

Article

Anti-Obesity Potential of *Sargassum horneri* and *Ulva australis* Extracts: Study In Vitro and In Vivo

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Abstract: Current research highlights the use of natural products or phytochemicals as drugs and functional additives to treat obesity with few side effects. *Sargassum horneri* (SH) and *Ulva australis* (UA) are marine waste resources on Jeju Island, Republic of Korea. In this study, we analyzed their antioxidant and anti-obesity efficacies to confirm their potential as functional additives. We prepared SH and UA extracts using 80% ethanol and observed that free radical scavenging activity and total phenol content were high in SH extracts, and total flavonoid content was elevated in UA extracts. Additionally, 3T3-L1 cells were treated with SH and UA extracts, and the ability of the extracts to inhibit adipocyte differentiation was examined using Oil Red O staining and analysis of neutral fat content. We confirmed that the mRNA expression of the C/EBP α , PPAR γ , and SREBP1c genes that act on adipocyte differentiation, and of FAS, a fatty acid synthase, was suppressed. Experiments in a mouse model of obesity showed that 12-week administration of a high-fat diet with 1% extract added to drinking water resulted in lower weight gain compared to the high-fat diet alone. These results suggest that SH and UA extracts have antioxidant properties and are effective in obesity prevention. Therefore, the two marine waste resources are potential functional additive candidates for preventing obesity.

Keywords: seaweed; extract; antioxidant; obesity; hyperlipidemia



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1. Introduction

According to the definition by the World Health Organization (WHO), obesity is an unhealthy condition resulting from the abnormal or excessive accumulation of fat in adipose tissue [1,2]. The body mass index (BMI) is a person's weight (kg) divided by the square of their height (m); individuals with BMI ranges between 25 and 30 are considered to be overweight and obese [1–3]. According to the WHO report on obesity, over 1.9 billion adults aged over 18 years old are overweight, and 600 million are reported to be obese, which makes obesity a problem in both developed and developing countries [4,5]. In Korea, according to the National Health and Nutrition Examination Survey, the prevalence of obesity among adults aged 19 years old and older is reported to be ~40% for men and ~26% for women. By 2030, the prevalence rate is expected to reach ~62% for men and 37% for women [6].

Among the factors that cause obesity are lack of exercise, stress, genetic factors, and endocrine disorders. Obesity is mainly caused by excessive body fat accumulation due to

excessive calorie intake beyond consumption [7]. Adipocytes are endocrine organs that control several metabolic processes that regulate the body's physiology. Adipocytes secrete various hormones that regulate metabolism, such as leptin and adiponectin, which are essential for maintaining homeostasis [8]. When excessive fat accumulates in adipocytes, functions such as intracellular endoplasmic reticulum stress and mitochondrial dysfunction are lost, and the risk of complications of obesity, such as hypertension, cardiovascular disease, cancer, and fatty liver, increases through insulin resistance and inflammatory responses [9,10]. Oxidative stress that affects obesity-related diseases increases the probability of various diseases by inducing chemical changes in the human body. Therefore, it is necessary to find antioxidant substances to protect cells in the human body from free radicals and prevent obesity.

Obesity-related studies have used mouse fibroblast 3T3-L1 preadipocytes to analyze the life cycle mechanisms of adipogenesis, lipogenesis, and lipolysis. The production and differentiation of adipocytes are regulated by transcription factors, such as C/EBP α /enhancer-binding protein α (C/EBP α) and peroxisome proliferator-activated receptor γ (PPAR γ) [11]. C/EBP α levels increase in the late stage of differentiation along with those of PPAR γ and induce the expression of genes that induce adipogenesis such as fatty acid synthase (FAS) and adiponectin (ADIPOQ). Their expression levels tend to markedly increase in differentiated adipocytes [12]. The regulation of genes related to adipocyte differentiation is one of the most important strategies for obesity prevention and treatment [13]. Therefore, therapeutic agents effective for adipocyte differentiation and the inhibition of adipogenesis are being released. The available drugs against obesity include orlistat, topiramate, sibutramine, rimonabant, and phenylpropanolamine; however, these drugs have side effects such as insomnia, anorexia, gastrointestinal disorders, increased blood pressure, and heart disease [14]. Overall, the only two drugs currently approved by the Food and Drug Administration (FDA) for long-term use are orlistat and sibutramine, while others are available only for short-term or limited use [15]. To address this problem, research on the development of treatments against obesity using safer natural products or phytochemicals that have fewer side effects is active [16].

Among such natural products, *Sargassum horneri* (SH) and *Ulva australis* (UA), used in the present study, are marine waste resources generated in large quantities on Jeju Island, Republic of Korea. SH, that belongs to the Sargassaceae algae, is a yellow-brown plant with thin, spatula-shaped leaves that grows mainly in winter. UA is a green alga in the Ulvaceae family and inhabits the middle and lower intertidal zones. Due to climate changes and rising water temperatures, the populations of these two plants are increasing and causing environmental and economic issues, including introduction to the coast, odor due to decomposition, the pollution of surrounding ecosystems, and damage to fishing boats [17].

SH, which causes various problems in Jeju, is a food rich in fucoidan and a known delicacy in East Asia, including China and Japan [18]. UA is known to enhance immune activity [19] and promote collagen biosynthesis [20]. Additionally, UA has been used in oriental medicine to treat hyperlipidemia, heat stroke, and urinary disorders [21]. To explore the use of seaweed as a valuable resource, several studies have analyzed the antioxidant [22], immune activity [23,24], whitening and anti-wrinkle [25], and anti-obesity effects [26] of SH. Similarly, some research has evaluated the antioxidant [27] and anti-inflammatory [28] effects of UA and its potential for bioethanol production [29]; yet, studies related to the effects of UA on obesity are scarce.

In this context, the present study compares the antioxidant effects of SH and UA extracts by analyzing their 2,2-diphenyl-1-picrylhydrazyl (DPPH) radical activity and total polyphenol and flavonoid contents. In addition, the inhibitory effect of treatment with SH and UA extracts on 3T3-L1 adipocyte differentiation was analyzed. In an animal model of obesity, induced by a high-fat diet, the effect of extract administration on preventing obesity was investigated.

2. Results

2.1. Total Phenol and Flavonoid Contents

The total phenol content of the SH and UA ethanol extracts was analyzed using gallic acid as a reference material. The total phenol content of the SH extract was 95.5 ± 1.4 mg gallic acid equivalent (GAE)/g, and that of the UA extract was 62.4 ± 1.8 mg GAE/g. Thus, of the two materials, SH showed a higher phenol content than UA (Table 1).

Table 1. Total phenol and flavonoid contents in *S. horneri* and *U. australis* ethanol extracts.

	<i>S. horneri</i>	<i>U. australis</i>
Total phenolic content (mg GAE/g extract)	95.5 ± 1.4	62.4 ± 1.8
Total flavonoid content (mg QE/g extract)	19.4 ± 1.0	20.9 ± 1.2

Data are expressed as the mean \pm standard deviation (n = 3).

The total flavonoid contents of the SH and UA ethanol extracts were analyzed using naringin as a reference material. The results revealed that the total flavonoid contents of the SH and UA extracts amounted to 19.4 ± 1.0 mg quercetin equivalent (QE)/g and 20.9 ± 1.2 mg QE/g, respectively. Thus, regarding total flavonoid content, UA had a higher content than SH (Table 1).

