

Quercetin promotes long noncoding RNA IncSHGL expression via estrogen receptor α to suppress type 2 diabetes mellitus in mice

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Research

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Abstract

Background Type 2 diabetes mellitus (T2DM) is the most common type of metabolic disorder involving multiple organ systems. Grape has been reported to improve the symptoms of T2DM, the precise mechanism of its action is unclear. Our study was aimed to determine the effect and mechanism of grape pomace extract in T2DM mice induced by high fat diet (HFD). **Materials and methods** Ultra-performance liquid chromatography and quadrupole time-of-flight mass spectrometry were used to identify the main active compounds in grape pomace extract to improve T2DM. C57BLK/6J mice induced by HFD supplemented with or without quercetin were used to show the effects of quercetin improving T2DM. By online database research, bioinformatics analysis and molecular biology experiments, Estrogen receptor alpha (ER α)-lncSHGL (lncRNA suppressor of hepatic gluconeogenesis and lipogenesis) pathway was identified as the target for quercetin. **Results** Quercetin was identified as one of the most active compounds in grape pomace extract to improve T2DM. Quercetin could inhibit HFD-induced T2DM in mice by activating ER α . lncSHGL was identified as the downstream of ER α and inhibited HFD-induced T2DM. **Conclusions** Quercetin could be beneficial for T2DM by promoting lncSHGL transcription and activating the lncSHGL pathway, and may be used as a drug component to treat T2DM.

Introduction

Type 2 diabetes mellitus, the fifth leading cause of death in the world in 2000, is a global epidemic [1]. The prevalence of type 2 diabetes mellitus worldwide in adults older than 18 years has increased from 4.7% in 1980 to 8.5% in 2014. In 2016, type 2 diabetes mellitus was estimated to cause 1.6 million deaths [2], and it is expected to affect 366 million individuals by 2030 [3, 4]. According to the reports from the Chinese Center for Disease Control and Prevention, the prevalence of type 2 diabetes mellitus in Chinese individuals older than 18 years increased from 9.7% in 2010 [5] to 10.4% in 2013 [6].

Presently, several drugs can suppress the rise of blood glucose level by maintaining it within a normal range. Because diet, activity, stress and metabolism kinetics can affect the blood glucose level, which eventually perturbing the insulin sensitivity, diabetic treatment is challenging [7]. In addition, the complexity of the treatment and the side effects of the drug itself are often unacceptable, as they negatively impact the quality of life [8]. An increasing number of individuals are turning to natural substances and foods to treat type 2 diabetes mellitus. Grapes are rich in vitamins, minerals and polyphenols, including flavonoids, proanthocyanidins, and procyanidins [9]. Many studies have shown that GPE can reduce postprandial hypertriglyceridemia and hypercholesterolemia as well as prevent insulin resistance (IR) [10, 11], due to its inhibitory effects on the intestinal α -glucosidase and pancreatic α -amylase [12]. Another study showed that GSE significantly improved obesity and hyperglycemia and reduced the serum glucosamine level in obese type 2 diabetes mellitus patients [13]. In acute-dose studies, grape polyphenols were reported to reduce the serum glucose concentration in animals and humans [14, 15]. Although studies have shown that GPE can effectively reduce hyperglycemia, the precise mechanism of action is unclear.

With the advancement of RNA-seq technology, nearly 3 billion non-coding RNAs (nc-RNAs) accounting for 97–98% of the human genome with unclear gene functions have been discovered, [16, 17]. Among these, 80–90% of ncRNA are longer than 200 nucleotides and not traditional protein-encoding RNAs, these ncRNAs are defined as long non-coding RNAs (lncRNAs) [18–20]. Increasing evidence has shown that lncRNAs have different biological functions, and several lncRNAs have functions in glucose and lipid metabolism [8, 21].

Here, we report that quercetin, one of the most active compounds in grape pomace extract (GPE), can improve the symptoms of HFD-induced type 2 diabetes mellitus via the regulation of lncRNA suppressor of hepatic gluconeogenesis and lipogenesis (lncSHGL). Our results show that lncSHGL might be a new therapeutic target for type 2 diabetes mellitus.

Materials And Methods

Mouse food preparation

Rodent chow and western style high fat diet (HFD) were purchased from Huafukang co. (Beijing, China). For quercetin containing HFD diet, 0.1g/kg quercetin (Sigma, USA) of HFD, was tailor-made by the same company. All mouse food was sterilized by irradiation to minimize bacterial contamination. Macronutrient and selected micronutrient content in the mouse HFD is summarized in Table 1.

Animals

Six-week male C57BLK/6J mice (Charles River, China) were housed in clean cages at 12 h light dark cycle at 20°C to 22°C, with 4 mice/cage. After adaptive feeding for 1 week, all mice were randomized in each group. All animal experimental protocols are in line with the Animal Management Rules of the Ministry of Health of the People's Republic of China and the China Agricultural University Guide for the Care and Use of the Laboratory Animals. The ethical committee No. is CAU20180217-4.

Type 2 diabetes mellitus mouse model

Type 2 diabetes mellitus mouse model was reported by Houssay and colleagues previously [22, 23]. Briefly, the islet B cells were kept in a high-load state for a long time through a HFD, and finally induced a mouse model characterized by obesity, impaired glucose tolerance, IR, and abnormal lipid metabolism [22]. The model has been reported to mimic the early human and progression of type 2 diabetes mellitus [24].

Purification and identification of active compounds by ultra-performance liquid chromatography (UPLC) combined with a quadrupole time-of-flight mass spectrometry (Q-TOF-MS)

UPLC system and chromatographic conditions and identification of chemical constituents by UPLC/Q-TOF-MSE analysis the Waters ACQUITY UPLC system (Acquity UltraPerformance Liquid Chromatography™, USA). Grape pomace was analyzed for its chemical profiles using a Waters Xevo G2

Q-TOF mass spectrometer (Milford, USA). To collect data in TOF MS experiments, MSE technology was used in which two separate scan functions were programmed for the MS acquisition method. Data were acquired and analyzed with Waters MassLynx v4.1 software. The sample analysis was performed in triplicate to test the repeatability.

Experimental group assignment

In this experiment, since body weight is a critical experimental indicator, a block randomization method was used. Briefly, the mice were divided into 4 zones according to their weight from light to heavy, and then grouped according to the random number table.

