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Running Title: Inhibition of hepatic lipogenesis by hemistepsin A

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ABSTRACT

Hemistepsin A (HsA) is a guaianolide sesquiterpene lactone that inhibits hepatitis and liver fibrosis. We evaluated the effects of HsA on liver X receptor (LXR)-mediated hepatic lipogenesis *in vitro* and *in vivo*. Up to 10 μ M, HsA did not affect the viability of HepG2 and Huh7 cells. Pretreatment with 5–10 μ M HsA significantly decreased the luciferase activity of the LXR response element, which was transactivated by T0901317, GW3965, and LXR α /retinoid X receptor α overexpression. In addition, it significantly inhibited the mRNA expression of *LXR α* in HepG2 and Huh7 cells. It also suppressed the expression of sterol regulatory element-binding protein-1c and lipogenic genes and reduced the triglyceride accumulation triggered by T0901317. Intraperitoneal injection of HsA (5 and 10 mg/kg) in mice significantly alleviated the T0901317-mediated increases in hepatocyte diameter and the percentage of regions in hepatic parenchyma occupied by lipid droplets. Furthermore, HsA significantly attenuated hepatic triglyceride accumulation by restoring the impaired expression of LXR α -dependent lipogenic genes caused by T0901317. Therefore, based on its inhibition of the LXR α -dependent signaling pathway, HsA has prophylactic potential for steatosis.

Keywords: Fatty liver, Hemistepsin A (HsA), Liver X receptor, Sterol regulatory element-binding protein-1c, T0901317

INTRODUCTION

Steatosis (fatty liver) is the pathologic state in which hepatocytes contain more than 5% fat. This can be caused by overeating, alcohol consumption, viral infection, hepatotoxins, and other causes. The prevalence of nonalcoholic fatty liver is increasing worldwide as a result of an increasingly sedentary lifestyle and excess food intake. Although nonalcoholic fatty liver is considered not to cause serious liver damage, it can progress to nonalcoholic fatty liver disease (NAFLD), which involves nonalcoholic steatohepatitis, fibrosis, cirrhosis, and cancer (1). Because there are no approved drugs for advanced NAFLD, it must be appropriately managed at an early stage.

Liver X receptors (LXRs) are type II nuclear receptors that activate *de novo* lipogenesis in the liver (2). In the absence of ligands, LXRs interact with retinoid X receptor (RXR) α and bind to the LXR response element (LXRE), which has a 5'-AGGTCA-3' direct repeat separated by four nucleotides. The conformational change in LXR, which occurs by ligand binding, promotes recruitment of the coactivator complex, resulting in transactivation of target genes. These include sterol regulatory element-binding protein (SREBP)-1c, a transcription factor essential for hepatic lipogenesis (2). In addition, LXR directly or in conjunction with SREBP-1c increases lipogenesis by inducing the production of fatty acid synthase (FASN), stearoyl-CoA desaturase-1 (SCD-1), and acetyl-CoA carboxylase (ACC) (2, 3). Two isoforms of LXRs are expressed in mammals, each with different tissue distributions: LXR α is primarily expressed in metabolically active organs/tissues including the liver, whereas LXR β is ubiquitously expressed (4). Although both LXR α and LXR β are expressed in the liver, the hepatic expression of LXR α is closely associated with the severity of NAFLD in humans (5). In addition, genetic ablation of LXR α impairs T0901317 (a synthetic LXR ligand)-mediated hypertriglyceridemia in mice (6), suggesting that LXR α is the predominant isoform regulating

lipogenesis in the liver. Hence, inhibition of LXR α activation and its triggering of the expression of target genes shows potential as a strategy for managing fatty liver.

Sesquiterpene lactones (STLs) are secondary metabolites widely distributed in the plant kingdom and are anti-feeding agents; that is, they protect plant from herbivory (7). Although all STLs have three isoprenoids with an α -methylene- γ -lactone moiety, more than 5,000 different structures and 30 subtypes have been identified from diverse plants including *Hemistepta lyrata* Bunge (Bunge) (7, 8). Hemistepsin A (HsA) belongs to the guaianolide subtype of STL; it is isolated from *H. lyrata* and shows antifungal activity and cytotoxicity toward cancer cells (8, 9). We recently reported that HsA decreases proinflammatory responses in macrophages by inhibiting the nuclear factor-kappa B (NF- κ B) signaling pathway and inducing the expression of nuclear factor E2-related factor 2 (Nrf2)-dependent antioxidant genes (10). The end result is attenuation of the fulminant hepatitis induced by lipopolysaccharide/galactosamine in mice (10). In addition, HsA promotes apoptosis of activated hepatic stellate cells by inhibiting the NF- κ B and Akt signaling pathways, which in turn helps alleviate carbon tetrachloride-induced fibrosis in mice (11). Although HsA may have hepatoprotective potential, its effect on nonalcoholic steatosis is unknown. Therefore, we evaluated the effects of HsA on T0901317-mediated hepatic lipogenesis *in vitro* and *in vivo*.

RESULTS

HsA inhibits T0901317-induced LXR α expression

We first assessed the viability after HsA-pretreated HepG2 and Huh7 cells (1–10 μ M, 1 h) were exposed to 10 μ M T0901317 for 24 h. HsA at up to 10 μ M did not affect cell viability compared to T0901317-treated cells (Fig. 1A). Next, reporter gene assays using an LXRE-driven

luciferase plasmid were conducted to investigate whether HsA inhibits the transactivation of LXR α . Treatment with T0901317 (10 μ M, 18 h) increased the LXRE luciferase activity in HepG2 cells. However, pretreatment with 5 and 10 μ M HsA decreased LXRE luciferase activity compared to that in T0901317-treated cells. HsA (10 μ M) alone did not affect luciferase activity (Fig. 1B). Additional LXRE reporter gene assays were conducted to confirm the inhibitory effects of HsA on LXR α transactivation. HsA (5 and 10 μ M) inhibited the LXRE-driven luciferase activity, which was increased by GW3965 (3 μ M; another LXR ligand) (Fig. 1C, left) and by ectopic expression of LXR α /RXR α (Fig. 1C, right).

