Effects of liraglutide on lipolysis and the AC3/PKA/HSL pathway

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Background: Liraglutide reduces blood glucose, body weight and blood lipid levels. Hormone-sensitive lipase (HSL) is a key enzyme in lipolysis. Evidence from our and other studies have demonstrated that adenylate cyclase 3 (AC3) is associated with obesity and can be upregulated by liraglutide in obese mice. In the present study, we investigated whether hepatic HSL activity is regulated by liraglutide and characterized the effect of liraglutide in the AC3/protein kinase A (PKA)/HSL signalling pathway.

Methods: Obese mice or their lean littermates were treated with liraglutide or saline for 8 weeks. Serum was collected for the measurement of insulin and lipids. We investigated hepatic AC3, HSL and phosphorylated HSL Ser-660 (p-HSL(S660)) protein expression levels and AC3 and HSL mRNA expression levels and cyclic adenosine monophosphate (cAMP), PKA activity in liver tissue.

Results: Liraglutide treatment decreased triglycerides (TGs) and free fatty acids (FFAs), increased glycerol, and upregulated hepatic AC3 and p-HSL(S660) levels and cAMP and PKA activities.

Conclusion: The results suggest that liraglutide can upregulates AC3/PKA/HSL pathway and may promotes lipolysis.

Keywords: liraglutide, obesity, lipolysis, adenylate cyclase 3, hormone-sensitive lipase

Introduction

In recent years, as the proportion of obese people in the world’s population has grown rapidly, it has attracted more and more attention. A key feature of obesity is excessive lipid accumulation in adipose tissue and ectopic localizations (especially liver and muscle). Liraglutide treatment can augment weight loss and improve blood lipid levels. Lipolysis is considered to be the process of hydrolysis of triglycerides (TGs) to free fatty acids (FFAs) and glycerin. Hormone-sensitive lipase (HSL) in adipose tissue has been suggested to be a key regulatory enyzm in controlling lipolysis. Endogenous glucagon-like peptide 1 (GLP-1) binds to and activates the glucagon-like peptide 1 receptor (GLP-1R). Upon its activation, GLP-1R stimulates adenylate cyclase (AC) to promote cAMP production, leading to activation of protein kinase (PKA). Activated PKA phosphorylates hormone-sensitive lipase (p-HSL). AC3 is a member of the ACs family, genome-wide association studies suggest that obesity-related genes are located on or near the AC3 gene. In both Swedish and Han Chinese populations, additional genetic evidence supports that single nucleotide variation of AC3 gene was found to be closely related to obesity. It is also emphasized in animal models that AC3 signalling plays an important role in maintaining energy homeostasis, AC3 mutation can protect mice from diet-induced obesity. Adult weight of AC3 knockout mice is
significantly higher than that of wild-type mice. It is also found that TG of AC3 knockout mice is increased, and the fat mass of epididymal adipocytes of AC3 knockout mice is more than that of control mice.\textsuperscript{18} Our previous studies have shown that liraglutide can upregulate hepatic GLP-1R and AC3 levels in obese mice, and the levels of AC3 were negatively correlated with body weight.\textsuperscript{19,20} Thus, we hypothesize that liraglutide treatment can enhance lipolysis and affect the AC3-cAMP-PKA-HSL signalling pathway. To test our hypothesis, we employed an obese mouse model in which C57BL/6J mice were fed a high-fat diet (HFD). TGs, glycerol, FFA in serum and PKA activity were analysed in the liver, and the expression levels of proteins associated with lipolysis were determined in liver. The results of this study will increase our knowledge of the effect of liraglutide on lipolysis, leading to a more comprehensive understanding of the mechanisms underlying the physiological actions of liraglutide on obesity.