In the pre-experiment, the effective dose for GPE in type 2 diabetes mellitus mouse model was optimized at 2.4g/Kg. As the median lethal dose of quercetin for mice was about 160 mg/kg [25], and a single intravenous injection of 100-150 mg/kg of quercetin was reported did not have any symptoms [26] 100 mg/kg quercetin in HFD was chosen in this study. Each mouse was randomly assigned to one of the following groups: 1) group-1, mice fed with regular normal chow (NC) and served as control (n=16); 2) group-2, mice fed with HFD (n=16); 3) group-3, mice fed with HFD+2.4 g/kg GPE (GHFD, n=16); 4) group-4, mice fed with HFD+100 mg/kg quercetin (QHFD, n=16). All mice were fed with the respective diets for 12 weeks with water available *ad libitum*.

Food intake, body weight and fasting glucose measurements during the experiment

During the 12-week feeding period, food intake was recorded daily, and body weight was recorded weekly. Fasting blood glucose levels were also monitored weekly using a blood glucometer and the accompanying test strips (ACCU-CHEK Meter®, Germany).

Mouse glucose tolerance test (GTT)

Briefly, the mice were fasted overnight, about 12-15 h; glucose solution (20%) was prepared, syringe, blood glucose meter, blood glucose test strip, timer and other laboratory supplies were prepared; The body weight of each mouse was recorded and fasting blood glucose was measured at 0 min. The volume of intraperitoneal injection of 20% glucose solution per mouse by 2 g glucose per kg body weight was calculated, as follows: 20% glucose (μl) = $10 \times$ body weight (g). After the glucose was injected, the blood glucose was measured at 15, 30, 60, 90, 120 min by cutting off the tail of each mouse and recorded, respectively.

Mouse insulin tolerance test (ITT)

Briefly, the mice were fasted for 5 h; the short-acting insulin stock solution (100 U/ml, Eli Lilly) was prepared, syringes, blood glucose meters, blood glucose test strips and other laboratory supplies were got ready. The body weight of each mouse was recorded and fasting blood glucose was measured at 0 min. The volume of insulin injected intraperitoneally in each mouse by injecting 0.75 U per kg body weight was

calculated. After injected with insulin, the mice were measured blood glucose at 15, 30, 60, 90, 120 min by cutting off the tail of each mouse and recorded, respectively.

Animal sacrifice and tissue collection

The mice were then injected intraperitoneally with 40 mg/kg Ketamine. Blood samples were collected and the serum was separated. Liver, heart, kidney, epididymal fat and pancreas were dissected, weighed and immediately frozen in liquid nitrogen and stored at -80°C, respectively.

Histological analysis of animal adipocytes

Freshly isolated epididymal fat and liver were fixed in 10% formalin (Sigma, USA) for 24 h followed by embedding the tissues in paraffin (Citoglas, China). The paraffin-embedded samples were sectioned and stained with hematoxylin-eosin following the procedures used by Ma et al., 2016 [27].

Mouse primary hepatocyte isolation

Hepatocytes were isolated from adult male C57BLK/6J mice, using a two-step collagenase perfusion procedure reported previously .

Cell culture

The human hepatoma cell line HepG₂, human renal epithelial cell line 293T and the mice normal liver cell line NCTC-149 were obtained from the Cell Resource Center, Peking Union Medical College (licensed from ATCC). The HepG₂ cell line and 293T cell line were maintained in Dulbecco's modified Eagle's medium – high glucose (DMEM-HG, Gibco Invitrogen, USA) supplemented with 10% fetal bovine serum (FBS, Gibco Invitrogen, USA). The NCTC-149 cell line was maintained in DMEM-HG supplemented with 10% horse serum. The mouse primary hepatocyte was maintained in Roswell Park Memorial Institute – 1640 (RPMI-1640, Gibco Invitrogen, USA) supplemented with 10% FBS. 100 units/ml penicillin (Sigma, USA), 100 µg/ml streptomycin (Sigma, USA), and 0.16% NaHCO₃ (Sigma, USA) was added to all cell culture media. All cells were incubated in a humidified atmosphere of 95% O₂ and 5% CO₂ at 37 °C.

Cell processing

When the cells were grown to about 70%, the cells were treated with quercetin, palmitic acid (P), estradiol (E₂), ICI 182780 (ICI), propylpyrazoletriol (PPT), and diarylpropionitrile (DPN), and samples were taken 24 h later for subsequent experiments. All of the reagents used above were purchased from Sigma (Sigma, USA). The working concentration of quercetin is 10µM. The working concentration of P is 500µM. The working concentrations of E₂, ICI, PPT and DPN are both 10⁻⁹ M.

Construction of plasmids

The coding sequence (CDS) of mice ER α and ER β were obtained from cDNA of mice by PCR. ER α was inserted into pCDNA3.0 through endonuclease site *EcoRI* and *BamHI*. ER β was inserted into pCDNA3.0 through endonuclease site *EcoRI* and *XhoI*. The promoter of mouse LncSHGL was obtained from the mouse genomic DNA by PCR and inserted into pGL3 vector by endonuclease site *HindIII* and *KpnI*. All of the endonucleases used above were purchased from NEB Corporation of the USA.

Transfection and luciferase activity analysis

Approximately 3×10^5 cells were seeded per well in a 6-well plate. When the cells were grown to about 70%, plasmids were transfected into 293T cell line using PEI (Polyscience, USA) transfection reagent according to the manufacturer's protocol. pGL3- LncSHGL-promoter vector and ER α (with or without) or ER β (with or without) overexpression vector were transfected into 293T cell line, and pGL3 basic vector was used as a negative control. Luciferase activity was detected by using Firefly Luciferase Detection Kit (YPH-bio, China) through Microplate reader (Bio-tec, USA) 48 h later.

RNA extraction and reverse transcription-quantitative PCR (RT-qPCR)

Total RNA was extracted from cells and tissues using TRIzol reagent (Kangwei Biomedical Technology, China). The total RNA was reversed transcribed by M-MLVRT (Takara Biomedical Technology, Japan). Relative expression was calculated by using $\Delta\Delta Ct$ (ABI, USA). Sequences of the primers used for this reaction are in Table 2.

Protein extraction and western blotting

Whole cells were lysed in RIPA buffer with 1mM PMSF (Sigma, USA), 0.5% sodium deoxy cholate. Extracted proteins were resolved by 12% SDS-PAGE and transferred to a polyvinylidenedifluoride (PVDF, Pall, USA) membrane. Then proteins were blocked with TBS-T containing 5% skim milk (Biotopped, China) at RT for 1 h and incubated overnight at 4°C with the primary antibody (1:1000). The PVDF membrane was washed in TBS-T and incubated for 2 h at room temperature with secondary antibody (1:5000). The bands were detected using enhanced chemiluminescence analysis system (GE, USA), and quantified by densitometry with Amersham Imager 600 uv (GE, USA) software. For quantitation of phosphorylated protein, phosphorylated protein was normalized to the corresponding total protein and then was normalized to the control value.