Because LXR α ligands autonomously increase LXR α expression (12), next, we evaluated the expression level of LXR α . The *LXR α* mRNA level was increased in HepG2 and Huh7 cells exposed to T0901317 (10 μ M, 12 h). However, the T0901317-mediated induction of *LXR α* was inhibited by pretreatment with 5 and 10 μ M HsA (Fig. 1D).

HsA attenuates T0901317-induced lipogenesis *in vitro*

Next, we explored the effects of HsA on the expression of SREBP-1c. HsA (5 and 10 μ M) inhibited the T0901317-induced mRNA level of *SREBP-1c* in Huh7 cells (Fig. 2A). It also decreased the SREBP-1c protein level in HepG2 and Huh7 cells (Fig. 2B). The results from an SREBP response element (SREBP_{RE}) reporter gene assay revealed that T0901317-mediated transactivation of SREBP-1c was inhibited by 5 and 10 μ M HsA (Fig. 3A). Moreover, induction of the expression of LXR α /SREBP-1c-dependent lipogenic genes (e.g., FASN, SCD-1, and ACC) was suppressed by HsA pretreatment (Fig. 3B and 3C). Although pretreatment with 5 μ M HsA tended to decrease the T0901317-mediated elevated FASN and ACC protein

levels in HepG2 cells, there were no significant differences compared to T0901317-treated cells (Fig. 3C).

To further explore HsA-mediated inhibition of lipogenic genes, Huh7 cells were stained with oil red O and their triglyceride levels were quantified after treatment with T0901317 in the presence or absence of HsA. **HsA (10 μ M, 24 h) reduced** the T0901317-mediated accumulation of oil red O-stained lipid droplets (Fig. 3D) and triglycerides (Fig. 3E, right). An HsA-mediated reduction in triglycerides was also noted in HepG2 cells (Fig. 3E, left).

HsA decreases T0901317-induced lipogenesis in mice

Next, 5 or 10 mg/kg HsA was intraperitoneally injected into C57BL/6 mice once daily for 3 days, and 25 mg/kg T0901317 was orally administered at 1 h after HsA injection (Fig. 4A). Biochemical analyses of serum indicated that **there were no differences in alanine aminotransferase (ALT) and aspartate aminotransferase (AST) levels between vehicle- and T0901317-treated mice. In addition, HsA injection in combination with T0901317 did not affect the ALT and AST levels compared to T0901317-treated mice.** The AST level was slightly, but significantly, increased in mice injected with only HsA (10 mg/kg) (Fig. 4B). Moreover, the T0901317-mediated increase in the relative liver weight **was attenuated** by 10 mg/kg HsA (Fig. 4C). Hematoxylin and eosin- and oil red O-staining indicated that **T0901317 increased** the diameter of hepatocytes and the accumulation of lipid droplets, but **HsA (5 and 10 mg/kg) inhibited** these T0901317-mediated changes (Fig. 4D). Furthermore, impaired serum and hepatic levels of triglyceride by **T0901317 were restored** by HsA (10 mg/kg) injection (Fig. 4E).

To confirm the inhibitory effects of HsA on hepatic lipogenesis, the mRNA levels of genes associated with LXR α -dependent lipogenesis were determined (Fig. 4F). **T0901317 decreased**

the *LXRα* mRNA level, and HsA (10 mg/kg) partly, but significantly, prevented the reduction in the *LXRα* mRNA level in hepatic tissues. As expected, T0901317 increased the mRNA levels of *SREBP-1c*, *FASN*, and *SCD-1* in hepatic tissues. Compared to T0901317, 5 and 10 mg/kg HsA decreased the *SREBP-1c* mRNA. However, the *FASN* and *SCD-1* mRNA levels were only reduced in mice treated with 10 mg/kg HsA. The mRNA level of *ACC* did not differ among the treatment groups.

DISCUSSION

Because T0901317 activates *LXRα* more potently than *LXRβ* and other nuclear receptors (e.g., farnesoid X receptor and pregnane X receptor) (13, 14), it has been used as an *LXR* ligand in studies of the pathophysiological role of *LXRα*. In this work, we found that HsA alleviated hepatic lipogenesis by inhibiting T0901317-mediated *LXRα* activation and the expression of its target genes *in vitro* and *in vivo*. In addition, it inhibited the elevated LXRE luciferase activity induced by GW3965 treatment or *LXRα*/*RXRα* expression. Therefore, the inhibition of hepatic lipogenesis by HsA is not mediated by its regulation of other nuclear receptors.

HsA significantly restored the T0901317-mediated changes in the mRNA level of *LXRα*. In agreement with a previous report (12), T0901317 increased *LXRα* expression in hepatocyte-derived cells (Fig. 1D). However, three T0901317 administrations significantly reduced the hepatic *LXRα* mRNA level in mice (Fig. 4F). Although the role of T0901317 in insulin level is unclear, insulin increases hepatic lipogenesis by inducing *LXRα* expression (15). T0901317 reportedly impairs glucose-mediated insulin secretion in pancreatic islets (16) and decreases the insulin level in *db/db* mice fed a high-fat diet (17). In addition, T0901317 promotes

palmitate- and glucose-mediated apoptosis of pancreatic β cells (18, 19). Therefore, the reduction in hepatic *LXR α* mRNA level caused by T0901317 **may be** a result of decreased secretion of insulin in the pancreas. **The discrepant *in vitro* and *in vivo* results showing *LXR α* mRNA expression induced by T0901317, and the effects of HsA and T0901317 on other extrahepatic tissues need to be further resolved.**

LXR α -mediated gene expression can be regulated by post-translational modification (e.g., phosphorylation) of *LXR α* , and adenosine 5'-monophosphate-activated protein kinase (AMPK) downregulates the activity of *LXR α* and SREBP-1c via direct phosphorylation (3, 21). In addition, AMPK inhibits fatty acid synthesis by inhibitory phosphorylation of ACC and facilitates autophagy to reduce lipid accumulation in hepatocytes (20). Moreover, leucodin, another guaianolide-subtype STL, inhibits ethanol-mediated lipid accumulation by activating AMPK (22). To explore the possibility that HsA inhibits hepatic lipogenesis via AMPK and/or autophagy activation, Huh7 cells were preincubated with chemical inhibitors of AMPK or autophagy and subsequently treated with HsA and T0901317. Our additional results showed that the HsA-mediated inhibition of SREBP-1c was sustained after pretreatment with the chemical inhibitors (Supplementary Fig. 1A and 1B). This suggests that HsA alleviates T0901317-mediated activation of the *LXR α* -SREBP-1c signaling pathway in a manner independent of AMPK and autophagy activation.