2.2. DPPH Radical Scavenging Effects

The DPPH radical scavenging activity of the SH and UA ethanol extracts was evaluated. The SH and UA extracts were prepared at concentrations of 0.01, 0.1, 1, and 10 mg/mL, and the radical scavenging activity was measured. After that, the IC₅₀ values were 0.64 ± 0.29 mg/mL for SH and 1.58 ± 0.17 mg/mL for UA. Both extracts showed lower activity than L-ascorbic acid, used as a positive control. However, when SH and UA were compared, the SH extract was found to have higher radical scavenging activity (Table 2).

Table 2. DPPH radical scavenging activity of *Sargassum horneri* and *Ulva australis* ethanol extracts.

	<i>S. horneri</i>	<i>U. australis</i>	Ascorbic Acid
DPPH (IC ₅₀ mg/mL)	0.64 ± 0.29	1.58 ± 0.17	0.03 ± 0.00

Data are expressed as the mean \pm standard deviation (n = 3).

2.3. Cell Viability of 3T3-L1 Cells

To evaluate the potential cytotoxic effect of the SH and UA extracts on 3T3-L1 cells, the extracts were prepared at concentrations of 62.5, 125, 250, 500, and 1000 μ g/mL and used to treat 3T3-L1 cells for 48 h. Treatments with either extract at a concentration of 1000 μ g/mL resulted in a cell viability of 95%, whereas, at 500 μ g/mL, cell viability was 100% (Figure 1). Thus, we confirmed that neither extract exhibited cytotoxicity at concentrations lower than 500 μ g/mL. For subsequent analyses, the highest concentration used was 500 μ g/mL.

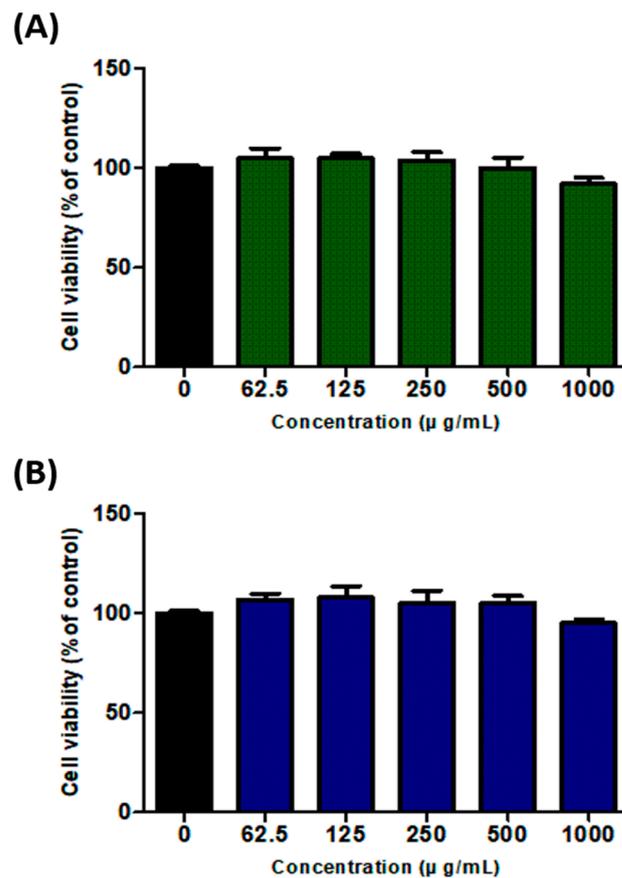


Figure 1. Effects of *S. horneri* and *U. australis* extracts on 3T3-L1 cell viability. (A) *S. horneri* ethanol extract; (B) *U. australis* ethanol extract. The extracts were diluted to 0, 62.5, 125, 250, 500, and 1000 µg/mL and used to treat 3T3-L1 cells for 48 h. Cell viability was measured using the WST-1 assay. Data are expressed as the mean ± standard deviation (n = 3).

2.4. Inhibition of Adipogenesis in 3T3-L1 Cells

To confirm the inhibitory effects of SH and UA extracts on adipocyte differentiation in 3T3-L1 cells, the extracts were applied for 8 d at concentrations of 250 and 500 µg/mL, and Oil Red O staining was performed. In the cells treated with 250 and 500 µg/mL of SH extract, lipid accumulation was inhibited by 94.4% and 84.9%, respectively, compared to the control cells. In the case of the UA extract, 93.5% and 83.0% of the Oil Red O elution amounts were observed in each concentration. Importantly, treatment with the two extracts at a concentration of 500 µg/mL significantly inhibited the differentiation of 3T3-L1 cells compared to control cells (Figure 2).

2.5. Analysis of TG Content in 3T3-L1 Cells

To investigate the effect of SH and UA extracts on modifying TG content, which is a significant marker of adipogenesis, 3T3-L1 cells were treated with the extracts for 8 d and then analyzed. The TG content in the control cells was measured as 904.2 ± 35.4 mg/g of protein. However, in the cells treated with 250 and 500 µg/mL of SH extract, they amounted to 739.7 ± 26.7 and 688.4 ± 22.6 mg/g of protein, respectively; these differences were statistically significant ($p < 0.01$). Similarly, significant differences ($p < 0.01$) were noted in the case of the UA extract, with 705.9 ± 20.9 and 651.3 ± 29.0 mg/g of protein measured in the cells treated with 250 and 500 µg/mL of UA extract, respectively. Therefore, the results confirmed that the cells treated with both extracts showed inhibition of TG content in a concentration-dependent manner compared to the control cells (Figure 3).

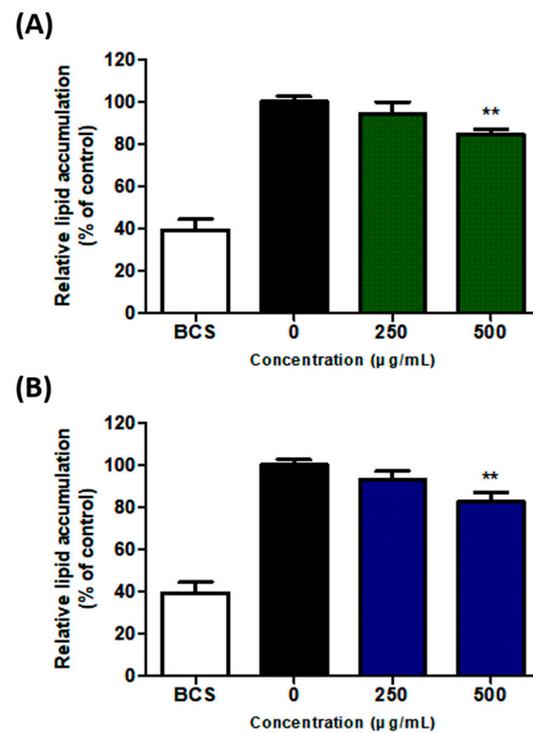


Figure 2. Inhibition of adipocyte differentiation in 3T3-L1 cells treated with *S. horneri* and *U. australis*. (A) *S. horneri* ethanol extract; (B) *U. australis* ethanol extract. After an 8-day differentiation process, the Oil Red O stain was used; the dye was eluted using 100% isopropanol, and optical density (OD) at 500 nm was measured. Data are expressed as the mean \pm standard deviation ($n = 3$). Significant differences: ** $p < 0.01$ vs. control cells.