Antibodies

The specific primary antibodies were purchased from Cell Signaling Technology (CST, USA) and Snata Cruz (Snata Cruz, USA). The secondary antibodies were purchased from Beijing zhongshanJinqiao Biotechnology Co., Ltd, China. The antibodies were listed in the Table 3.

Online analysis

The main mode of action (MOA) of quercetin as an estrogen receptor agonist was analysed from the "Large Scale Visualization of Drug Induced Transcriptomic Signatures" (<http://amp.pharm.mssm.edu/dmoa/report/BRD-K97399794>) website. The LncSHGL promoter sequence was searched from the website of the National Center for Biotechnology Information (NCBI, <https://www.ncbi.nlm.nih.gov/>). Binding sites for LncSHGL and ER α were predicted on the Jasper website (<http://jaspar.genereg.net/analysis>).

Correlation analysis

The Fragments Per Kilobase of exon model per Million mapped fragments (FPKM) expression data of liver cancer was download from TCGA database. The data was transformed from FPKM to TranscriptsPerKilobase of exonmodel per Million mapped reads (TPM). Genes or ncRNA related to diabetes were obtained by KEGG pathway analysis and literature search. Expression correlations between ESR1 and these genes or ncRNA were calculated and plotted in R package ggplot2.

Statistical analysis

The data were analyzed for statistical significance with the SPSS 12.0.1 Package (SPSS Inc., USA). Briefly, data for all groups were first tested for normality with the SPSS software. Before analyzing each set of data, the data was tested for their normal distribution by SPSS software, and all data were in a normal distribution. Homogeneity of variance test was used to test the homogeneity of the variance of the data before performing the analysis of variance. In the homogeneity test of variance, the variance of each group was equal. A one-way analysis of variance was tested. *P* values < 0.05 were regarded as statistically significant. All values are presented as the means \pm SEM (standard error of mean).

Results

Quercetin, one of the most abundant active compounds in GPE, inhibits HFD-induced type 2 diabetes mellitus in mice

GPE was found to inhibit HFD-induced type 2 diabetes mellitus (Fig. 1). GPE significantly reduced HFD-induced weight gain and fasting serum glucose level in mice (Fig. 1A, B). Additionally, GPE significantly reduced the weight of epididymal fat in HFD-induced mice (Fig. 1E). We also examined the food intake of mice and the expression changes of food intake-related genes in the hypothalamus of mice. As show in Fig. 1C and Fig. 1D, GPE did not change the intake of food. To further understand the anti-diabetic effects, GPE was analyzed by ultra-performance liquid chromatography (UPLC) and quadrupole time-of-flight mass spectrometry (Q-TOF-MS) (Fig. 1F). We tested the inhibitory efficiency of several major components in grape extract on α -galactosidase, and found that among these components, quercetin can significantly inhibit α -galactosidase activity (Fig. 1G).

To examine the major efficacy of quercetin, 0.1g/kg quercetin was added to the HFD. No change in the body weight was observed in mice fed NC/quercetin. A daily intake of HFD was 1.54-2.1g, the energy per

gram of HFD was 4.057 Kcal; therefore each mouse consumed 6.25-8.52 Kcal per day. No difference in the intake of food between quercetin-treated group and control was noticed (Fig. 2C). However, quercetin improved hyperglycemia (Fig. 2A, B), IGT and IR in HFD-induced mice (Fig. 2D, E). Similar to GPE, quercetin significantly reduced the liver weight and the gonadal lipid content in HFD-induced mice (Fig. 2F). Quercetin significantly inhibited the HFD-induced body fat increase, as well as significantly inhibited the HFD-induced muscle percentage decrease (Fig. 2G). Quercetin can significantly reduce hepatic steatosis and gonadal coronal structures in HFD-induced mice (Fig. 2H). Moreover, by immunocytochemistry (IHC) analysis of the UCP1 (a marker for brown fat), we observed that quercetin can improve the whitening of brown fat in mice induced by HFD (Fig. 3A).

Quercetin significantly increases the expression of diabetes-associated lncRNA through ER signal

By using the “Large Scale Visualization of Drug Induced Transcriptomic Signatures” website, we noticed that quercetin is an estrogen receptor agonist; this was the verified data from the cell experiments (Fig. 3B).

Recently, a group of transcriptome sequencing analyses showed that lncRNA transcripts were found in the intergenic regions of the estrogen receptors binding site, and play a fundamental role in estrogen-dependent transcriptional responses [28], and are prerequisite conditions for the regulation of the downstream network of the steroid receptors [29, 30]. ERs have been reported to inhibit diabetes, but the molecular mechanism is unclear [31]. Through literature search and KEGG pathway analysis, genes and ncRNAs related to diabetes were obtained and calculated the correlation with ER α in TCGA liver cancer dataset (supplementary table 1 and supplementary Fig. 1). The correlation between ER α and these genes was plotted in table S1 and Fig. S1. SLC2A2, B4GALT1-AS1, FAS, IRS1, HNF4A and PIK3R1 were found to be significantly correlated with ER α ($R > 0.2$, $P < 0.05$). We examined the expression changes of these genes on the human liver cancer cell line HepG2 and found that quercetin can significantly increase the expression of B4GAT1-AS1 (Fig. S2). Next, we verified on mice liver and found that among these related genes; quercetin can significantly increase lncSHGL expression (Fig. S3). Several lncRNAs are involved in the etiology of type 2 diabetes mellitus and its complications [17, 32-34]. To investigate whether quercetin elicits its anti-diabetic effects via lncRNAs, we measured the levels of lncRNAs in HFD-induced and quercetin-treated mice. The results showed an increased lncSHGL expression after quercetin treatment (Fig. 3C). A previous study has reported that lncSHGL induces AKT phosphorylation by recruiting heterogeneous nuclear ribonucleoprotein A1 (hnRNPA1), thereby improving type 2 diabetes mellitus symptoms in mice [17]. Quercetin increased hnRNPA1 expression in the QHFD group compared with HFD group (Fig. 3D). Moreover, we observed that quercetin can increase the expression of lncSHGL and hnRNPA1 in the livers of mice (Fig. 3C, D).