We previously reported that inhibition of Akt as well as activation of p38 mitogen-activated protein kinase (MAPK) and c-Jun N-terminal kinase contribute to the HsA-mediated prevention of hepatic inflammation and fibrosis (10, 11). Sustained activation of the phosphoinositide 3-kinase/Akt signaling pathway by phosphatase and tensin homolog deficiency provokes nonalcoholic steatohepatitis in mice (23). In addition, p70 ribosomal S6 kinase, a downstream factor of Akt, enhances *LXR α* activity (3), suggesting that HsA inhibits

LXR α -mediated hepatic lipogenesis by modulating the Akt-dependent signaling pathway. However, at least under our experimental conditions, pretreatment with Akt inhibitors did not decrease T0901317-induced SREBP-1c expression. Moreover, MAPK inhibitors did not significantly alter the HsA-mediated SREBP-1c reduction (Supplementary Fig. 1C and 1D). Although HsA inhibited T0901317-mediated hepatic lipogenesis, further study is needed to identify the major signaling molecules associated with HsA-mediated inactivation of LXR α and SREBP-1c.

Lipid accumulation is necessary but not sufficient for progression from simple steatosis to the more serious NAFLD; other insults are also required. For instance, overproduction of fatty acids accelerates hepatocyte death by promoting oxidative stress and mitochondrial dysfunction, which amplifies the inflammatory response intended to clear damaged hepatocytes (1). Hence, oxidative stress and inflammation are implicated in the development of advanced NAFLD. Deletion of Nrf2, a master transcription factor for the protection of cells from oxidative stress, exacerbates hepatic steatosis and inflammation in mice fed a methionine- and choline-deficient diet (24). Moreover, several natural products inhibit NAFLD by activating Nrf2-dependent antioxidant genes (25). Interestingly, our results showed that HsA significantly increased nuclear expression of Nrf2, transactivated the luciferase activity of an antioxidant response element-harboring reporter gene, and increased the mRNA levels of Nrf2-dependent antioxidant genes (e.g., heme oxygenase 1, glutamate-cysteine ligase catalytic subunit, and sestrin 2) in hepatocyte-derived cells (Supplementary Fig. 2A–2C). In addition, T0901317-mediated hepatic lipid peroxidation and protein nitrosylation were significantly reduced in HsA-treated mice (Supplementary Fig. 2D). Moreover, HsA suppresses proinflammatory responses in Toll-like ligand-stimulated macrophages and induces apoptosis of activated hepatic stellate cells by downregulating the NF- κ B signaling pathway (10, 11).

Therefore, the antioxidant and anti-inflammatory effects of HsA may together prevent progression to advanced NAFLD. In conclusion, HsA ameliorated lipogenesis by inhibiting the LXR α -SREBP-1c signaling pathway in T0901317-stimulated hepatocyte-derived cells as well as in the mouse liver. Therefore, HsA has prophylactic and therapeutic potential for fatty liver.

MATERIALS AND METHODS

Reagents

HsA was isolated from *H. lyrata* as previously reported (11). T0901317 was supplied by Cayman Chemical (Ann Arbor, MI, USA). Anti-SREBP1 was purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Anti-FASN, anti-ACC, and horseradish peroxidase-conjugated secondary antibodies were obtained from Cell Signaling Technology (Beverly, MA, USA). Thiazolyl blue tetrazolium bromide (MTT), GW3965, β -actin antibody, oil red O, hematoxylin, eosin, and other reagents were supplied by Sigma-Aldrich (St. Louis, MO, USA).

Cell culture and treatment

HepG2 and Huh7 cells (human hepatocyte-derived cells) were supplied by American Type Culture Collection (Manassas, VA, USA) and Korean Cell Line Bank (Seoul, Korea), respectively. The cultured cells were pretreated with 1–10 μ M HsA for 1 h and subsequently exposed to either T0901317 (10 μ M) or GW3965 (3 μ M) for the indicated time periods.

Cell viability assay

The treated cells were incubated with MTT (0.5 mg/ml) for 2 h, and A_{570} was measured using

a Synergy HTX Multi-Mode Plate Reader (BioTek, Winooski, VT, USA). Relative cell viability was calculated as a percentage of untreated controls.

Reporter gene assay

Reporter plasmids (LXRE and SREBP_{RE}) and expression plasmids (LXR α and RXR α) were provided, as previously described (3). pRL-SV40, a reporter plasmid expressing *Renilla* luciferase, was obtained from Promega (Madison, WI, USA). LXRE or SREBP_{RE} reporter plasmid (300 ng) was transiently transfected with pRL-SV40 (40 ng) into HepG2 cells, as previously reported (11). For some experiments, LXR α and RXR α (300 ng each) were co-transfected with LXRE reporter plasmid, and equal amounts of pcDNATM 3.2/V5-DEST plasmid (Invitrogen, Carlsbad, CA, USA) were used for mock transfection. The relative luciferase activities were calculated as the firefly luciferase activity divided by *Renilla* luciferase activity.

Real-time polymerase chain reaction (PCR)

Isolation of RNA, reverse transcription, real-time PCR, and relative quantification of specific mRNAs were carried out as previously described (11). Gene-specific primers are listed in Supplementary Table 1.

Western blot analysis

Preparation of whole-cell protein lysates, Western blot analysis, and chemiluminescence detection were conducted, as previously described (11). Band intensities of proteins of interest were analyzed using ImageJ software (National Institutes of Health, Bethesda, MD, USA) and normalized to those of β -actin.

Oil red O staining of Huh7 cells

Treated cells were fixed in 100% ethanol, stained with oil red O (2 mg/ml in 60% isopropanol) for 10 min, and visualized under a light microscope (Eclipse Ti-U; Nikon, Kanagawa, Japan).

Quantification of triglycerides

Triglyceride levels in cultured cells or hepatic tissues were quantified using a Triglyceride Quantification Assay Kit (Abcam, Cambridge, UK) according to the manufacturer's instructions and normalized to the protein concentration.

