Lipolysis in 3T3-L1 Adipocytes

AMPKα, the energy metabolism sensor, also acts as a regulator of lipolysis by phosphorylating ATGL and HSL, the two major lipases working in the lipolytic process (Kim et al., 2016). Since we verified the effect of Rb1 on the AMPKα pathway and sirtuins, our next objective was to investigate its effect on lipolysis. CGI58, a critical regulator of ATGL, showed an increasing tendency in Rb1-treated adipocytes (**Figures 4A, B**). In addition, protein levels of ATGL was significantly increased by Rb1 treatment, while pHSL was not (**Figures 4A, C, D**). The lipolytic activation induced by Rb1 was re-confirmed when we performed an IF staining assay of ATGL and pHSL (**Figures 4E, F**).

Furthermore, intracellular FFA was increased by Rb1 treatment (p < 0.05), confirming its lipolytic effect (**Figure 4G**). As cAMP and its downstream target PKA are regulators of lipolysis, we then measured the change in these factors. Indeed, cAMP and PKA both were significantly induced in Rb1-treated 3T3-L1 cells (**Figures 4H, I, J**).

To confirm, we obtained SVF cells from iWAT of C57BL/6J mice and induced beige differentiation based on a previous report (Aune et al., 2013). Rb1 treatment in SVF cells also increased the mRNA expression of *Atgl*, the gene which transcripts ATGL (**Supplementary Figure S1**).

# **Rb1 Increases Thermogenic Capacity in 3T3-L1 Adipocytes and Induces Browning**

Lipolysis is known as the pre-step of brown and beige adipocyte thermogenesis. Importantly, lipolysis plays a central role in the catabolic activity of BAT and WAT. FAs induce oxidative phosphorylation and provide fuel that supports UCP1-mediated respiration (Granneman et al., 2003). Lipolysis also provides ligands for PPARa, which acts crucially in the browning of WAT (Li et al., 2005). As shown in Figures 5A, B, Rb1-treated adipocytes showed increased levels of PPARa, implying the thermogenic triggering by Rb1. PGC1a, the major transcription factor of both brown and beige adipocytes (Inagaki et al., 2016), was also increased by Rb1 treatment (Figures 5A, C). The thermogenic capacity was determined by assessing the changes in UCP1 (Figures 5A, D). Rb1 treatment induced a 2.06-fold change in the protein expression of UCP1 when compared to vehicle-treated adipocytes. Similar results were retrieved from Rb1-treated SVF cells (Supplementary Figure S1) and hAMSCs (Supplementary Figure S2) as well. Further real-time RT-PCR assays showed increased beige-specific gene levels of Prdm16, *Tmem26*, *Tbx1*, and *CD137* in Rb1-treated cells (Figure 5E).

Upregulated factors of mitochondrial beta oxidation such as mRNA of *Cpt1a* and *Cpt1b* also indicated the Rb1-induced thermogenic capacity (**Figure 5F**). Since the thermogenic action of UCP1 requires alteration of oxidative levels, we evaluated the change in  $O_2$  consumption, which in turn was found to be increased by Rb1 treatment (**Figure 5G**).

# Rb1 Induces Browning in iWAT of *db/db* Mice

We then conducted an animal study to evaluate the effect of Rb1 *in vivo*. After feeding obese *db/db* mice with Rb1 (10 mg/kg/ day) for 6 weeks, we observed unchanged food intake between vehicle-fed and Rb-fed *db/db* mice (**Figure 6A**). Surprisingly, not in accordance with the *in vitro* results, Rb1 did not affect



body weight change in *db/db* mice (data not shown). However, a significant reduction in iWAT was shown in Rb1-treated mice, while epididymal white adipose tissue (eWAT), BAT, and liver tissue weight were not affected (**Figure 6B**). A DXA scan analysis showed decreased fat body mass in the Rb1-fed group compared to the vehicle-treated *db/db* group (**Figures 6C, D**). Next, we evaluated the Rb1-induced histological changes in iWAT. As a result, the average size of lipid droplets was increased by 4.68-fold in *db/db* mice compared to WT mice, and this was decreased down to 61% by Rb1 administration (**Figures 6E, F**).

Further western blot assays were performed to evaluate the effect of Rb1 on thermogenesis-related factors. As shown in **Figures 6G**, **H**, Rb1 treatment increased the protein expressions of pAMPK $\alpha$ , ATGL, and UCP1 in iWAT of mice, suggesting that a browning effect was induced by Rb1.

# Thermogenic Effect of Rb1 Is Dependent on $\beta$ 3AR Pathway

As  $\beta$ 3AR is considered as one of the highest upstream signals of non-shivering thermogenesis (Arch, 2002; Dulloo, 2011),



we attempted to investigate whether Rb1 could regulate the expression of this receptor. As expected, Rb1 treatment in 3T3-L1 adipocytes resulted in a dose-dependent increase of  $\beta$ 3AR expression (**Figure 7A**). Similar results were observed in Rb1-treated SVF cells and hAMSCs (**Supplementary Figures S1** and **S2**). Furthermore, when L748337, a selective  $\beta$ 3AR antagonist, was pre-treated in adipocytes, the effect of Rb1 on UCP1 induction was abolished, down to nearly 67% (**Figure 7B**), suggesting that the thermogenic effect of Rb1 is dependent on the  $\beta$ 3AR signaling pathway, at least partially.