Through the Jasper website, we noticed that ER α has the predicted binding sites within the lncSHGL promoter region (Fig. 3E). To examine the quercetin-mediated mechanism responsible for the improvement of type 2 diabetes mellitus symptoms in mice fed a HFD, we carried out several experiments. We found that quercetin increased the expression of lncSHGL and hnRNPA1 in mouse

primary hepatocytes, mouse normal liver cell line NCTC-149 cells, and human hepatoma HepG2 cells. Furthermore, quercetin and the ER agonist increased the expression of lncSHGL and hnRNPA1 in the palmitic acid treated cells (Fig. 4A-4F). With regard to quercetin, opposing effects were observed in the presence of ER antagonists (Fig. 4A-4F). Palmitic acid decreased the phosphorylated AKT level in HepG2 cells, whereas quercetin increased it (Fig. 4G). Taken collectively, the results show that quercetin can activate AKT signaling by promoting the expression of lncSHGL and hnRNLA1, thereby inhibiting HFD-induced type 2 diabetes mellitus.

Quercetin increases lncSHGL expression by ER α

Data analysis performed with online tools indicated that quercetin is an ER agonist. As shown in Fig. 3B, both E₂ and ER agonists significantly increased lncSHGL expression in cells, whereas ER antagonists decreased lncSHGL expression. We predicted the possible binding regions of ER α and lncSHGL (Fig. 3E). We constructed the predicted lncSHGL promoter-binding region into a luciferase expression vector, co-transfected cells with ER α and ER β overexpression vectors, and measured the fluorescence. The results of the double luciferase experiment showed that quercetin and estrogen can increase the binding of ER α and lncSHGL promoter regions (Fig. 5A). The fluorescent signal was significantly higher in cells transfected with the ER α overexpression vector than in those transfected with the ER β overexpressing vector (Fig. 5B). We mutated the predicted binding site in the promoter region of lncSHGL, and then co-transfected cells with ER α to detect the luciferase activity. As shown in Fig. 5C, luciferase expression was significantly reduced following mutation of the binding site of the promoter region of lncSHGL.

Discussion

With increasing daily responsibilities and challenges, many individuals do not pay adequate attention to their health. For instance, many individuals are obese, which can cause type 2 diabetes mellitus, due to poor dietary choices and unreasonable social pressures. Currently, type 2 diabetes mellitus is a major disease that can severely endanger human health. Several drugs can effectively suppress the increased serum glucose levels in patients. However, the side effects of these drugs cannot be ignored, as they are a concern for many individuals. Several studies have reported that natural compounds, such as those found in grapes, can curb the symptoms of type 2 diabetes mellitus and other metabolic syndromes, but their mechanisms of action are unclear [35–37]. Consistent with the results of other studies, we report that GPE suppressed the symptoms of HFD-induced type 2 diabetes mellitus. However, the mechanism of action remains unclear.

By UPLC and Q-TOF-MS, we identified quercetin to be one of the most active compounds in GPE. Quercetin is a flavonoid. These compounds have received much attention due to their structural diversity, richness, strong pharmacological activity, and safety [38]. Numerous studies have reported the biological effects of quercetin, and its potential benefits as an anti-cancer, anti-oxidative, and anti-inflammatory agent, especially in the treatment of type 2 diabetes mellitus [39–42]. Quercetin can improve the symptoms and eliminate the complications of diabetes [38, 43, 44]. Liu and colleagues have reported the

involvement of quercetin in the insulin-mediated regulation of the glucose level in skeletal muscle through AMPK [43]. Soo-Mi and colleagues have demonstrated that quercetin can improve hyperglycemia and dyslipidemia in type 2 diabetes mellitus db/db mice [44]. Consistent with the results of previous studies, our findings indicated that quercetin reduced HFD-induced hyperglycemia and improved IR.

Recently, a group of transcriptome sequencing analyses showed that LncRNA transcripts were found in the intergenic regions of the estrogen receptors binding site, and play a fundamental role in estrogen-dependent transcriptional responses [28], and are prerequisite conditions for the regulation of the downstream network of the steroid receptors [29, 30]. Several lncRNAs are involved in the etiology of type 2 diabetes mellitus and its complications [17, 32–34]. To define the mechanism of action by which quercetin improved the symptoms of type 2 diabetes mellitus in mice fed a HFD, we followed previous studies that have reported an association between lncRNAs and type 2 diabetes mellitus [45–47]. lncRNAs have key roles in various biological processes [16, 19]. For example, seven lncRNAs have been described to function in metabolic processes in the liver. We report that quercetin significantly increased lncSHGL expression after validating the expression of the seven lncRNAs in the livers of the four groups of mice. lncSHGL sequesters hnRNPA1 to inhibit hepatic gluconeogenesis and lipogenesis, which improves the symptoms of diabetes [17].

Our *in vitro* and *in vivo* results demonstrate that quercetin improved the symptoms of type 2 diabetes mellitus in mice fed a HFD. In addition, our results illustrate for the first time an association between ER α and lncSHGL. Quercetin increased lncSHGL expression by promoting the binding of ER α to the lncSHGL promoter region.

The limitation of this study is that it lacks clinical data; essentially, there is no evidence to indicate that quercetin is effective in diabetic patients. The effects of quercetin in diabetic patients will be investigated in a subsequent study.

Here, we reported the mechanism of action by which GPE improved the symptoms of type 2 diabetes mellitus in mice fed a HFD. Quercetin is one of the most active compounds in GPE, and it promoted ER α binding to the lncSHGL promoter region, thereby increasing lncSHGL expression and suppressing the symptoms of type 2 diabetes mellitus induced by a HFD (Fig. 5C). Our results show that quercetin could be beneficial for type 2 diabetes mellitus by promoting lncSHGL transcription and activating the lncSHGL pathway, and may be used as a drug component to treat type 2 diabetes mellitus.

Conclusions

We found that GPE could improve T2DM and identified quercetin as one of the most abundant and active compounds in GPE. Quercetin could inhibit HFD-induced T2DM in mice by activating ER α . lncSHGL was identified as the downstream of ER α and inhibited HFD-induced T2DM. Our results show that quercetin could be beneficial for T2DM by promoting lncSHGL transcription and activating the lncSHGL pathway, and may be used as a drug component to treat T2DM.

Abbreviations

DPN
diarylpropionitrile
ER α
estrogen receptor α
ER β
estrogen receptor β
ERs
estrogen receptors
E2
estradiol
FPKM
Fragments Per Kilobase of exon model per Million mapped fragments
GSE
Grape seed extract
GPE
Grape pomace aqueous
HFD
high fat diet
hnRNPA1
heterogeneous nuclear ribonucleoprotein A1
ICI
ICI 182780
IHC
immunocytochemistry
IR
insulin resistance
LncRNA
long non-coding RNA
LncSHGL
LncRNA suppressor of hepatic gluconeogenesis and lipogenesis
PPT
propylpyrazoletriol
Q-TOF-MS
quadrupole time-of-flight mass spectrometry
TPM
Transcripts Per Kilobase of exon model per Million mapped reads
T2DM
type 2 diabetes mellitus

Declarations

1. Ethics approval and consent to participate

All animal experimental protocols are in line with the Animal Management Rules of the Ministry of Health of the People's Republic of China and the China Agricultural University Guide for the Care and Use of the Laboratory Animals. The ethical committee No. is CAU20180217-4.