Animal experiments and measurement of relative liver weight

Animal experiments were approved by the Institutional Animal Care and Use Committee of Daegu Haany University (Approval No. DHU2018-016). Forty C57BL/6 mice (6-weeks-old males; Saeron Bio, Euiwang, Republic of Korea) were maintained under standard conditions as previously described (11). Mice (N = 8 per group) were intraperitoneally injected with 5 or 10 mg/kg HsA dissolved in corn oil once daily for three consecutive days. To induce steatosis, T0901317 (25 mg/kg) in 1% carboxymethylcellulose and 1% Tween 20 was orally administered three times at 1 h after HsA injection. An equal volume of corn oil and carboxymethyl cellulose/Tween 20 were administered to vehicle group. **Dosages of HsA and T0901317 were chosen based on previous studies (10,11) and a preliminary study (data not shown), respectively.** All mice were euthanized 24 h after the last T0901317 administration, and the liver and blood were collected. The relative liver weights were calculated as the percentage of liver weight to body weight.

Serum biochemistry

Serum levels of ALT, AST, and triglyceride were measured using a blood chemistry analyzer (Dri-Chem NX500i; Fuji Medical System, Tokyo, Japan).

Tissue staining and histopathology

Hepatic sections were stained with hematoxylin and eosin or oil red O, as previously described (26). The hepatocyte diameter ($\mu\text{m}/\text{hepatocytes}$) was calculated as the mean diameter of 10 hepatocytes in hematoxylin and eosin-stained tissues. The regions occupied by lipid droplets ($\%/\text{mm}^2$ of hepatic parenchyma) were calculated as the proportion of oil red O stained area in a restricted-view field of hepatic parenchyma.

Statistical analysis

All numerical results are expressed as the mean \pm standard deviation (SD) of at least three independent experiments (*in vitro* assay) or eight mice (*in vivo* assay). The One-way analysis of variance was conducted to assess the significance of differences among experimental groups, followed by Tukey's honestly significant difference or Dunnett's T3 test. *P* values less than 0.05 were considered statistically significant.

ACKNOWLEDGMENTS

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CONFLICT OF INTEREST

The authors declare that they have no conflict of interests.

FIGURE LEGENDS

Fig. 1. Hemistepsin A (HsA) inhibits T0901317-induced expression of LXR α . Relative viabilities of cells treated with HsA in the presence of T0901317 were determined using an MTT assay (A). HepG2 cells were transfected with the LXRE reporter plasmid with or without the LXR α /RXR α expression plasmids. Luciferase activities were determined after transfected cells were treated with HsA, T0901317 (B), and GW3965 (C, left). pcDNATM 3.2/V5-DEST was used for mock transfection (C, right). Relative mRNA levels of LXR α were determined by real-time PCR (D). ** $P < 0.01$ versus untreated controls or mock transfection; # $P < 0.05$, ### $P < 0.01$ versus T0901317, GW3965, or LXR α /RXR α . NS, not significant.

Fig. 2. HsA decreases T0901317-induced SREBP-1c expression. Huh7 and HepG2 cells were treated with 5–10 μ M HsA (1 h) and 10 μ M T0901317 (12 h). Relative mRNA (A) and protein (B) levels of SREBP-1c determined by real-time PCR and Western blotting, respectively. Lines in the right panel of (B) indicate cropped images of the same membrane with the same exposure. ** $P < 0.01$ versus untreated controls; # $P < 0.05$, ### $P < 0.01$ versus T0901317.

Fig. 3. HsA attenuates T0901317-induced lipogenesis. HsA-pretreated cells (5–10 μ M, 1 h) were exposed to T0901317 for 18 (A), 12 (B), or 24 h (C–E). An SREBP_{1c} reporter gene assay was conducted using HepG2 cells (A). The relative mRNA and protein levels of lipogenic

genes (B and C) were determined. The cells were stained with oil red O (D), and levels of triglycerides (E) were measured. * $P < 0.05$, ** $P < 0.01$ versus untreated controls; # $P < 0.05$, ## $P < 0.01$ versus T0901317.

Fig. 4. HsA alleviates T0901317-induced lipogenesis in mice. Animal experimental procedure (A). ALT (B, left), AST (B, right), and triglyceride (E, left) levels in serum and relative liver weights (C) were measured after euthanizing the mice. The mean diameters of hepatocytes and the lipid droplet-occupied regions (D, right) were calculated from hepatic tissues stained with hematoxylin and eosin and oil red O (D, left). Scale bars indicate 60 μm . The levels of triglycerides (E, right) and the mRNA levels of lipogenic genes (F) were measured in hepatic tissues. * $P < 0.05$, ** $P < 0.01$ versus vehicle group; # $P < 0.05$, ## $P < 0.01$ versus T0901317. CV, central vein; NS, not significant; PT, portal triad.