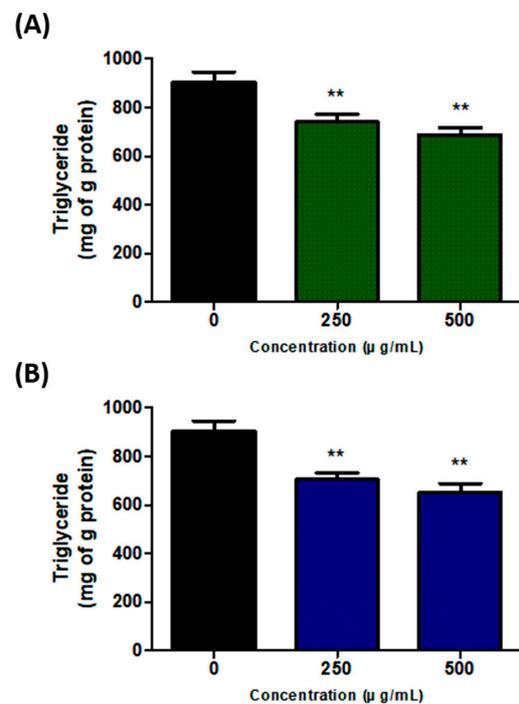


Figure 3. Effects of *S. horneri* and *U. australis* treatment on triglyceride (TG) inhibition during adipocyte differentiation. (A) *S. horneri* ethanol extract; (B) *U. australis* ethanol extract. After 8 d of differentiation, the protein was extracted, and the intracellular TG content was determined using enzyme colorimetry. Data are expressed as the mean \pm standard deviation ($n = 3$). Significant differences: ** $p < 0.01$ vs. control cells.

2.6. Effect on Adipogenic Gene Expression in 3T3-L1 Cells

To examine whether SH and UA inhibit the expression of adipogenic transcription factors, 3T3-L1 preadipocytes were treated with the extracts at various concentrations (250 and 500 $\mu\text{g/mL}$), cultured for 8 d, and analyzed. The SH and UA extracts inhibited the expressions of *C/EBP α* , *PPAR γ* , *SREBP1c*, and *FAS* genes (Figure 4). Of the SH and UA extracts, the expression of adipogenic transcription factors was more highly inhibited at a concentration of 500 $\mu\text{g/mL}$ of the UA extract.

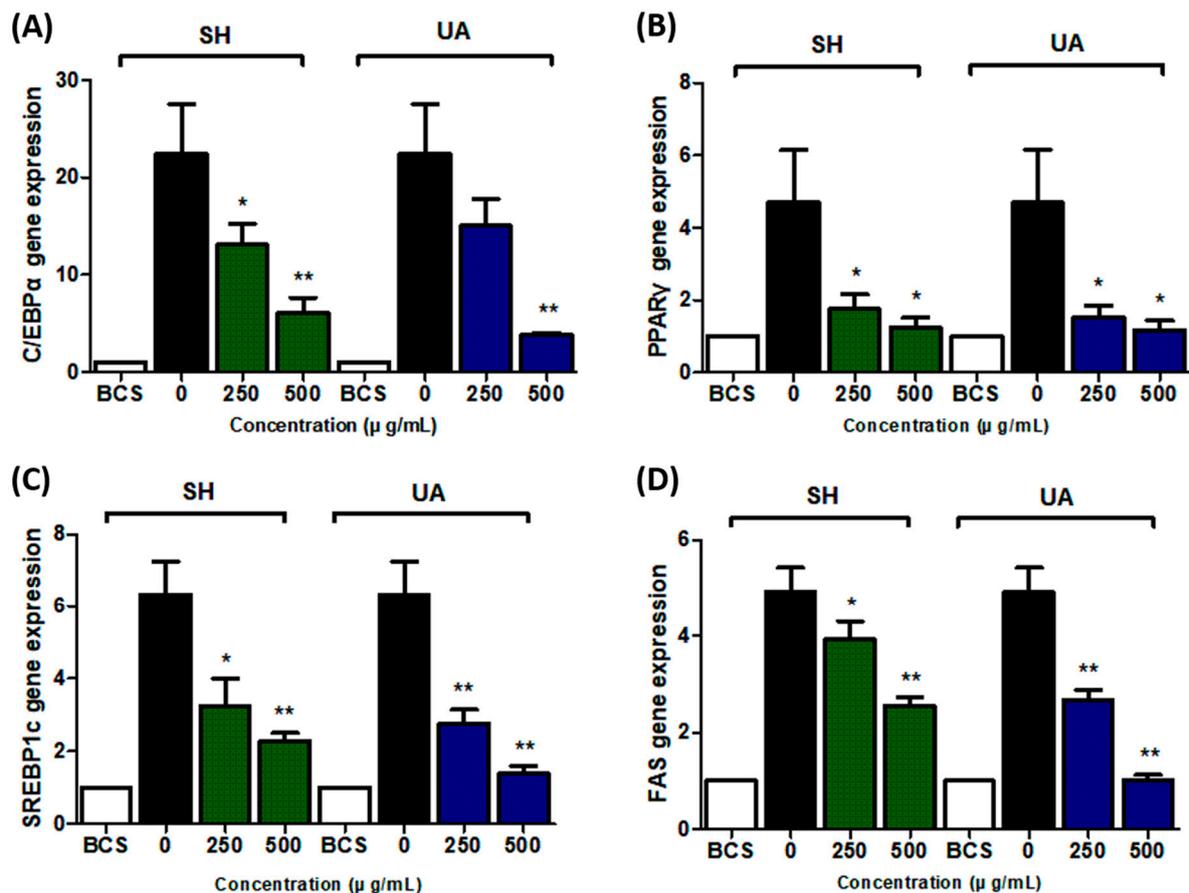


Figure 4. Effects of *S. horneri* and *U. australis* on adipogenic mRNA expression in 3T3-L1 cells. (A) *C/EBP α* , (B) *PPAR γ* , (C) *SREBP1c*, and (D) *FAS*. 3T3-L1 preadipocytes were differentiated in the presence of different concentrations (250 and 500 $\mu\text{g/mL}$) of SH and UA for 8 d. The mRNA expression level was measured using qRT-PCR and normalized using that of the housekeeping gene *GAPDH*. Data are expressed as the mean \pm standard deviation ($n = 3$). Significant differences: * $p < 0.05$, and ** $p < 0.01$ vs. control cells.

2.7. Changes in Body Weights and Food and Drink Intake of Mice

The anti-obesity effect was analyzed by administering SH and UA extracts in drinking water at 1% concentration together with a high-fat diet (HFD). In the group administered SH and UA in the drinking water, the rate of weight gain was lower than that in the HFD group (Figure 5A). The analysis of the weight gain showed that the SH and UA extract administration groups had a weight gain of 21.1 ± 2.1 and 19.6 ± 1.3 g, respectively, which were lower than the 24.9 ± 2.0 g in the HFD group. Significantly lower rates of weight gain were measured in all the groups that received the extracts, and of the two extracts, UA showed a lower rate of weight gain (Figure 5B).