2. Consent for publication

Not applicable.

3. Availability of data and materials

The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

4. Competing interests

The authors declare that they have no competing interests.

5. Funding

No funding.

6. Author's Contributions

Fan hanlu and Li xiangdong designed these experiments; Fan hanlu, Li haiwen, Liu huijiao, Li peng, Jia xiaomeng and Guo peilan did these experiments; Fan hanlu and Li xiangdong wrote the main manuscript text; Fan hanlu prepared figures. All authors reviewed the manuscript.

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Tables

Table 1. Macronutrient and selected micronutrient content in the mouse high fat diet

	gm%	kcal%
Protein	26.2	20
Carbohydrate	26.3	20
Fat	34.9	60
Ingredient	gm	kcal
Casein, 30 Mesh	200	800
L-Cystine	3	12
Corn Starch	0	0
Maltodextrin 10	125	500
Sucrose	68.8	275.2
Cellulose, BW200	50	0
Soybean Oil	25	225
Lard*	245	2205
Mineral Mix S10026	10	0
DiCalcium Phosphate	13	0
Calcium Carbonate	5.5	0
Potassium Citrate, 1 H ₂ O	16.5	0
Vitamin Mix V10001	10	40
Choline Bitartrate	2	0
FD&C Blue Dye#1	0.05	0
Total	773.85	4057

Table 2 List of oligonucleotide primer pairs used in RT-PCR analyses

Name	Sequences
H-GAPDH F	TGTGTCCGTCGTGGATCTGA
H-GAPDH R	TTGCTGTTGAAGTCGCAGGAG
M- β -actin F	AGCCATGTACGTAGCCATCC
M- β -actin R	GCTGTGGTGGTGAAGCTGTA
M-LncSHGL F	GACGCGTTTTCTATCGTGC
M-LncSHGL R	CCAGGCCCTCCATCATTGT
M-hnRNPA1 F	ATGTCTAAGCCGAGTCTCCCA
M-hnRNPA1 R	TGTTAGTGTTCCCATGCTC
H-B4GALT1 F	AATCCATGTCTATAACTTCCT CAGT
H-B4GALT1 R	AAGGATTCAGTTTATTCTCTGGCT
H-hnRNPA1F	GGAACGCTCACGGACTGT
H-hnRNPA1R	GTGGGCACCTGGTCTTTG

Table 3 List of the antibodies for western blot

Antibody	Dilution	Cat.nr.	Source	Company
P-AKT	1:2000	#4060S	Rabbit monoclonal IgG	CST
AKT	1:2000	9272S	Rabbit monoclonal IgG	CST
β -actin	1:1000	sc-47778	Mouse monoclonal IgG	Santa Cruz

Figures

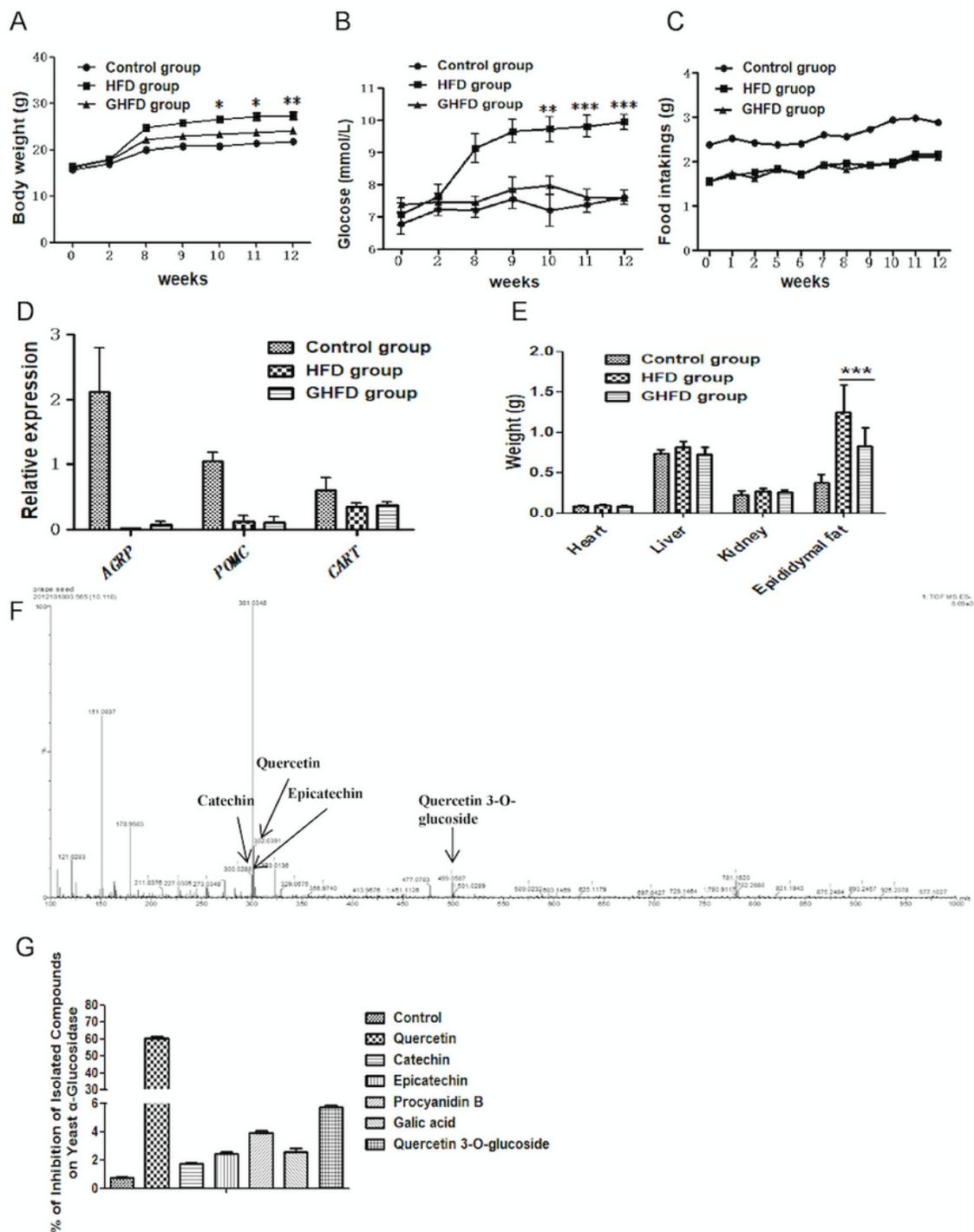


Figure 1

Grape pomace extract improves the symptoms of type 2 diabetes mellitus in mice fed a HFD. Grape pomace extract (GPE) improves the symptoms of type 2 diabetes mellitus in mice fed a HFD. (A) Body weight (n = 16) (GHFD group versus HFD group, **, p < 0.01; *, p < 0.05), (B) fasting plasma glucose level (n = 16) (GHFD group versus HFD group, **, p < 0.01; *, p < 0.05), (C) food intake (n = 15), (D) expression

of hypothalamic feed intake-related genes, (E) organ weight of each group of mice (n = 16) , (F) UPLC and Q-TOF-MS results and (G) detect the efficiency of inhibition of α -galactosidase activity.