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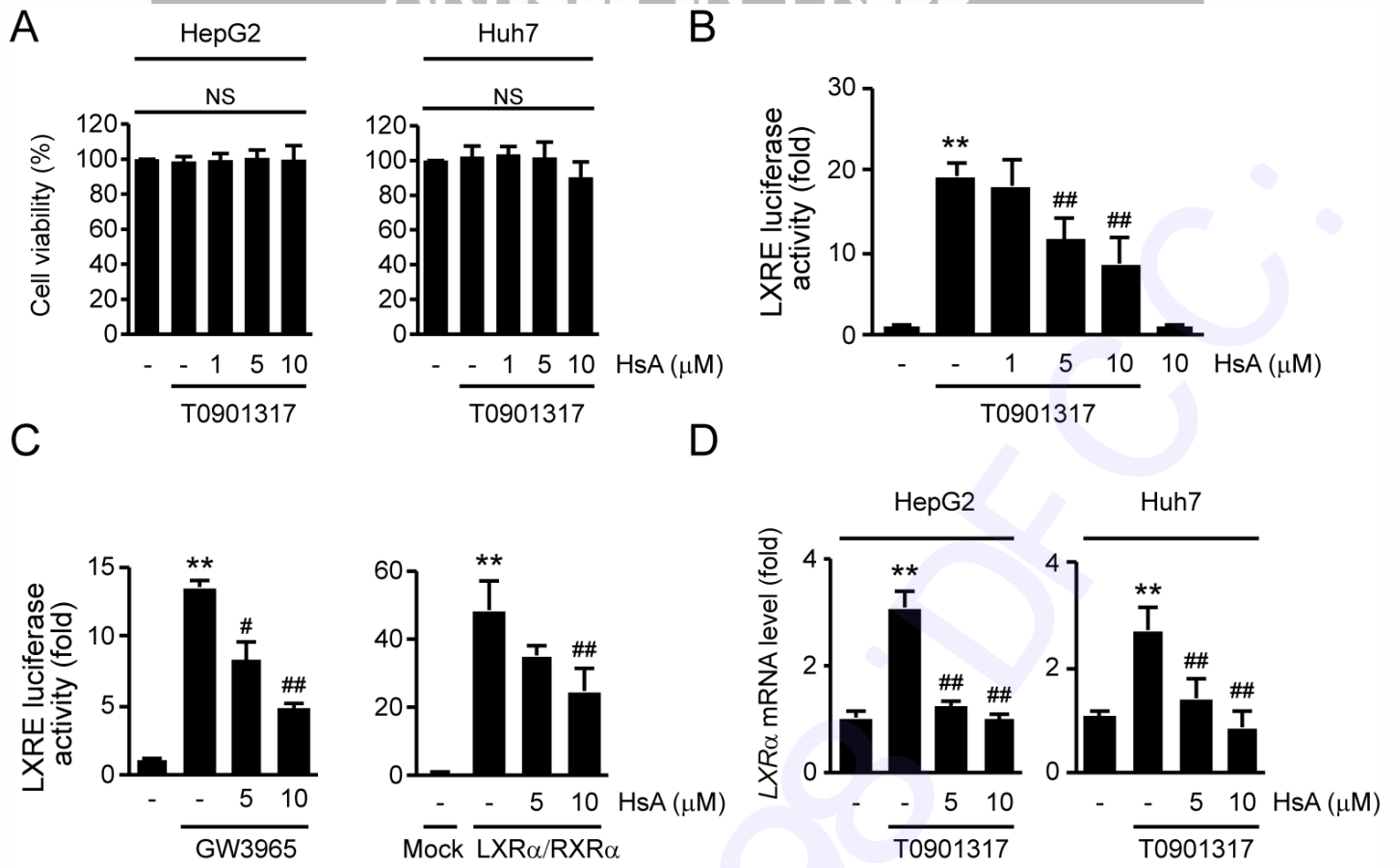