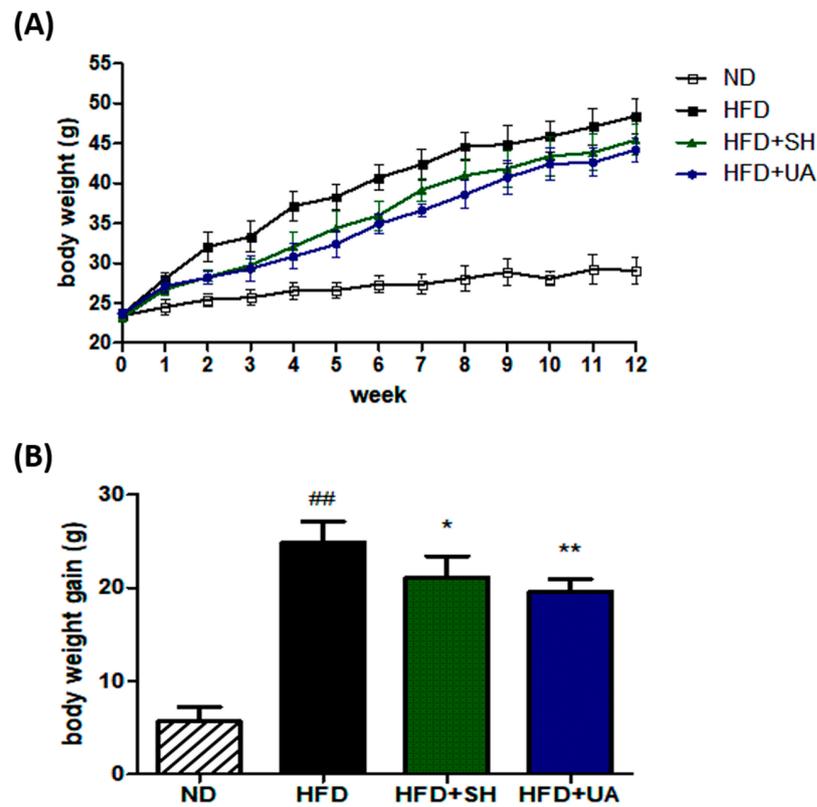


Figure 5. Effects of *S. horneri* and *U. australis* on body weight in C57BL/6 mice fed a high-fat diet for 12 weeks. (A) Weekly changes in body weight; (B) body weight gain. ND, normal diet; HFD, high-fat diet; HFD+SH, administration of 1% *S. horneri* extract in the drinking water and HFD; HFD+UA, administration of 1% *U. australis* extract in the drinking water and HFD. Each value represents the mean \pm standard deviation ($n = 6$). Significant differences: * $p < 0.05$, ** $p < 0.01$ vs. the HFD group, and ## $p < 0.01$ vs. the ND group.

Food and drink intake were analyzed to exclude the possibility that the extracts led to decreases in food or drinking water intake. Food and drink intake were measured at the same time every week and the results were analyzed. The results revealed no differences in food and drink intake between the HFD group and the two extract administration groups; we thus concluded that administration of the extracts had a weight loss effect (Table 3).

Table 3. Changes in food and drink intake in C57BL/6 mice during 12 weeks of HFD-induced obesity.

Groups	Food Intake (g/Day)	Drink Intake (mL/Day)
ND	2.5 \pm 0.3	4.9 \pm 0.6
HFD	2.3 \pm 0.1	5.2 \pm 0.5
HFD+SH	2.3 \pm 0.1	4.9 \pm 0.6
HFD+UA	2.2 \pm 0.1	5.0 \pm 0.4

Data are expressed as the mean \pm standard deviation ($n = 6$).

2.8. Serum Biochemical Analysis

The results of the serum biochemical analysis showed that aspartate aminotransferase (AST) and alanine aminotransferase (ALT), which are liver function test components, tended to increase in the HFD group compared to those in the normal group. However, AST and ALT levels in the group administered the two types of seaweed extracts significantly decreased compared to those in the HFD group. Gamma Glutamyl Transferase (GGT) levels did not identify significant differences in all groups. The level of total cholesterol (CHO) was high at 193.7 \pm 11.3 mg/dL in the HFD group compared to that in the normal

group. However, it significantly decreased to 142.4 ± 6.6 and 156.3 ± 9.7 mg/dL in the SH and UA extract administration groups, respectively. TG levels were 50.9 ± 6.0 , 32.6 ± 6.1 , and 39.0 ± 5.1 mg/dL, in the HFD, SH, and UA groups, respectively; TG levels significantly decreased in the groups treated with the extracts compared to those in the HFD group. HDL cholesterol (HDL-C) levels tended to increase in the high-fat diet group compared to the normal diet group. LDL cholesterol (LDL-C) levels, analyzed by total cholesterol, triglyceride, and HDL cholesterol levels, were higher in the HFD group at 71.9 ± 4.2 mg/dL compared with the normal group. However, in the SH and UA extract treatment groups, they were significantly reduced to 40.7 ± 4.4 and 40.8 ± 3.1 mg/dL, respectively. Blood glucose (GLU) levels, too, significantly decreased in the experimental groups compared to those in the HFD group (Table 4).

Table 4. Changes in serum biochemical parameters in obese C57BL/6 mice induced by a high-fat diet.

	ND	HFD	HFD+SH	HFD+UA
AST (U/L)	211.4 ± 16.0	312.8 ± 15.7 #	244.7 ± 16.5 *	217.2 ± 11.1 *
ALT (U/L)	55.4 ± 8.1	138.5 ± 15.4 #	114.2 ± 12.1 *	89.8 ± 8.8 *
GGT (U/L)	0.8 ± 0.6	0.5 ± 0.5	0.6 ± 0.4	0.7 ± 0.4
CHO (mg/dL)	113.3 ± 6.3	193.7 ± 11.3 #	142.4 ± 6.6 *	156.3 ± 9.7 *
TG (mg/dL)	34.1 ± 5.1	50.9 ± 6.0 #	32.6 ± 6.1 *	39.0 ± 5.1 *
HDL-C (mg/dL)	71.0 ± 3.7	111.8 ± 8.9 #	95.0 ± 9.3	107.7 ± 10.2 #
LDL-C (mg/dL)	35.1 ± 7.4	71.9 ± 4.2 #	40.7 ± 4.4 *	40.8 ± 3.1 *
GLU (mg/dL)	204.8 ± 5.9	281.2 ± 9.3 #	231.2 ± 8.1 *	223.7 ± 6.9 *

AST, aspartate aminotransferase; ALT, alanine aminotransferase; GGT, gamma-glutamyl transferase; CHO, total cholesterol; TG, triglyceride; HDL-C, HDL cholesterol; LDL-C, LDL cholesterol; GLU, glucose. Each value represents the mean ± standard deviation (n = 6). Significant differences: * $p < 0.05$ vs. HFD group, and # $p < 0.05$ vs. ND group.

2.9. Weight Analysis of Liver and White Adipose Tissue

The comparison of the weights of liver, epididymal fat, mesenteric fat, and retroperitoneal fat between the ND and HFD groups demonstrated that the weights of all tissues were significantly higher in the HFD group. The comparison of the epididymal fat mass between the HFD group and the extract-administered groups showed that the HFD SH and UA groups had significantly lower values than the ND group (3.01 ± 0.42 , 2.25 ± 0.24 , and 2.32 ± 0.17 g, respectively). Comparison of the weights of the liver, retroperitoneal fat, and mesenteric fat showed that tissue weights in the extract-administered groups were lower than those in the obese group; however, this difference did not reach statistical significance (Table 5). In the comparison between the two extract-administered groups, most of the tissue weights were lower in the UA-administered group than in the SH-administered group.

Table 5. Changes in liver and white adipose tissue weight in obesity prevention test animals.

	ND	HFD	HFD+SH	HFD+UA
Liver	1.12 ± 0.07	1.80 ± 0.18 [#]	1.54 ± 0.29	1.44 ± 0.27
Epididymal	0.59 ± 0.13	3.01 ± 0.42 [#]	2.25 ± 0.24 [*]	2.32 ± 0.17 [*]
Mesenteric	0.59 ± 0.06	2.00 ± 0.29 [#]	1.66 ± 0.46	1.40 ± 0.45
Retroperitoneal	0.24 ± 0.08	1.83 ± 0.26 [#]	1.54 ± 0.20	1.48 ± 0.19

Each value represents the mean ± standard deviation (n = 6). Significant differences: ^{*} $p < 0.05$ vs the HFD group, and [#] $p < 0.05$ vs. the ND group.