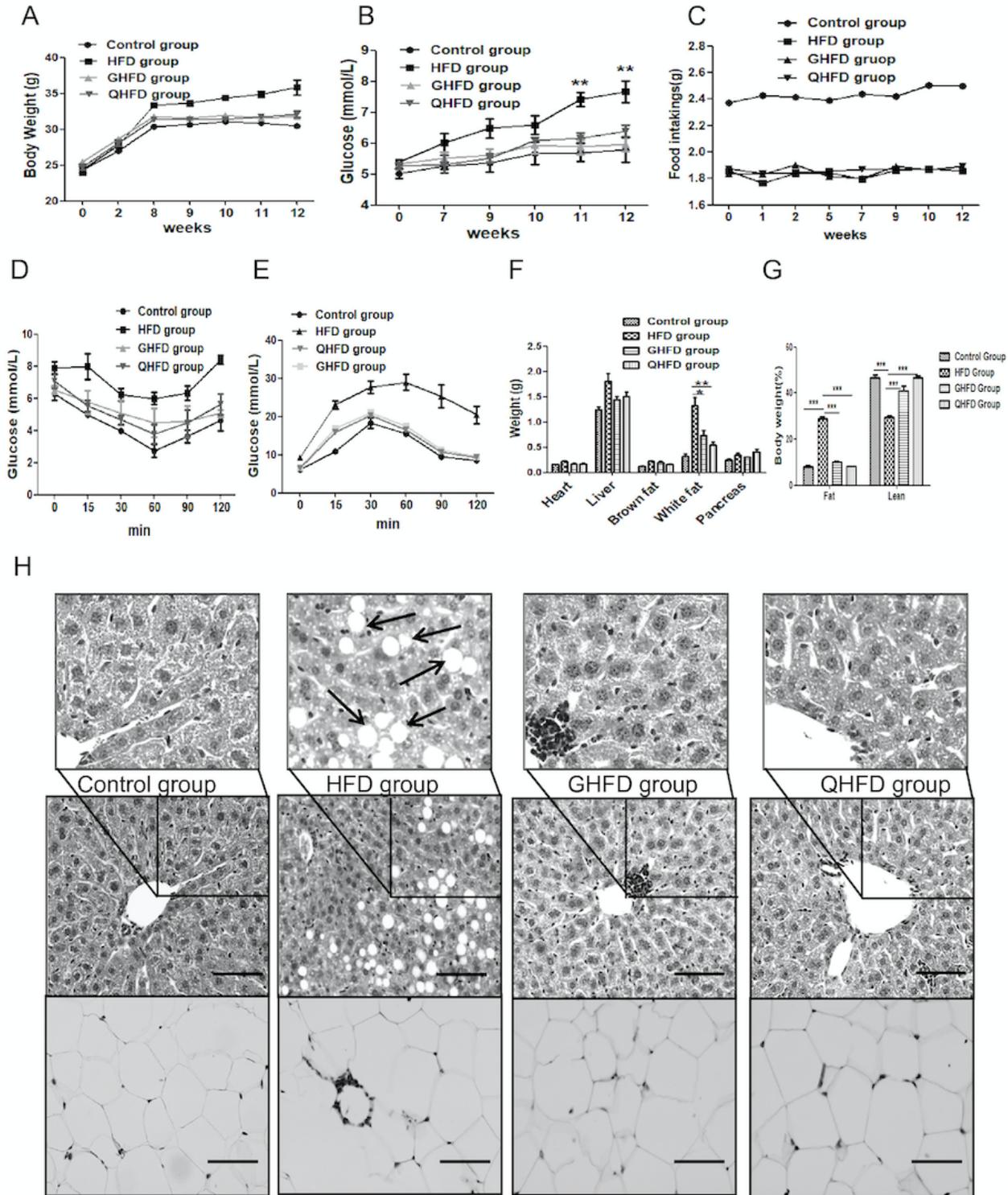


Figure 2

Quercetin can reduce HFD-induced type 2 diabetes mellitus in mice. Quercetin can reduce HFD-induced type 2 diabetes mellitus in mice. (A) Body weight (n = 16) (QHFD group versus HFD group, **, p < 0.01; *, p < 0.05), (B) fasting plasma glucose level (n = 16) (QHFD group versus HFD group, **, p < 0.01; *, p < 0.05),

(C) food intake (n = 16) , (D) ITT results of each group of mice (n = 16), (E) GTT results of each group of mice (n = 16) (F) organ weight of each group of mice (n = 16) (QHFD group versus HFD group, **, p < 0.01; *, p < 0.05), (G) body fat percentage and lean rate in mice and (H) paraffin-embedded, hematoxylin and eosin stained sections of the liver and gonads above the liver, and below the epididymal fat (magnification bar, 10µm).