Since  $\beta$ 3AR is closely related to the activation of classical brown fat as well, we attempted to figure out whether Rb1 treatment regulates thermogenesis in BAT. As in **Supplementary Figures S3A, B,** Rb1-treated *db/db* mice displayed a significantly higher level of  $\beta$ 3AR in BAT. Consequently, lipolysis and non-shivering thermogenesis seemed to be induced, when confirmed by the protein levels of ATGL and UCP1. An *in vitro* study with primary cultured brown adipocytes showed similar results (**Supplementary Figure S3C**).

## DISCUSSION

The negative impact of obesity on mankind health forces clinicians and researchers to seek a promising anti-obese strategy. In 2016, the World Health Organization (WHO) reported that around 2 billion adults were overweight, and 650 million obese (WHO, 2018). However, currently available strategies, mostly medications, display unwelcome side effects (Daneschvar et al.,

2016). Thus, the task of safe and effective anti-obese agents is still an ongoing challenge. In this context, the potential of natural products, also proved by the steady growth of their market size (Brown, 2017), may give an advantage for the next promising strategy for obesity care.

There are some obvious clues to the anti-obese action of Rb1 from previously published literature. Park et al. reported that Rb1 and another saponin of P. ginseng, ginsenoside Rg1, suppressed TG accumulation in vitro (Park et al., 2008), while Xiong et al. showed that the anti-obese effect of Rb1 was effective in vivo as well (Xiong et al., 2010). Shen et al. suggested that AMPK was responsible for this effect (Shen et al., 2013), Lin and colleagues explained that it resulted from decreased appetite by Rb1-regulated neuropeptide Y (NPY) and peptide YY (PYY) (Lin et al., 2014), and Yu et al. reported that perilipin expression was the clue (Yu et al., 2015). Further evidence also supports the potential thermogenic effect of Rb1. Several studies have recently reported the browning or lipolytic effect of a wide variety of ginsenoside isoforms besides Rb1. Regarding ginsenoside Rg1, a quantitively important isoform (Lü et al., 2009), it is known to reduce TG accumulation in adipocytes (Park et al., 2008), suppress hepatic glucose production in HepG2 cells (Kim et al., 2010), and increase glucose uptake in muscle cells (Lee HM et al., 2012). Furthermore, Li et al. reported its beneficial effect in glucose metabolic disorder (Li et al., 2018), and Liu et al. showed the AMPK-mediated anti-obese effect of ginsenoside Rg1 (Liu et al., 2018). A recent study also reported the browning potential of ginsenoside Rg1 (Lee et al., 2018). Ginsenoside Rg3 also shows improvement in metabolic diseases in vivo and in vitro (Hwang et al., 2009; Kim SN et al., 2009; Lee et al., 2011; Lee et al., 2017; Zhang et al., 2017). Ginsenoside Rb2,



another abundant isoform of ginsenosides, is reported to possess beneficial effects on lipid accumulation. Cholesterol and TG levels were attenuated by ginsenoside Rb2 treatment in 3T3-L1 adipocytes (Kim EJ et al., 2009), and similar results were shown in high-fat diet–fed obese mice (Dai et al., 2018). Moreover, significant activation of thermogenic factors in BAT and WAT were observed in ginsenoside Rb2–fed obese mice (Hong et al.,

2019). Through these obvious hints from previous literature, we could expect the potential benefits from Rb1; however, to date, the exact cascade mechanism of the thermogenic effect of Rb1 still remains to be elucidated. Therefore, in this study, we aimed to investigate the detailed action mechanism of Rb1 on beige adipocyte recruitment and thermogenesis induction using db/db mice and 3T3-L1 adipocytes.



AMPK is a metabolism regulator protein consisting of three subunits: catalytic  $\alpha$ , regulatory  $\beta$ , and  $\gamma$  (Hardie, 2018). As its name suggests, the elevation of AMP/ADP associated with reduced ATP triggers the activating phosphorylation of AMPK specifically at Thr172 of the  $\alpha$  subunit (Oakhill et al., 2012). Once phosphorylated, AMPK regulates activation of metabolic proteins and transcription factors, leading to an energy production. Numerous nature-derived materials such as berberine (Kim WS et al., 2009), quercetin (Ahn et al., 2008), and resveratrol (Wang et al., 2015) are shown to induce AMPK activation and thus ameliorate metabolic diseases. Our results also suggest the possible use of Rb1 as an AMPK $\alpha$ -activating anti-obesity agent. Rb1 increased phosphorylation of not only AMPK $\alpha$  but also its upstream kinase LKB1 and downstream factor ACC. In addition, Rb1 increased the levels of SIRT1, which is known to facilitate metabolic AMPK action (Ruderman et al., 2010).