2.10. Histopathological Analysis of Liver and Adipose Tissues

Histological analysis of liver and adipose tissues was performed using hematoxylin and eosin (H&E) staining. Regarding the liver tissue, we observed that the number of fat droplets increased in the HFD group compared to those in the normal group. The number and size of lipid droplets in the group administered the SH and UA extracts tended to decrease compared to those in the HFD group (Figure 6). In addition, in the white adipose tissue, the size of adipocytes in the HFD group was enlarged compared to that in the normal group, and adipocyte size in the group administered the SH and UA extracts tended to decrease (Figure 7).

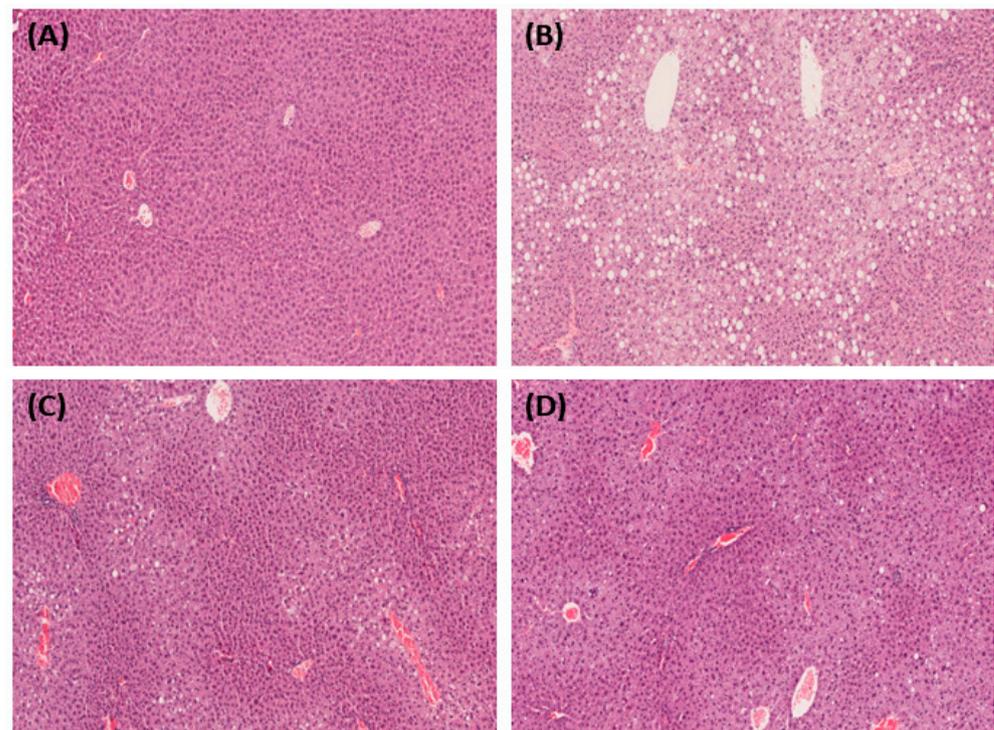


Figure 6. Microphotographs of the liver in HFD-fed C57BL/6 mice showing the effect of treatment with *S. horneri* and *U. australis*. Representative liver sections stained for hematoxylin and eosin (H&E) (magnification × 200). (A) ND; (B) HFD; (C) HFD+SH; and (D) HFD+UA.

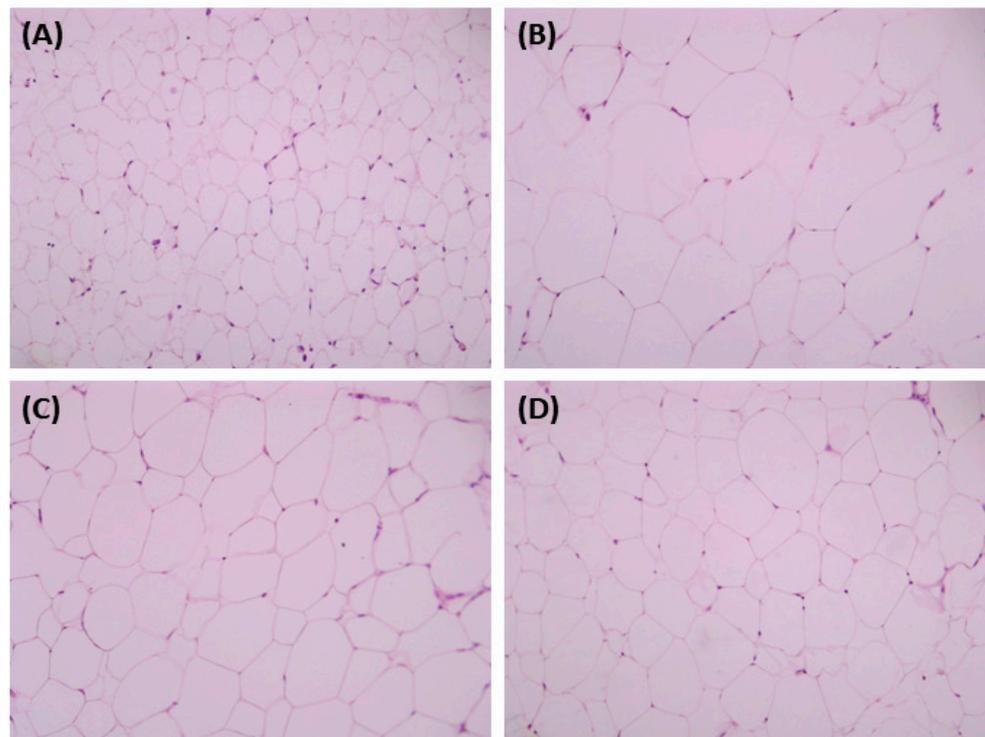


Figure 7. Microphotographs of the adipose tissue in HFD-fed C57BL/6 mice showing the effect of treatment with *S. horneri* and *U. australis*. Representative sections of adipose tissue stained for H&E (magnification $\times 200$). (A) ND; (B) HFD; (C) HFD+SH; and (D) HFD+UA.

2.11. Bioactive Compounds Present in the Extracts

The bioactive compounds present in the SH and UA extracts are shown in Table 6. The main compounds present in the SH extract were analyzed as Neophytadiene, 3,7,11,15-Tetramethyl-2-hexadecen-1-ol, and Hexadecanoic acid ethyl ester. The main compound present in the UA extract was analyzed as 2-Hexadecen-1-ol, 3,7,11,15-tetramethyl-,acetate, [R-[R*,R*-(E)]]-.

Table 6. Compounds isolated from *S. horneri* and *U. australis* extracts.

Extracts	RT	Name of Compounds (Molecular Formula)	Area (%)
<i>S. horneri</i>	22.8	Neophytadiene (C ₂₀ H ₃₈)	25.3
	23.1	3,7,11,15-Tetramethyl-2-hexadecen-1-ol (C ₂₀ H ₄₀ O)	14.0
	24.4	Hexadecanoic acid, ethyl ester (C ₁₈ H ₃₆ O ₂)	24.6
<i>U. australis</i>	23.3	2-Hexadecen-1-ol, 3,7,11,15-tetramethyl-,acetate, [R-[R*,R*-(E)]]-(C ₂₂ H ₄₂ O ₂)	40.4

3. Discussion

Obesity is a chronic metabolic disorder that threatens the health of people in developed and developing countries due to excessive body fat accumulation [30]. Obesity is related to various diseases, such as diabetes, hypertension, hyperlipidemia, stroke, and angina, and is known to be a major cause of aging [31]. Accordingly, prevention and treatment of obesity are important. To treat obesity, a combination of diet and drug approaches has been used. However, orlistat and rimonabant, commercially available drugs to treat obesity, have limitations in their use in children, patients with myocardial infarction, liver disease, and kidney disease, as well as various side effects, such as headache, abdominal pain, and panic disorder [32,33]. In this context, interest in food products based on natural substances,

such as anti-obesity substances, oriental medicines, and medicinal plants, has been steadily growing [34].