Materials and methods

Animal husbandry and analytic procedures

4-week-old C57BL/6J mice (male) were purchased from the Medical Laboratory Animal Centre of Guangzhou Province (Guangzhou, China). The experimental mice were maintained at the Experimental Centre of Guangxi Medical University. The experimental mice were maintained in a specific pathogen-free (SPF) room with a 12-h light/dark cycle. In the first week, all mice were fed with a normal rodent chow diet (5% fat wt/wt). After then, the mice were randomly divided into two groups. ie N group: the mice fed with a normal chow diet (5% fat wt/wt) and O group: the mice fed with HFD (34.9% fat wt/wt). Body weight (BW) and blood glucose of all experimental mice were maintained at the Animal Experiment Centre of Guangxi Medical University. The experimental mice were considered obese. After 12 weeks of chow diet or HFD feeding, the obese mouse model was successfully established, all mice were divided into the following four groups: N + saline (N + S), N + liraglutide (N + L), O + saline (O + S) and O + liraglutide (O + L). In N + L and O + L groups, liraglutide was injected subcutaneously at a dose of 0.1 mg/kg/12 h. N + S and O + S groups were subcutaneously injected with the same volume of saline as controls. After 8 weeks of treatment, the experimental mice were fasted overnight and anaesthetized with sodium pentobarbital (50 mg/kg, i.p.). The blood was obtained through the angular vein and centrifuged by a 4 °C centrifuge to separate the serum, which was stored at −20 °C. At the same time, the liver was dissected immediately and frozen in liquid nitrogen, then transferred to −80 °C refrigerator for preservation until analysis. All animal experiments and care procedures were conducted under the Guidelines of the Animal Ethics Committee of First Affiliated Hospital of Guangxi Medical University (the National Standard GB/T35892-2018 of the People’s Republic of China). The Animal Ethics Committee of First Affiliated Hospital of Guangxi Medical University also approved this study. Liraglutide was kindly provided by Novo Nordisk (Bagsvaerd, Denmark).

Biochemical analysis of serum and liver

The mouse TGs, glycerol, FFA and insulin ELISA kit (Shanghai JiNing Industrial Co., Ltd., China) was used to measure serum TGs, glycerol, FFA and insulin levels. Liver samples were prepared according to the instructions provided. A PKA Kinase Assay Kit (Enzo Life Sciences Inc. USA) was used to measure PKA activity in liver samples. A glucose meter (Johnson & Johnson, New Brunswick, NJ, USA) was used to measure blood glucose levels. The HOMA-IR was used to assess IR. The HOMA-IR score was calculated as [(fasting insulin (mU l\(^{-1}\)) × fasting glucose (mmol l\(^{-1}\)))/22.5].\textsuperscript{22}

Real-time reverse transcription-PCR

Total RNAs in liver were extracted, and a Thermo Reverse Transcriptase Kit (Thermo Scientific, Waltham, MA, USA) was used to synthesize cDNA. Quantitative real-time PCR was performed using FastStart Universal SYBR Green Master (ROX) (Roche Diagnostics, Indianapolis, IN, USA) on an Applied Biosystems StepOnePlus Real-Time PCR System (Thermo Fisher Scientific, Foster City, CA, USA). The primers included the following: AC3 forward primer (5’-GGACACCGCTCAACACATC-3’) and reverse primer (5’-GCCACATTGACCGTGATTGC-3’). HSL forward primer (5’-T CCTGGAACATAGTGAGCAAG-3’) and reverse primer (5’-CAGACACACTCTCCTGCGATAGAC-3’). The glyceraldehyde 3-phosphatedehydrogenase (gpdh) was used as the internal control. Gapdh forward primer (5’-TGGTGTCGGTGATCTGA-3’) and reverse primer (5’-TTGGCTTG TGAAGTCGCAGGAG-3’). The relative copy number was calculated using the threshold crossing point (Ct) as calculated by the LightCycler software combined with the \( \Delta \Delta \text{Ct} \) calculations.
Western blotting
Western blot analyses were performed as previously described.23 The primary antibodies included antibodies against the AC3 (1:1000; Sigma-Aldrich Co., LLC., China), cAMP (1:500; Abnova Co., Inc., USA), HSL (1:500; Bioworld Technology Co., Ltd., China) and phosphorylated-HSL Ser660 (p-HSL(S660)) (1:500; Cell Signal Technology Inc., USA). Horseradish peroxidase-conjugated goat anti-rabbit IgG was used as the secondary antibody. The membranes were scanned with the Odyssey infrared imaging system (LICOR Biosciences).

Statistical analysis
Quantitative values are shown as the means ± SD. Comparisons were performed using Student’s t-test, the one-way analysis of variance (ANOVA) or a factorial analysis, where appropriate. In all cases involving the use of Student’s t-test, one-way ANOVA analysis or factorial analysis, a homogeneity of variance test was applied. Values of p<0.05 were considered statistically significant. All analyses were performed using SPSS 17.0 (Chicago, IL, USA).