Non-shivering thermogenesis in BAT and WAT is a promising molecular target for obesity management. The recently discovered beige adipocytes within WAT differ from classic white adipocytes both by morphology and ability. By certain stimulation, such as cold exposure (Vitali et al., 2012; Wu et al., 2012) or pharmacological activation (Lee JY et al., 2012; Rachid et al., 2015), these beige adipocytes, which were normally acting as a storage unit for lipid, become capable of producing heat through mitochondrial UCP1 activation. Once activated in either BAT or WAT, the thermogenic action of mitochondria requires fuel: FAs. In order to supply FAs, the lipolysis signaling is activated within the adipocyte, and the lipases ATGL, HSL, and monoacylglycerol lipase (MG) subsequently process TG into FAs for  $\beta$ -oxidation (Steensels and Ersoy, 2019). In this study, we observed that Rb1 can increase ATGL, suggesting a possible role of Rb1 in induction of the lipolysis signaling for thermogenic actions.

β3ARs have a critical role in activation of non-shivering thermogenesis via PKA signaling, in both BAT and "beiged" WAT (Lowell and Spiegelman, 2000; Cao et al., 2001). Jimenez et al. showed that \$3AR knocked-out mice displayed a low mRNA level of Ucp1 and failed to induce thermogenesis by cold stimuli (Jimenez et al., 2003). In accordance,  $\beta$ 3AR activation by pharmacological agonists CL316,243 and isoproterenol results in enhanced UCP1 expression in BAT (Bachman et al., 2002) and higher thermogenic capacity in WAT (Wang et al., 2013). Naturederived nutritional agents are also candidates for β3AR-activated thermogenesis as well. Ephedrine, an active compound of genus *Ephedra*, is shown to induce  $\beta$ 3AR expression and glycerol release, which was potentiated with  $\beta$ 3AR agonist BRL37344 and inhibited by β3AR antagonist SR59230A co-treatment (Cheng et al., 2001). Another team reported that cinnamon extract induces browning in 3T3-L1 adipocytes and WAT of high-fat diet-induced obese mice by activating β3AR (Kwan et al., 2017). Our results suggest another natural product which can induce browning by  $\beta$ 3AR activation. Rb1 administration increased expressions of lipases and thermogenic factors including UCP1, and these effects were suppressed when a  $\beta$ 3AR antagonist, L748337, was pre-treated. Although activation of  $\beta$ 3AR may benefit metabolic diseases such as obesity, the risk of  $\beta$ 3AR in cardiovascular diseases cannot be neglected (Cossu et al., 1997; Sears, 2002). However, various studies report that the effect of Rb1 does not harm but even possibly improves the cardiovascular system. Rb1 can benefit myocardial ischemia (Guan et al., 2002; Wang et al., 2008; Wu et al., 2011; Xia et al., 2011; Kong et al., 2015; Yan et al., 2015; Li et al., 2016; Yan et al., 2016; Cui et al., 2017; Zheng et al., 2017a), atherosclerosis/vascular dysfunctions (Qiao et al., 2017; Zhou et al., 2017; Zhang et al., 2018), or other related diseases

(Jiang et al., 2007; Kong et al., 2010; Li et al., 2011; Li et al., 2012; Zhang et al., 2015; Zheng et al., 2017b), suggesting the potential safety of Rb1 in cardiovascular functions, despite its action on  $\beta$ 3AR.

In our study, we have shown that Rb1 can decrease lipid accumulation *in vivo* and *in vitro* by inducing the lipolysisthermogenesis cascade. This effect was probably due to activation of  $\beta$ 3ARs, as  $\beta$ 3AR inhibitor treatment decreased the thermogenic effect of Rb1. However, further studies are required to understand the whole-body significance of Rb1-activated  $\beta$ 3AR, as our study mainly focused on the beige adipocyte recruitment in WAT. Because the mechanisms of WAT browning and BAT activation are both related in the  $\beta$ 3AR-dependent thermogenesis, thus, to investigate the whole precise mechanism of the anti-obese effect of Rb1, relevant studies dealing with the role of BAT in the effect of Rb1 are necessary. Furthermore, as the non-adrenergic pathway is also capable of progressing lipolysis and thermogenesis (Braun et al., 2018), related investigation on Rb1 action has to be carried out as well.

Overall, our results demonstrate the effect of Rb1 on  $\beta$ 3ARdependent lipolysis and thermogenesis. Regarding the wellknown clinically beneficial features of *P. ginseng* in traditional Korean medicine, we suggest Rb1 as a potentially safe and effective therapeutic agent for treatment of metabolic diseases.

## DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this manuscript will be made available by the authors, without undue reservation, to any qualified researcher.

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### **ETHICS STATEMENT**

All animal experiments were performed according to the Guide for the Care and Use of Laboratory Animals and were approved by the Institutional Review Board of Kyung Hee University (confirmation number: KHUASP (SE)-13-012).

## **AUTHOR CONTRIBUTIONS**

SL, JP and J-YU designed the protocol and prepared the manuscript; SL and JP performed the experiments; J-YU was in charge of the whole experiment conduction and proofreading of the manuscript. All authors approved the final version to be published.

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fphar.2019.01154/ full#supplementary-material

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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