Figure 1

Fig. 1. figure 1

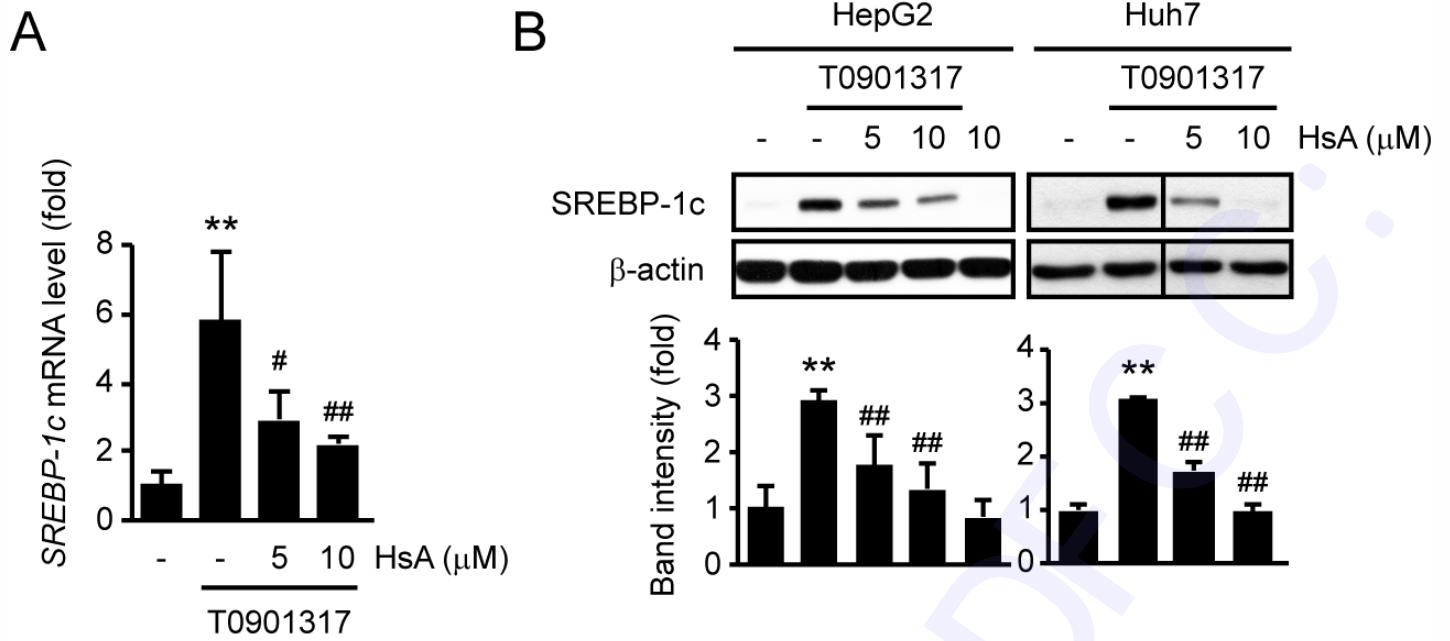


Figure 2

Fig. 2. figure 2

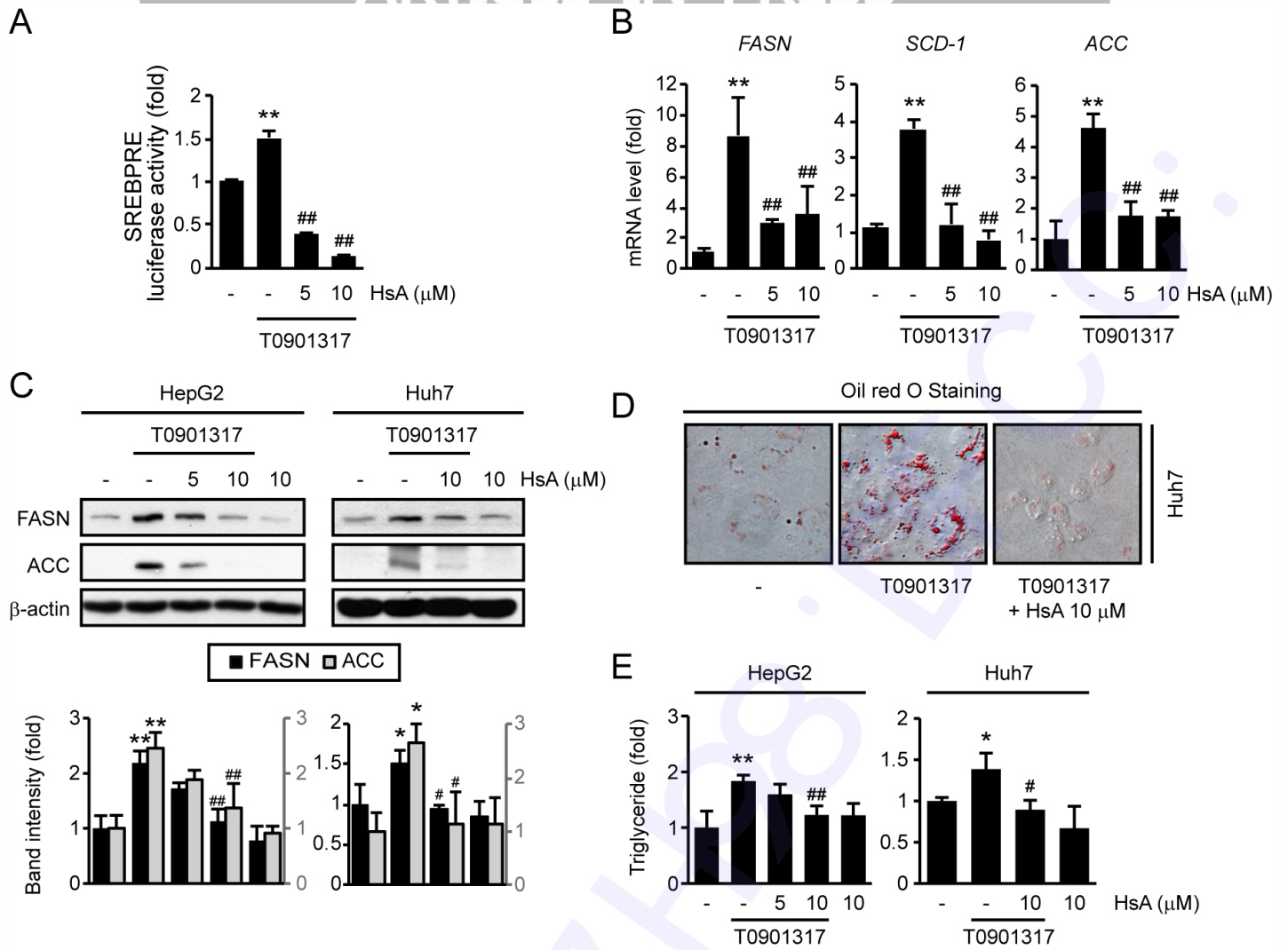


Figure 3

Fig. 3. figure 3

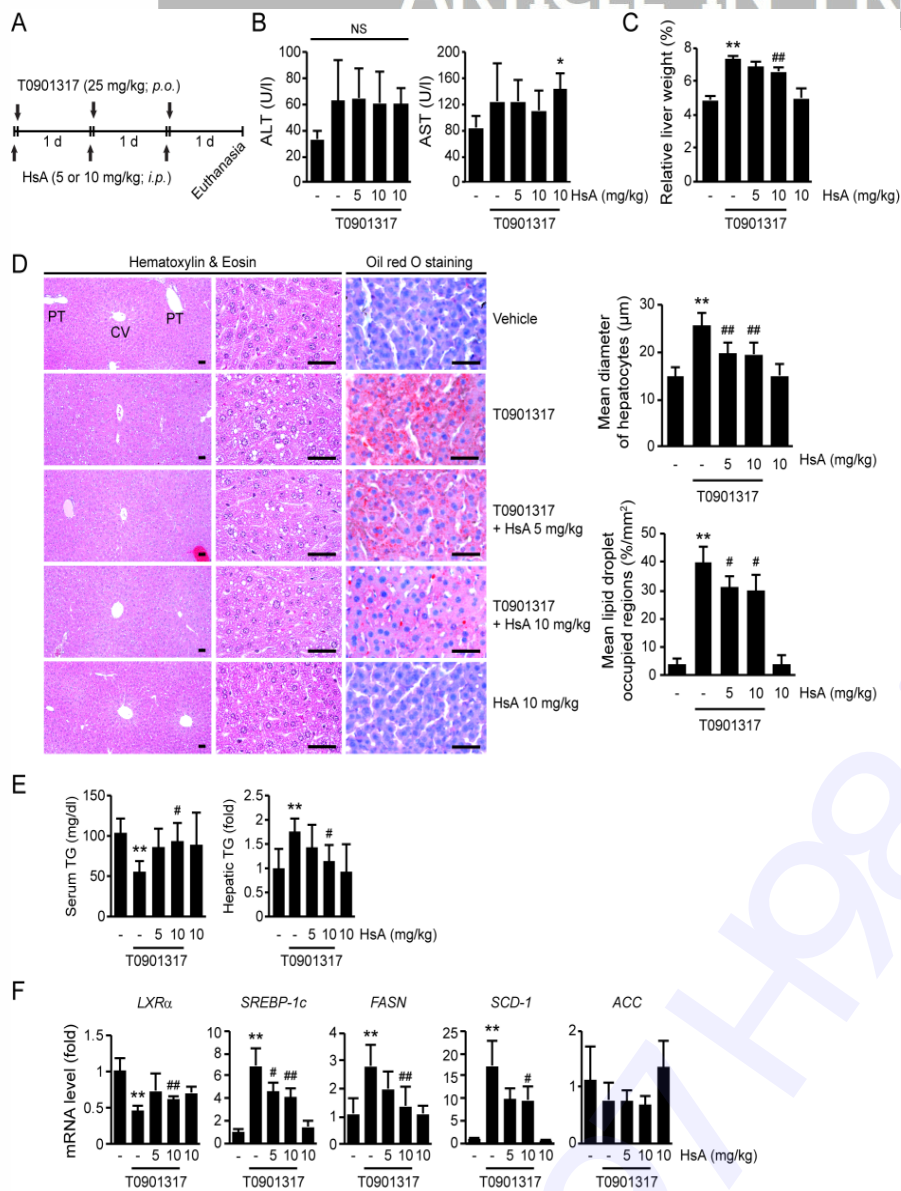


Figure 4

Fig. 4. figure 4

Title: Hemistepsin A inhibits T0901317-induced lipogenesis in the liver

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Running Title: Inhibition of hepatic lipogenesis by hemistepsin A

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J.K. Kim and I. J. Cho are contributed equally to this work.