Results
Effects of liraglutide on the general condition and metabolism of the mice
BW, FBG, HOMA-IR, and the serum levels of TGs, glycerol, FFA were to evaluate the metabolic response of liraglutide in mice. Mice in the N group exhibited significantly reduced BW compared with mice in the O group (Table 1). FBG levels were elevated in the O group relative to those in the N group (Table 1). Factorial analysis showed that BW and HOMA-IR scores were significantly different between normal control mice and obese mice, and there were significant differences between liraglutide-treated mice and saline-treated mice (Table 2). There was no significant difference in FBG between normal control group and obesity group, nor between liraglutide-treated mice and saline-treated mice (Table 2). The serum levels of TGs, glycerol and FFA were significantly higher in obese mice than in normal control mice. Liraglutide treatment significantly decreased the serum levels of TGs, FFA play a pivotal role in insulin resistance. Many studies have demonstrated that excessive FFAs play a pivotal role in insulin resistance.26–28 Our data show that liraglutide treatment decreased serum TGs and FFAs and increased serum glycerol. The decrease in TGs and the increase in glycerol indicated that liraglutide treatment promoted lipolysis. The decrease in FFAs is likely due to liraglutide increasing FFA uptake and increasing the oxidation rate of FFAs. Mells, Fu, Sharma, etc.29 found that liraglutide treatment significantly increased the mRNA and protein levels of genes related to fatty acid uptake and peroxisomal β-oxidation.

Effects of liraglutide on hepatic AC3, HSL, p-HSL(S660), and cAMP levels in mice
We examined AC3, HSL, p-HSL(S660) protein levels and AC3, HSL mRNA levels and cAMP levels in liver tissue; the results are summarized in Figures 1 and 2. The hepatic levels of AC3 and p-HSL(S660) protein and AC3 mRNA and cAMP were significantly different between normal control mice and obese mice. The hepatic levels of AC3 and p-HSL(S660) protein and AC3 mRNA and cAMP were lower in obese mice than in normal control mice. No significant differences in HSL protein and HSL mRNA levels were observed between normal control mice and obese mice. Liraglutide treatment significantly increased hepatic AC3, p-HSL(S660) and cAMP levels but not HSL protein and mRNA levels in normal control and obese mice. The increase of hepatic AC3, p-HSL(S660) and cAMP levels in obese mice were more obviously.

Effects of liraglutide on hepatic PKA activity levels
Liver PKA activity was significantly higher in normal control mice than in obese mice (Figure 3). Liraglutide treatment significantly increased liver PKA activity in normal control and obese mice (Figure 3). The increase in obese mice was more obviously.

Discussion
In this study, obesity in mice was induced by a high-fat diet. We found that treatment with liraglutide decreased the BW and HOMA-IR index in obese mice and normal control mice, though the decreases were stronger in obese mice. Our data showed that the serum levels of TGs, glycerol and FFA were significantly higher in obese mice than in normal control mice. Liraglutide treatment significantly decreased serum levels of TGs and FFA, and significantly increased serum glycerol levels.

It has been repeatedly demonstrated that liraglutide can reduce TG and FFA levels.2–5,24 Lipolysis is the process by which TGs are hydrolysed to FFAs and glycerol. Lipolysis was determined by assaying FFA and glycerol levels.25 Many studies have demonstrated that excessive FFAs play a pivotal role in insulin resistance.26–28 Our data show that liraglutide treatment decreased serum TGs and FFAs and increased serum glycerol. The decrease in TGs and the increase in glycerol indicated that liraglutide treatment promoted lipolysis. The decrease in FFAs is likely due to liraglutide increasing FFA uptake and increasing the oxidation rate of FFAs. Mells, Fu, Sharma, etc.29 found that liraglutide treatment significantly increased the mRNA and protein levels of genes related to fatty acid uptake and peroxisomal β-oxidation.
GLP-1 has been reported to reduce hepatic steatosis in animal models of non-alcoholic fatty liver disease (NAFLD). The mechanism underlying this effect of GLP-1 is not completely understood. Earlier studies have suggested that exenatide (exendine-4) had direct effects on the liver by enhancing lipid hydrolysis and oxidation. The mechanism by which liraglutide promotes lipolysis is not completely understood. Our previous study showed that liraglutide treatment upregulated GLP-1R and AC3 and that AC3 gene expression was negatively correlated with BW, HOMA-IR and the area ratio of hepatic fat deposition in the liver. We hypothesized that liraglutide treatment could 

### Table 1 Body weights and fasting blood glucose levels of mice before liraglutide treatment

<table>
<thead>
<tr>
<th></th>
<th>Body Weight (g)</th>
<th>Fasting Blood Glucose (mmol/L)</th>
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<tbody>
<tr>
<td>Normal Control Group</td>
<td>27.6±1.34</td>
<td>6.37±1.03</td>
</tr>
<tr>
<td>Obese Group</td>
<td>33.25±2.81***</td>
<td>9.9±2.21**</td>
</tr>
</tbody>
</table>

Notes: Values are means ± SD (n=12 per group). Student’s t-test was used to analyse differences. Statistically significant difference, p<0.01.