Seaweed has been widely used as diet food due to its low fat and calorie content and the presence of various essential nutrients, such as protein, vitamins, and minerals whose deficiency is commonly noted during dieting [13]. Most seaweeds have been used as food and functional additives; however, SH and UA are not consumed in Korea and have been classified as marine organic waste resources causing coastal pollution on Jeju Island [35]. In the present study, extracts were prepared using 80% ethanol to upcycle these two seaweed waste resources into functional materials. Their antioxidant effect was then identified, and anti-obesity effects were confirmed *in vitro* and *in vivo*.

The antioxidant activities of SH and UA extracts were analyzed based on their DPPH radical scavenging activity. DPPH radical scavenging activity measures antioxidant activity via a reduction process in amines and aromatic compounds due to the electron-donating ability of antioxidants, and the degree of purple discoloration has previously been used as an indicator [36]. Using this method, we confirmed that the SH and UA extracts show free radical scavenging activity in a concentration-dependent manner. Additionally, the SH extract was found to have a higher scavenging activity than the UA extract. Phenolic compounds play an essential role as antioxidants, possess phenolic hydroxyl groups, which are collectively referred to as flavonoids, anthocyanins, tannins, and catechins, and are widely distributed in the plant kingdom [37]. Flavonoids are pigments in various plants that have been reported to have antioxidant, UV-blocking, anti-mutagenic, antiviral, and anti-inflammatory effects [38]. In the present study, the total polyphenol contents in the SH and UA extracts were compared, and SH was found to have a higher content than UA (95.5 and 62.4 mg GAE/g, respectively). This finding is consistent with previous research showing that brown algae have a higher polyphenol content than green and red algae [39]. In addition, when we compared the flavonoid contents in the two materials, we found that SH was 19.4 and UA 20.9 mg QE/g, indicating that the two materials were analyzed similarly.

To confirm whether the SH and UA extracts affect adipocyte formation, they were applied at concentrations equal to or lower than 500 $\mu\text{g}/\text{mL}$, which do not exhibit cytotoxicity during the differentiation process. The results confirmed that adipocyte differentiation was inhibited in a concentration-dependent manner. During the differentiation of preadipocyte 3T3-L1 cells, fat is accumulated in the cells, mostly as neutral fat [40]. The TG content in the cells decreased in a concentration-dependent manner. After treatment with SH and UA extracts at a concentration of 500 $\mu\text{g}/\text{mL}$, TG content decreased by more than 30%. A previous study on the inhibition of adipocyte differentiation using SH reported that TG content decreased by 23% in the experimental group treated with SH at 1 mg/mL for 8 d [41]. In contrast, in our results, TG content was considerably reduced when the extract was applied at a concentration of 500 $\mu\text{g}/\text{mL}$. This difference may be due to differences in the drying method, extraction solvent, and seaweed extraction method.

Preadipocytes were treated with the 3-isobutyl-1-methylxanthine (IBMX) complex, and specific genes were expressed during the differentiation process into adipocytes. Several transcription factors, including *PPAR γ* , *C/EBP α* , and *SREBP1c*, play a pivotal role in regulating the differentiation and accumulation of adipocytes [42,43]. *PPAR γ* is a predominantly liver-distributed transcription factor that, when activated, produces marked changes in fat morphology, fat accumulation, and insulin sensitivity [44]. *C/EBP α* plays an important role in adipogenesis. Expressions of *PPAR γ* and *C/EBP α* are mutually regulated at the preadipocyte differentiation stage, and the two genes cooperate to promote adipogenesis and regulate the expression of *SREBP1c* and *FAS* [45,46]. The results of our analysis confirmed that the expression of transcription factors of adipogenesis was downregulated in a concentration-dependent manner. In addition, evidence suggests that differentiation into adipocytes is inhibited by the suppression of *FAS* expression [47], an important metabolic enzyme acting on adipogenesis and lipogenesis. Therefore, our results confirmed that the two seaweed extracts function as suppressors of the expression of key transcription

factors that act during the fat differentiation process and that UA regulates the expression of transcription factors more effectively than SH.

The HFD mouse model has been widely used to investigate anti-obesity effects and for the development of anti-obesity drugs [48]. Obese mice fed an HFD are characterized by increased body weight and adipocyte size, fatty liver development, and increased plasma GLU and TG levels [47]. In the present study, to confirm the effect of preventing obesity in C57BL/6 mice induced by an HFD, the extract was added to drinking water at 1% concentration and administered for 12 weeks. In general, extracts have been administered mainly via the oral route, but they may also be supplied through feed or water [26,49]. Since the extracts were added to the drinking water, we investigated potential differences between feed and drinking water intake. The results showed no differences in diet and water intake between the HFD and the extract administration groups, and no issues were observed regarding palatability. The weight gain rate significantly decreased in the group administered the SH and UA extracts compared to that in the HFD group. A previous obesity study using SH showed a protective effect against obesity when the substance was added to a high-fat diet at 2% and 6% and fed to mice [26]. However, this study found that adding the extract to drinking water at a concentration of 1% showed anti-obesity effects despite a lower treatment concentration than the previous results. Furthermore, the group that was administered the UA extract had a lower rate of weight gain than the SH group.

The serological analysis revealed that the levels of AST, ALT, CHO, TG, and blood GLU, which are related to liver function, were significantly reduced in the extract-administered groups compared to those in the HFD group. According to previous research results, Sargassaceae are rich in fucoxanthin, which has anti-obesity and anti-diabetic effects, while UA contains polysaccharides and dietary fiber, which is known to help with hypercholesterolemia [21,50,51]. In our results, we reasoned that the reduction in serological values in the two experimental groups compared to those in the obese group was due to the components in the SH and UA extracts. In addition, the results confirmed that accumulation of fat droplets in liver tissue was suppressed after administration of SH and UA extracts; similarly, the size of adipocytes in white adipose tissue was also suppressed. These results confirmed that SH and UA have antioxidant properties and can prevent fatty liver, TG, and blood GLU increases in obesity. Based on this evidence, we conclude that SH and UA have obesity-preventive effects.

Phytochemicals with traditional medicinal properties have been analyzed using various chemical analysis methods. Of the three biologically active compounds analyzed in SH extracts, Neophytadiene is known to have antimicrobial, antioxidant, and anti-inflammatory effects [52], and Hexadecanoic acid ethyl ester is known to have antioxidant, hypocholesterolemic, nematocidal, and pesticide effects [53]. Phytol and phytyl acetate were analyzed in SH and UA extracts, which are not only diterpene compounds [54] but have been shown in various studies to have nitric oxide inhibitory activity as well as plant-growth-regulating, hypolipidemic, antimicrobial, and antiviral properties, among others [55]. The antioxidant and anti-obesity effects of SH and UA extracts are thought to be due to the effects of the compounds analyzed and the components present in the seaweed, such as dietary fiber.

Therefore, the results of this study suggest that SH and UA, typically considered marine waste resources, can be effectively used as functional additive candidates for obesity prevention. In addition, our results can be considered primary data for the recycling of seaweed waste.