Supplementary materials

Compound C was supplied by Calbiochem (San Diego, CA, USA). Bafilomycin A1, LY294002, and anti-Nrf2 antibody were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). PD98059, SB203580, SP600125, and anti-nitrotyrosine antibody were obtained from Millipore (Temecula, CA, USA). Antibodies against 4-hydroxynonenal and histone H3 were provided by Abcam (Cambridge, UK) and Cell Signaling Technology (Danvers, MA, USA), respectively. Chloroquine, 3-methyladenine, and wortmannin were supplied by Sigma-Aldrich (St. Louis, MO, USA).

Isolation of nuclear proteins

Nuclear proteins were extracted from HsA (5 or 10 μ M, 6 h)-treated HepG2 cells using NE-PERTM Nuclear and Cytoplasmic Extraction Reagents (ThermoFisher Scientific; Somerset, NJ, USA) as described previously (1).

Reporter gene assay

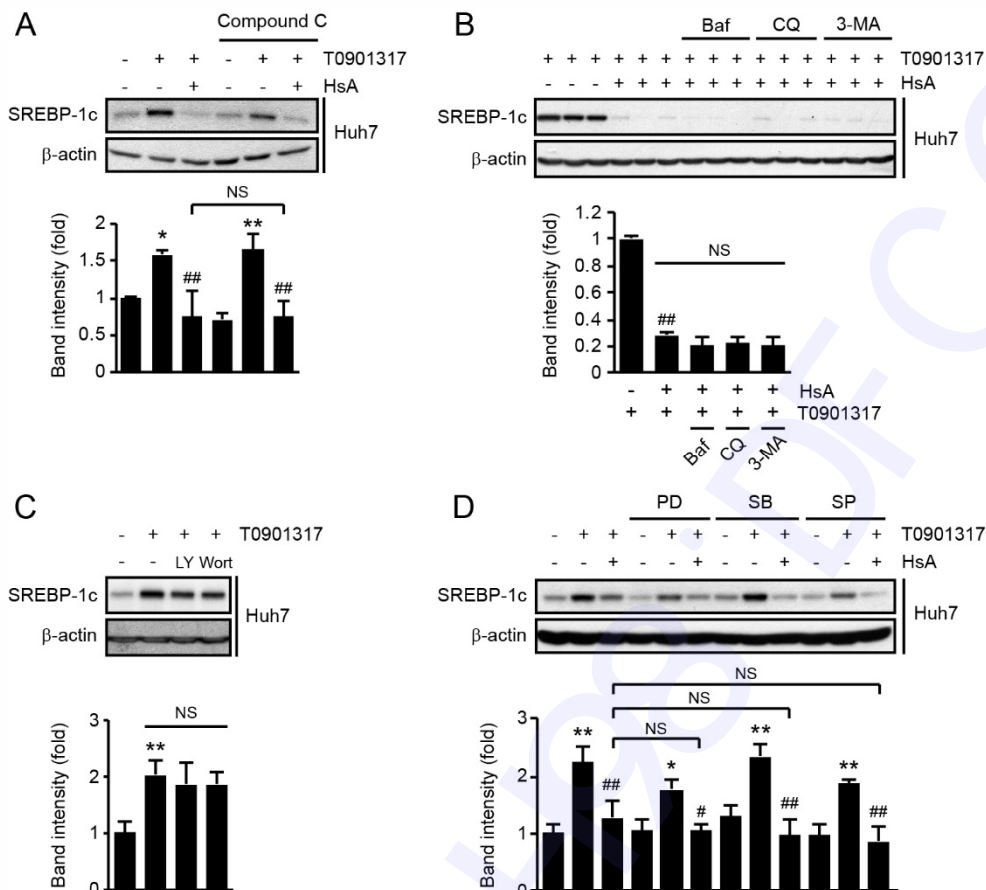
Recombinant HepG2 cells which stably express antioxidant response element (ARE) harboring reporter gene were established, as previously reported (2). After the cells were treated with 5 and 10 μ M HsA for 18 h, luminescence intensities were normalized by protein concentration.

Immunohistochemistry

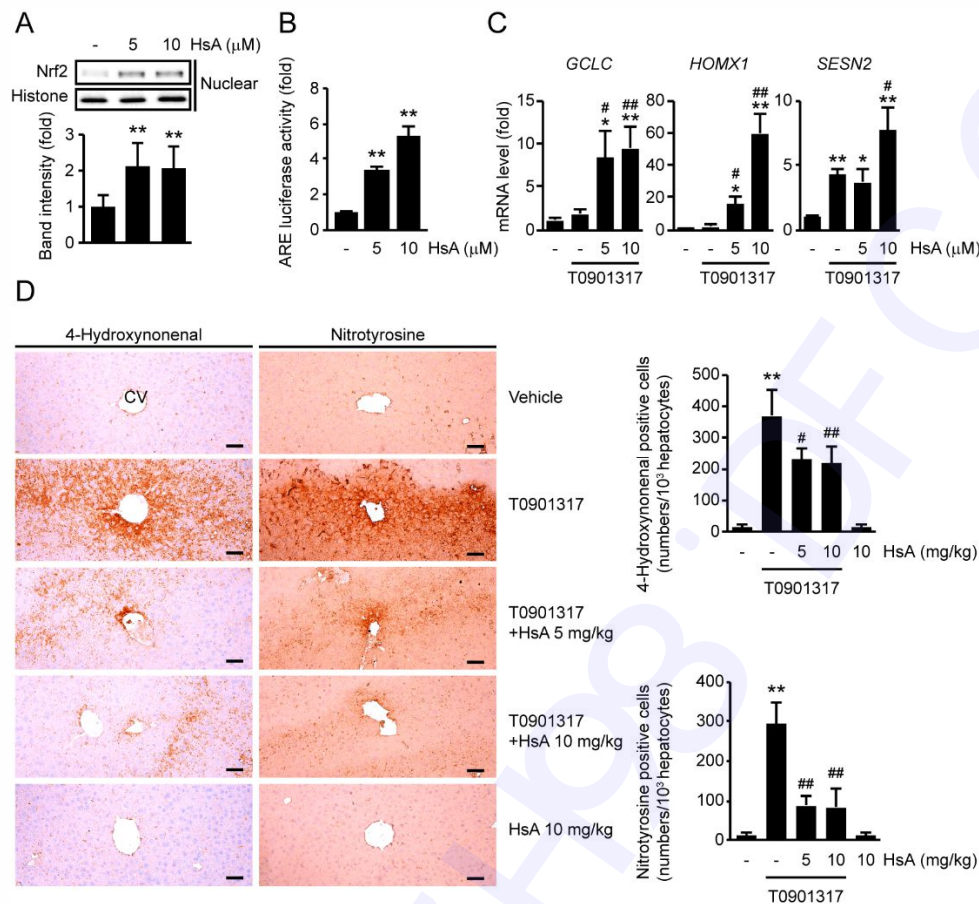
Hepatic tissues were immunostained using anti-nitrotyrosine or anti-4-hydroxynonenal as described previously (3), and cells exhibiting > 20% immunoreactivity in hepatic parenchyma were regarded as positive.