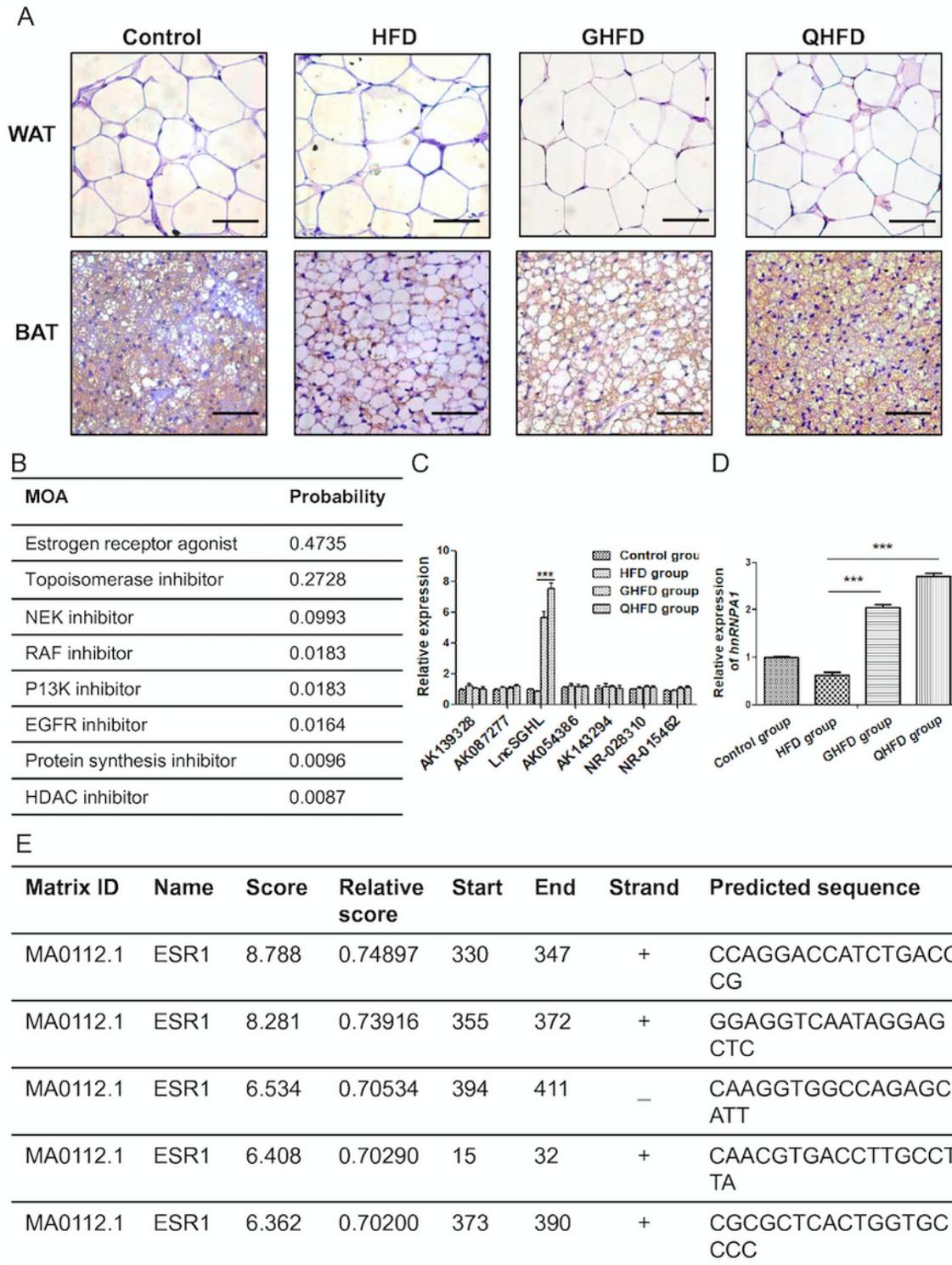


Figure 3

Quercetin promotes LncSHGL expression via ER. Quercetin promotes LncSHGL expression via ER. (A) Immunohistochemical results that UCP1 of four groups WAT and BAT (magnification bar, 10 μ m), (B) quercetin MOA analysis (C) LncRNA expression in the liver, (n = 5) (QHFD group versus HFD group, **, p < 0.01; *, p < 0.05), (D) hnRNPA1 expression in the liver, (n = 5) (GHFD group versus HFD group and QHFD group versus HFD group, **, p < 0.01; *, p < 0.05), and (E) prediction of the binding site between ER and the LncRNA promoter region.