4. Materials and Methods

4.1. Materials

Dulbecco's modified Eagle's medium (DMEM), bovine calf serum (BCS), fetal bovine serum (FBS), and penicillin–streptomycin (PS) were purchased from Welgene (Daegu, Republic of Korea). Folin & Ciocalteu phenol reagent, IBMX, insulin from bovine pancreas, gallic acid, Oil Red O, and isopropanol were obtained from Sigma-Aldrich (St. Louis,

MO, USA). Dexamethasone was purchased from the Tokyo Chemical Industry (Tokyo, Japan). The Cellvia cell viability assay kit was obtained from AbFrontier (Seoul, Republic of Korea). The TG colorimetric assay kit was purchased from Asan pharmaceuticals (Seoul, Republic of Korea). The AccuPrep[®] Universal RNA Extraction Kit was supplied by Bioneer (Daejeon, Republic of Korea). The One-Step TB Green[®] PrimeScript[™] reverse transcriptase polymerase chain reaction (RT-PCR) and Bradford Protein Assay kits were purchased from Takara (Tokyo, Japan). The Korean Cell Line Bank (KCLB; Seoul, Republic of Korea) provided the 3T3-L1 cells (KCLB: 10092.1). All chemicals and solvents used in the present study were of analytical grade.

4.2. Preparation of *S. horneri* and *U. australis* Extracts

SH and UA were collected from the coast of Jeju Island at the Jeju Techno Park Biodiversity Research Center, washed and dried, and then grinded into powder. After immersing 50 g of the dried sample in 1 L of 80% ethanol, the extract was prepared by being stirred at room temperature for 24 h. The extracted material was filtered using a 0.45 µm bottle-top vacuum filter (Corning Inc, Corning, NY, USA). The filtered material was lyophilized after the ethanol was volatilized using a rotary evaporator (LabSciTech, Corona, CA, USA) to prepare the extract powder. Sample extraction was performed by repeating the process twice. The prepared sample was stored at −20 °C and used after being suspended in a cell culture medium or sterile distilled water.

4.3. Analysis of Total Phenol and Flavonoid Content

Total phenol content was measured using the Folin–Denis method [56]. One milliliter of the extract at a concentration of 1 mg/mL was mixed with 1 mL of Folin–Ciocalteu reagent and allowed to stand for 3 min. Next, 1 mL of 10% Na₂CO₃ solution was added and reacted at room temperature for 1 h. The absorbance at 760 nm was measured using a microplate reader (Varioskan LUX). The procedure was repeated using the standard substance gallic acid to prepare the calibration curve ($y = 0.0025x$; $R^2 = 0.9933$). Total phenol content was expressed as mg GAE/g of extract sample.

Total flavonoid content was quantified using the diethylene glycol colorimetric method [57]. One milliliter of diethylene glycol and 100 µL of 1 N NaOH were mixed with 100 µL of SH and UA extracts (1 mg/mL), blocked from light, and reacted at 37 °C for 1 h. The absorbance at 420 nm was measured using a microplate reader (Varioskan LUX). The procedure was repeated using the standard substance quercetin to prepare a calibration curve ($y = 0.0018x - 0.0019$; $R^2 = 0.9977$). Total flavonoid content was expressed as mg QE/g of extract sample.

4.4. DPPH Radical Scavenging Assay

The antioxidant activity of SH and UA extracts was evaluated using the DPPH (1,1-diphenyl-2-picrylhydrazyl) method [58]. Each extract was prepared at concentrations of 0.01, 0.1, 1, and 10 mg/mL using ethanol. Next, 190 µL of a substance of 10 µL and a 0.2 mM DPPH solution were added to a 96-well plate and mixed. The reaction was performed at 37 °C for 30 min, and the absorbance at 550 nm was measured using a microplate reader (Varioskan LUX, Thermo Fisher Scientific, Waltham, MA, USA). DPPH radical scavenging ability was calculated according to the following equation:

$$\text{DPPH free radical scavenging effect (\%)} = [1 - (\text{absorbance of experimental group} / \text{absorbance of control group})] \times 100$$

Half the maximum inhibitory concentration (IC₅₀) was calculated using linear regression analysis and expressed as the mean of three measurements.

4.5. Cell Culture and Differentiation of 3T3-L1 Adipocyte Cells

3T3-L1 cells, a white adipocyte-like cell line, were cultured at 37 °C and 5% CO₂ by adding 1% PS and 10% BCS to DMEM. The cells were cultured to approximately 70% confluency and then periodically passaged and used for experiments. For differentiation,

the 3T3-L1 cells were first cultured using DMEM supplemented with 10% BCS until they reached confluence. After 2 d of confluence, the differentiation induction medium containing 10% FBS, 1% PS, 0.5 μ M IBMX, 1 μ M dexamethasone, and 10 μ g/mL insulin in DMEM was replaced, and differentiation was induced for 48 h. The cells were then cultured for 6 d using DMEM containing 10% FBS, 1% PS, and 10 μ g/mL of insulin; the culture medium was replaced every 2 d. The extract samples were applied from the beginning of differentiation and were used for further analysis after 8 d of treatment.

4.6. Cell Viability Assay

A water-soluble tetrazolium-1 (WST-1) assay was used to measure the cytotoxicity of the extracts and determine the range of concentrations to be used in the subsequent experiment. 3T3-L1 preadipocytes were dispensed at 1×10^5 cells/mL in 96-well plates and cultured for 24 h at 37 °C and 5% CO₂. After sample treatment, Cellvia solution (WST-1) was added at 1/10 volume of the medium, and the cells were incubated for 30 min at 37 °C in a 5% CO₂ incubator. The absorbance at 450 nm was measured using a microplate reader (Varioskan LUX). Subsequent experiments were performed in a concentration range that did not induce toxicity.

4.7. Oil Red O Staining of 3T3-L1 Adipocytes

Adipogenesis in 3T3-L1 cells was measured using Oil Red O staining. Differentiated 3T3-L1 cells were washed thrice with 1 \times phosphate-buffered saline (PBS, pH 7.4) and then fixed with 10% formaldehyde for 1 h at room temperature. After the cells were washed twice with distilled water (DW), they were treated with 60% isopropanol for 5 min at room temperature. Next, the isopropanol was removed, and the cells were allowed to dry at room temperature. A solution of 500 mg Oil Red O dissolved in 100 mL of 2-propanol was mixed with distilled water at a ratio of 6:4 and filtered through a 0.45 μ m filter. The Oil Red O solution was added to the plates, and the cells were incubated for 1 h at room temperature. After staining, the cells were washed 4 times with DW, and the dye was eluted using 100% isopropanol. The optical density (OD) at 500 nm was then measured using a spectrophotometer (Milton Roy Company, New York, NY, USA).

4.8. Measurement of Intracellular Triglyceride Levels

After adipocyte differentiation, 3T3-L1 cells were washed with cold 1 \times PBS. After adding DW, the cells were collected using a cell scraper, and chloroform and methanol were added to lyse the cells. The cell lysate was incubated in a shaking incubator at 37 °C for 1 h and then centrifuged at 3000 rpm for 5 min. After transferring the bottom layer to an Eppendorf tube, the lid was opened and left for more than 24 h; the samples for analysis were prepared by adding 20 μ L of 100% ethanol. Intracellular TG content was measured using the TG colorimetric assay kit. The cellular protein concentration was measured using the Bradford Protein Assay Kit, and the TG content was normalized to that of cellular protein.