### Table 2 Serum metabolic parameters of mice after treatment with liraglutide or saline

<table>
<thead>
<tr>
<th></th>
<th>N+S</th>
<th>N+L</th>
<th>O+S</th>
<th>O+L</th>
</tr>
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<tbody>
<tr>
<td>Body Weight (g)</td>
<td>27.70±1.49</td>
<td>26.2±1.49**</td>
<td>31.98±2.47##</td>
<td>27.1±1.52##</td>
</tr>
<tr>
<td>FBG (mmol/L)</td>
<td>6.33±1.24</td>
<td>6.9±1.22</td>
<td>7.48±1.42</td>
<td>6.10±0.97</td>
</tr>
<tr>
<td>insulin (mLU/L)</td>
<td>5.42±0.98**</td>
<td>11.52±0.94#</td>
<td>3.84±0.84##</td>
<td>1.69±0.23##</td>
</tr>
<tr>
<td>HOMA-IR</td>
<td>1.98±0.49</td>
<td>1.69±0.53##</td>
<td>32.29±1.75##</td>
<td>16.21±1.76##</td>
</tr>
<tr>
<td>TGs</td>
<td>22.92±3.19</td>
<td>11.28±3.09##</td>
<td>66.51±9.87##</td>
<td>115.00±9.34##</td>
</tr>
<tr>
<td>Glycerol</td>
<td>52.94±4.06</td>
<td>60.53±6.77##</td>
<td>115.17±3.93##</td>
<td>619.56±100.26##</td>
</tr>
<tr>
<td>FFA</td>
<td>498.33±75.22</td>
<td>301.85±70.80##</td>
<td>1151.73±33.93##</td>
<td>619.56±100.26##</td>
</tr>
</tbody>
</table>

Notes: Values are means ± SD (n=6 per group). A 2×2 factorial analysis was used to analyse differences. The p-value of the interaction between obesity and liraglutide treatment for body weight, fasting blood glucose, HOMA-IR, TG, glycerol and FFA are 0.030, 0.065, 0.000, 0.001, 0.016, 0.000, respectively. Between saline and liraglutide, *p<0.05; between normal and obese, #p<0.05, ##p<0.01.

![Figure 1](https://www.dovepress.com/)

**Figure 1** AC3, hormone-sensitive lipase (HSL) and p-HSL(S660) expression at the protein levels and cAMP levels in the liver after treatment with liraglutide or saline. Values are means ± SD (n=6 per group). A 2×2 factorial analysis was used to analyse differences. (A-D) present the levels of AC3 protein, cAMP, HSL protein and p-HSL(S660) protein, respectively, in the liver tissues of mice after treatment with liraglutide or saline. The p-value of the interaction between obesity and liraglutide treatment for AC3 protein, cAMP, HSL protein and p-HSL(S660) protein levels in the livers are 0.049, 0.044, 0.354 and 0.049, respectively. (E) Western blots show hepatic AC3, cAMP, HSL and p-HSL (S660) levels.

Notes: between saline and liraglutide, *p<0.05, **p<0.01; between normal and obese, #p<0.05, ##p<0.01.
enhance lipolysis and upregulate the AC3-cAMP-PKA-HSL pathway. Phosphorylation of HSL at Ser660 is critical for activation of the protein’s intrinsic enzymatic activity.

Therefore, we compared the protein expression levels of hepaticAC3, HSL and p-HSL(S660), and the mRNA expression levels of AC3 and HSL, and cAMP levels and PKA activity in livers of obese and normal control mice with and without liraglutide treatment. We observed that the protein and mRNA expression of hepatic AC3 and the cAMP levels and PKA activity were significantly different in obese mice than in normal control mice, and the protein and mRNA expression of hepatic AC3 and the cAMP levels and PKA activity were increased after liraglutide treatment in obese mice and in normal control mice. The increase in obese mice was more obviously. Liraglutide treatment did not affect HSL protein and mRNA expression, and hepatic HSL expression was not different between obese and normal control mice. Although total HSL expression was not affected, p-HSL(S660) was significantly increased by liraglutide treatment. Human genetic and animal models indicate that AC3 may play an important role in energy homeostasis. Our previous study data provide evidence that supports this concept.

In this present study, AC3 was also upregulated by liraglutide treatment, and the increase in AC3 was paralleled by increases in cAMP, PKA activity and p-HSL(S660). Whether AC3 is the key regulator of this pathway remains to be investigated in future studies.

**Conclusion**

In this study, we found that liraglutide treatment decreased TGs and increased glycerol. Liraglutide treatment upregulated AC3 and p-HSL(S660) levels and cAMP and PKA activities in liver. Thus, we conclude that liraglutide can upregulate the AC3/PKA/HSL pathway and may promote lipolysis.

**Acknowledgment**

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**Author contributions**

All authors contributed to data analysis, drafting or revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.
Disclosure
The authors report no conflicts of interest in this work.

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