Supplementary Table 1. Primer sequences used in the present study

Species	Gene symbol	Forward primer	Backward primer	Amplicon size (bp)
Human	<i>LXRα</i>	5'-GATCGAGGTGATGCTTCTGGAG-3'	5'-CCCTGCTTTGGCAAAGTCTTC-3'	109
	<i>SREBP-1c</i>	5'-CGACATCGAAGACATGCTTCAG-3'	5'-GGAAGGCTTCAAGAGAGGAGC-3'	179
	<i>FASN</i>	5'-GACATCGTCCATTCGTTTGTG-3'	5'-CGGATCACCTTCTTGAGCTCC-3'	467
	<i>SCD-1</i>	5'-CCTCTACTTGGAAGACGACATTCGC-3'	5'-GCAGCCGAGCTTTGTAAGAGCGGT-3'	281
	<i>ACC</i>	5'-GCTGCTCGGATCACTAGTGAA-3'	5'-TTCTGCTATCAGTCTGTCCAG-3'	270
	<i>HMOX1</i>	5'-CAGGAGCTGCTGACCCATGA-3'	5'-AGCAACTGTCGCCACCAGAA-3'	195
	<i>SESN2</i>	5'-CAAGCTCGGAATTAATGTGCC-3'	5'-CTCACACCATTAAGCATGGAG-3'	323
	<i>GCLC</i>	5'-GAAGTGGATGTGGACACCAG-3'	5'-TTGTAGTCAGGATGGTTTGCGA-3'	128
	<i>GAPDH</i>	5'-GAAGGTGAAGGTCGGAGTC-3'	5'-GAAGATGGTGATGGGATTTC-3'	226
Mouse	<i>LXRα</i>	5'-TGCCATCAGCATCTTCTCTG-3'	5'-GGCTCACCAGCTTCATTAGC-3'	161
	<i>SREBP-1c</i>	5'-ACCGTCACTTCCAGCTAGAC-3'	5'-CCACTAAGGTGCCTACAGAGC-3'	181
	<i>FASN</i>	5'-AGCGGCCATTTCATTGCCC-3'	5'-CCATGCCCAGAGGGTGGTTG-3'	302
	<i>SCD-1</i>	5'-CCGGAGACCCTTAGATCGA-3'	5'-TAGCCTGTAAAAGATTTCTGCA-3'	88
	<i>ACC</i>	5'-GTCAGCGGATGGGCGGAATG-3'	5'-CGCCGGATGCCATGCTCAAC-3'	270



Supplementary Fig. 1. Effect of chemical inhibition of AMPK, autophagy, Akt, or MAPKs signaling pathway on the HsA-mediated SREBP-1c reduction. Huh7 cells were pretreated with compound C (10 μ M; A), bafilomycin A1 (2 nM; B), chloroquine (10 μ M; B), 3-methyladenine (2.5 mM; B), LY294002 (10 μ M; C), **wortmannin** (1 μ M; C), PD98059 (30 μ M; D), SB203580 (30 μ M; D), or SP600125 (30 μ M; D) for 1 h and then exposed to T0901317 (10 μ M) with or without HsA (10 μ M) for 12 h. Equal protein loading was verified by β -actin Western blotting. Results are presented as mean \pm SD of at least three independent experiments. * P < 0.05, ** P < 0.01 *versus* untreated controls; # P < 0.05, ## P < 0.01 *versus* T0901317. Baf, bafilomycin A1; CQ, chloroquine; LY, LY294002; 3-MA, 3-methyladenine; NS, not significant; PD, PD98059; SB, SB203580; SP, SP600125; Wort, **wortmannin**.



Supplementary Fig. 2. HsA activates Nrf2-dependent antioxidant genes and inhibits T0901317-mediated oxidative stress. Nrf2 expression in the nucleus were determined after HepG2 cells were treated with 5 or 10 μM HsA for 6 h. Equal protein loading was verified by histone H3 Western blotting (A). Recombinant HepG2 cells which had been stably transfected with ARE-driven reporter gene were treated with 5 and 10 μM HsA for 18 h. Luciferase activities were normalized by protein concentration (B). After HsA (5 and 10 μM)-pretreated Huh7 cells were incubated with T0901317 (10 μM) for 12 h, mRNA levels of *HMOX1*, *GCLC*, and *SESN2* were analyzed by real-time PCR (C). Hepatic tissues were immunostained using either 4-hydroxynonenal or nitrotyrosine antibody (D, left), and immunoreactive cells were counted using an automated image analyzer (D, right). Scale bars indicate 60 μm (D, left). Results are presented as mean ± SD of three independent experiments (A–C) or eight mice (D). **P* < 0.05, ***P* < 0.01 versus untreated controls or vehicle group; #*P* < 0.05, ##*P* < 0.01 versus T0901317. CV, central vein; GCLC, glutamate-cysteine ligase catalytic subunit; HMOX1, heme oxygenase 1; SESN2, sestrin 2.

SUPPLEMENTARY REFERENCES

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