4.9. Total RNA Extraction and Quantitative (q)RT-PCR

Differentiated preadipocytes were washed with 1 \times PBS, and total RNA was extracted using an Accuprep[®] universal RNA extraction kit (Bioneer). The extracted RNA was quantified using NanoDrop 2000 (Thermo Fisher Scientific). The qRT-PCR was performed using a One-Step TB Green[®] PrimeScript[™] RT-PCR Kit (Takara) with 1 μ g of RNA on a 7500 Fast Real-time PCR system (Applied Biosystems, Waltham, MA, USA). The conditions were 42 °C for 50 s, 95 °C for 10 s, followed by 40 cycles of 95 °C for 5 s, 60 °C for 30 s, and 72 °C for 30 s. The primers used in this laboratory are listed in Table 7 [1,2]. The expression of target genes was normalized to that of the housekeeping gene glyceraldehyde-3-phosphate dehydrogenase (GAPDH). Changes in gene expression were calculated using the $2^{-\Delta\Delta CT}$ method, and the relative mRNA amount was quantified.

Table 7. Target genes and their primer sequences used for qRT-PCR.

Gene	GenBank Accession Number	Forward Primer (5' to 3')	Reverse Primer (5' to 3')	Amplicon (bp)
<i>SREBP1c</i> ¹	NM_011480.4	GCG CTA CCG GTC TTC TAT CA	TGT GTG CAC TTC GTA GGG TC	176
<i>PPARγ</i> ²	NM_011146.3	CCG TGC AAG AGA TCA CAG AG	GGC CCT CTG AGA TGA GGA C	159
<i>C/EBPα</i> ³	NM_007678.3	GCT GGA GTT GAC CAG TGA CA	CCT TGA CCA AGG AGC TCT CA	116
<i>FAS</i> ⁴	NM_007988.3	CTA CCA GGC CAT CCG TAG TG	ACA ATA TCC ACT CCC TGA ATC	157
<i>GAPDH</i> ⁵	NM_008084.3	CCC CTC TGG AAA GCT GTG G	ACA TTG GGG GTA GGA ACA CG	150

¹ *SREBP1c*, Sterol regulatory element binding protein 1c; ² *PPAR γ* , peroxisome proliferator activated receptor gamma; ³ *C/EBP α* , CCAAT/enhancer-binding protein alpha; ⁴ *FAS*, fatty acid synthase; ⁵ *GAPDH*, glyceraldehyde 3-phosphate dehydrogenase.

4.10. Animal Experiment

Five-week-old male C57BL/6 mice were purchased from KOSA BIO Inc. (Sunnam, Republic of Korea) and were housed in specific-pathogen-free conditions. During the experiment, the mice were maintained in a stable breeding environment with a light/dark cycle of 12 h at a temperature of 23 ± 3 °C, relative humidity of $50 \pm 10\%$, and 150–300 Lux. A standard diet (Samtako Bio Korea, Seoul, Republic of Korea) and water were freely consumed by the mice for one week before the experiment, while the animals were allowed to adapt to the environment. The C57BL/6 mice were divided into the following four groups (six animals per group): (1) control group fed a normal diet; (2) HFD+DW group; (3) HFD + 1% SH drinking administration group; (4) HFD + 1% UA drinking administration group. The experiment was conducted for 12 weeks, and the mice in all HFD groups were fed a diet that contained 60% kcal from fat purchased from Research Diets (catalog number D12492: protein 20% kcal, fat 60% kcal, carbohydrate 20% kcal, energy density 5.21 kcal/g; New Brunswick, NJ, USA). This study was approved by the Animal Experimentation Ethics Committee of the Catholic University of Pusan (CUP AEC 2021-001).

4.11. Analysis of Body Weight Change and Food and Drink Intake

To analyze the changes in the weight of the experimental animals, weight was measured using an electronic balance once a week at a fixed time (10:00 A.M.) during the experimental period. Body weight gain was expressed as a percentage by comparing the weight measured at the beginning and end of the experimental period. Food and drink intake were also measured once a week at a fixed time. The daily dietary and liquid intake per animal were calculated by measuring the serving and remaining amounts of food and water.

4.12. Serum Biochemical Analysis

The experimental animals were fasted for 12 h before necropsy, and a mixture of 25 mg/kg of alfaxalone (Alfaxan[®]; Careside, Seongnam-si, Republic of Korea) and 5 mg/kg of xylazine (Rompun[®]; Bayer Korea, Seoul, Republic of Korea) was used as anesthetic. After anesthesia, blood was collected from the heart and coagulated at room temperature for serum biochemical tests. The serum obtained after centrifugation at 2500 rpm for 30 min was analyzed using a blood biochemical analyzer (BT 1500; Biotechnica Instruments SpA, Roma, Italy). The levels of the following components were analyzed: AST, ALT, GGT, CHO, TG, HDL-C, and GLU. LDL cholesterol was calculated using the Friedewald equation: LDL-cholesterol = total cholesterol—HDL cholesterol—(triglycerides/5).

4.13. Organ Weight and Histopathological Analysis

After deep anesthesia was achieved, the liver, epididymal fat, retroperitoneal fat, and mesenteric fat of the mice were extracted, weighed, and analyzed as absolute organ weights against body weight. The excised organs were washed with physiological saline, and, after removing moisture with a filter paper, a part of the excised tissue was fixed in 10% formalin and embedded in a paraffin block for sectioning (3 μ m thickness) and subsequent

histopathological examination. Tissue sections were stained using H&E and evaluated using a microscope (Olympus BX51; Olympus Optical Co., Tokyo, Japan).

4.14. Gas Chromatography–Mass Spectroscopy Analysis

After GC–MS analysis was performed by sending a request to the Korea Polymer Testing and Research Institute (Koptri, Seoul, Republic of Korea), analysis was performed using Agilent Technologies GC-MS (GC-7890A, MS 5975C) and Agilent J&W DB-5MS Ultra Inert column (30 m × 0.25 mm × 0.25 μM). The GC oven was programmed to heat as follows: An initial temperature of 40 °C was set and held at this temperature for 5 min; then, the oven temperature was increased to 280 °C at a rate of 10 °C /min and held for 5 min. The injector temperature was 250 °C, the flow rate of the carrier gas, helium, was 1.0 mL/min, and the split ratio was 1:10. The mass spectrum scan range was set to 30–500 (*m/z*). The analyzed compounds were compared to the spectra of the samples using the Koptri mass spectral library to identify the compound name, molecular weight, and structure of the test material.

4.15. Statistical Analysis

GraphPad Prism 5.0 software (GraphPad Software Inc., San Diego, CA, USA) was used for data analysis. Values were expressed as the mean ± standard deviation and tested using one-way ANOVA for significant differences; *p* values < 0.05 were considered to indicate significant differences.

5. Conclusions

The results of this study confirmed the antioxidant effects of SH and UA extracts and the inhibition of transcription factors involved in lipid accumulation and adipogenesis in 3T3-L1 cells. They have also been shown to improve anti-obesity effects and serum biochemical markers in mice given a high-fat diet and a 1% concentration of the extract in drinking water. Based on these results, we have obtained basic data suggesting the possibility of their use as a functional additive candidate for obesity prevention that can be applied not only to humans but also to pets. However, further research is necessary to analyze the bioactive compounds of SH and UA, their molecular mechanisms, and the synergistic effects of mixed administration of the two extracts in preventing obesity. However, there is a lack of data analyzing the bioactive compounds and mechanisms of action of SH and UA extracts for their anti-obesity effects. Therefore, it is considered necessary to study the analysis of the bioactive compounds, the mechanisms of action, and the synergistic effects of the mixed administration of the two extracts when studying their anti-obesity effect by preparing fractions of the two extracts.

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