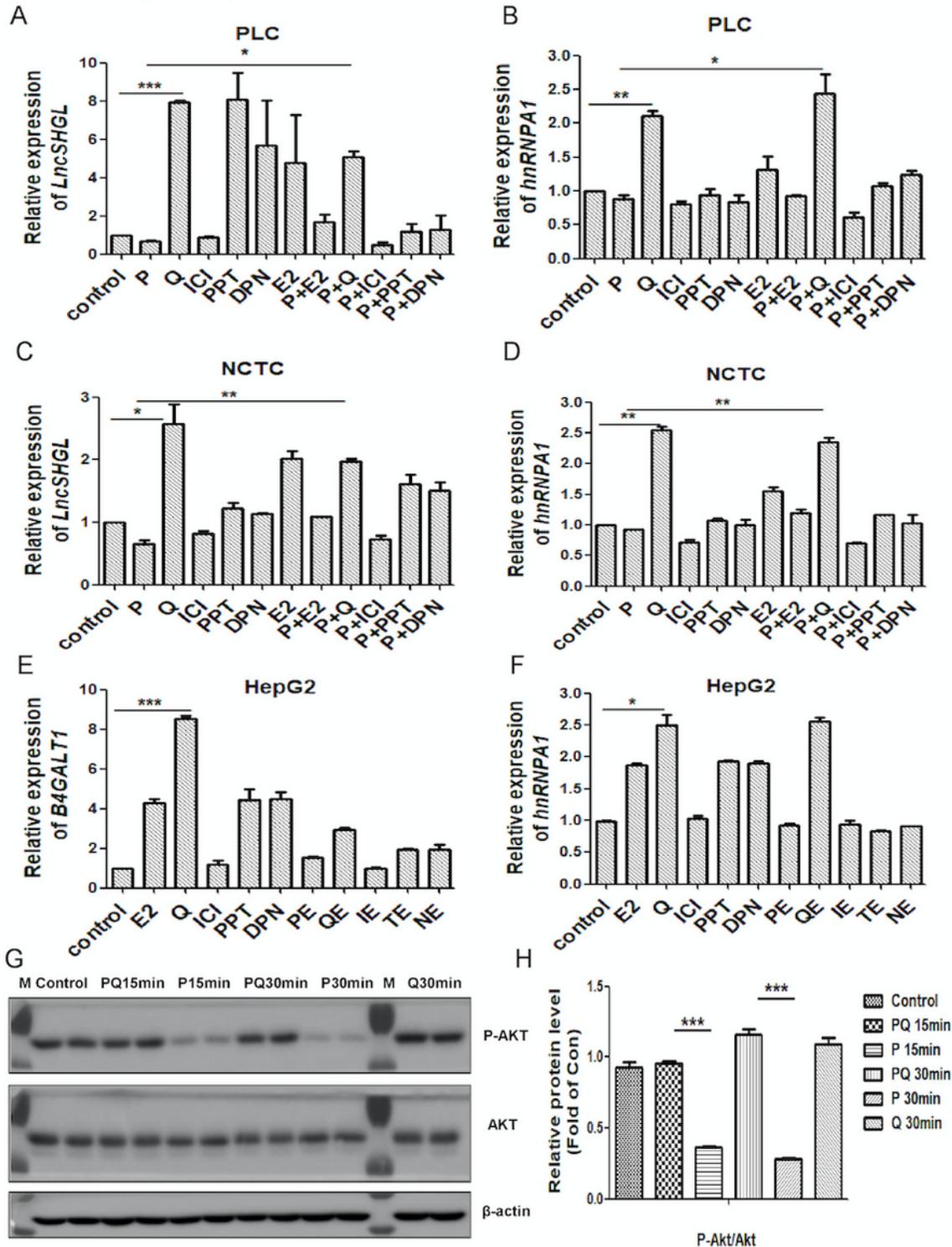


Figure 4

Quercetin works with estrogen and ER agonists to increase lncSHGL expression. Quercetin works with estrogen and ER agonists to increase lncSHGL expression. (A) lncSHGL expression in primary hepatocytes after various treatments, (n = 5) (Control versus Q and P versus P+Q, **, p < 0.01; *, p <0.05), (B) hnRNPA1 expression in primary hepatocytes after various treatments, (n = 5) (Control versus Q and P versus P+Q, **, p < 0.01; *, p <0.05), (C) lncSHGL expression in NCTC cells after various treatments, (n = 5) (Control versus Q and P versus P+Q, **, p < 0.01; *, p <0.05), (D) hnRNPA1 expression in NCTC cells after various treatments, (n = 5) (Control versus Q and P versus P+Q, **, p < 0.01; *, p <0.05), (E) lncSHGL expression in HepG2 cells after various treatments, (n = 5) (Control versus Q, **, p < 0.01; *, p <0.05), (F) hnRNPA1 expression in HepG2 cells after various treatments, (n = 5) (Control versus Q, **, p < 0.01; *, p <0.05), (G) changes in the phosphorylated AKT protein level and (H) corresponding quantification of protein.

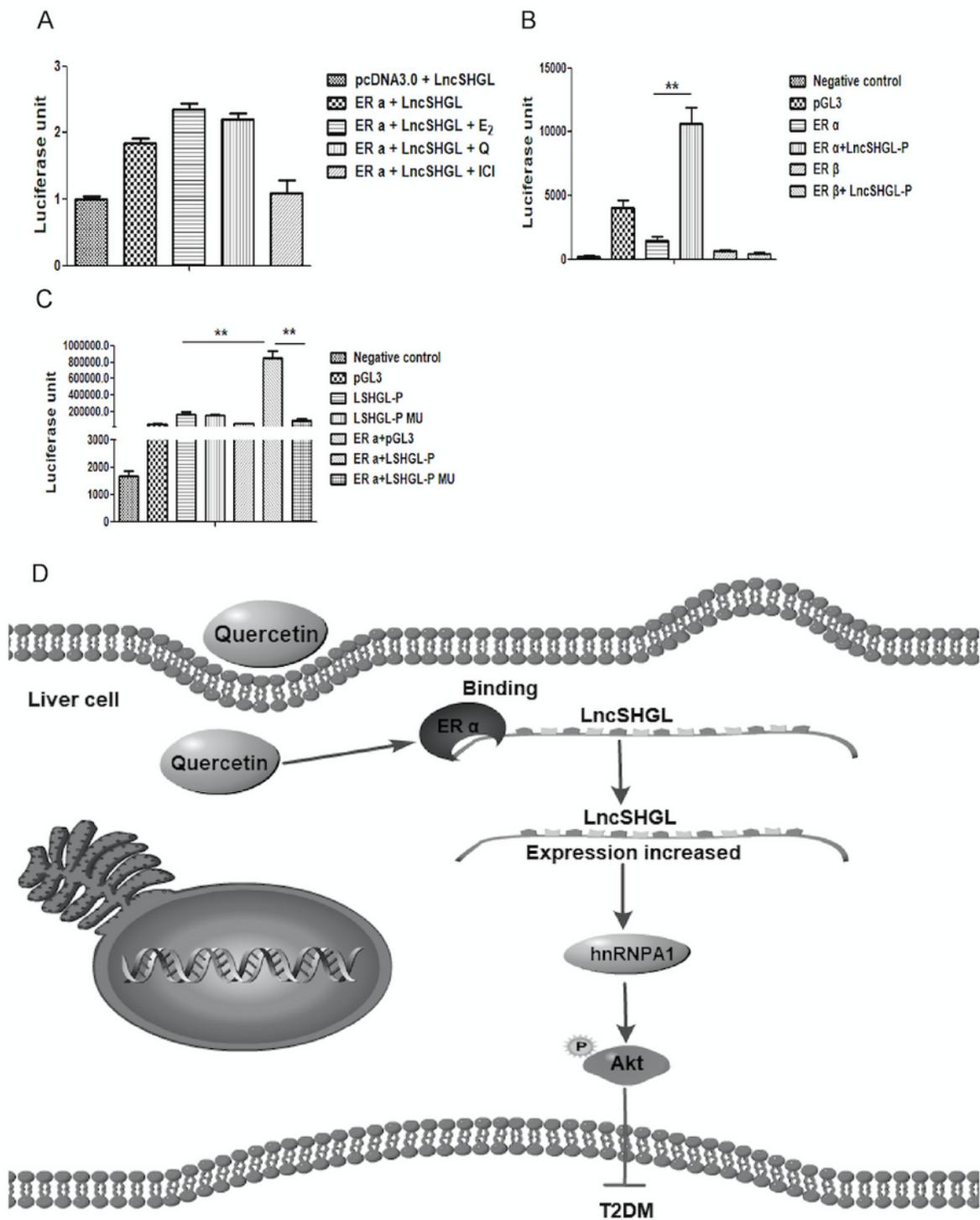


Figure 5

The LncSGHL promoter region binds to ERα. The LncSGHL promoter region binds to ERα. (A) Luciferase results, (n=4) (ER α + LncSHGL versus ER α + LncSHGL + E₂ ; ER α + LncSHGL versus ER α + LncSHGL + Q and ER α + LncSHGL versus ER α + LncSHGL + ICI, **, p < 0.01; *, p < 0.05)(B)Luciferase results, (n = 6) (ER α versus ER α + LncSHGL-P, **, p < 0.01; *, p < 0.05), (C) luciferase results, (n = 6) (LncSHGL-P versus ER α + LncSHGL-P and ER α + LncSHGL-P versus ER α + LncSHGL-P MU, **, p < 0.01; *, p < 0.05) and (D)

quercetin promoted ER α binding to the IncSHGL promoter region, thereby increasing IncSHGL expression and suppressing the symptoms of type 2 diabetes mellitus induced by a HFD.

Supplementary Files

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