Research Article

Erchen Decoction Ameliorates the Metabolic Abnormalities of High-Fat Diet-Fed Rats

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Objective. Brown adipose tissue (BAT) dissipates chemical energy to protect against obesity. In the present study, we aimed to determine the effects of Erchen decoction on the lipolysis and thermogenesis function of BAT in high-fat diet-fed rats.

Methods. Sprague-Dawley rats were randomly divided into four groups, which were fed a control diet (C) or a high-fat diet (HF), and the latter was administered with high and low doses of Erchen decoction by gavage once a day, for 12 weeks. Body weight, the serum lipid profile, serum glucose, and insulin levels of the rats were evaluated. In addition, the phosphorylation and protein and mRNA expression of AMP-activated protein kinase (AMPK), adipose triglyceride lipase (ATGL), peroxisome proliferator-activated receptor γ coactivator-1α (PGC-1α), and uncoupling protein 1 (UCP-1) in BAT were measured by immunoblotting and RT-PCR.

Results. Erchen decoction administration decreased body weight gain and ameliorated the abnormal lipid profile and insulin resistance index of the high-fat diet-fed rats. In addition, the expression of p-AMPK and ATGL in the BAT was significantly increased by Erchen decoction. Erchen decoction also increased the protein and mRNA expression of PGC-1α and UCP-1 in BAT.

Conclusion. Erchen decoction ameliorates the metabolic abnormalities of high-fat diet-fed rats, at least in part via activation of lipolysis and thermogenesis in BAT.

1. Introduction

The prevalence of obesity has reached epidemic proportions, and 600 million people worldwide are estimated to be obese by 2025 [1]. The prevalence of obesity in China has risen rapidly in the past four decades [2]. Obesity is associated with unemployment, social disadvantages, and reduced socioeconomic productivity, thus increasingly creating an economic burden [3]. Furthermore, obesity has been shown to play an important role in the pathophysiology of numerous chronic diseases which include cardiovascular disease (CVD), dyslipidemias, type 2 diabetes (T2D), and cancer [4–6]. The fifth risk for attributable deaths in females was high body mass index (hBMI), while for the males, hBMI accounts for the sixth risk attributed to death in 2019 [7]. Although tremendous progress over the last decade has been made in treating obesity, the development of additional classes of therapeutics and drugs is still needed to be explored to reduce its burden.

Obesity develops when the metabolic energy input of the body exceeds the energy output. Human beings have two kinds of adipose tissue. White adipose tissue (WAT) stores the excess energy tissue in the form of triglycerides in a fat-storing droplet. In addition to WAT, mammals possess brown adipose tissue (BAT), which plays a nearly opposite function [8]. BAT has multilocular lipid droplets and large numbers of mitochondria that contain uncoupling protein-1 (UCP-1). UCP-1 in BAT uncouples the respiratory chain from oxidative phosphorylation. This process allows brown adipocytes to actively oxidize substrates to produce heat [8]. Numerous studies reported that BAT-mediated thermogenesis can protect against obesity by promoting energy...
2.2. Preparation of ECD and LC-MS/MS Analyses. The reduced pressure ECD was stored under 4°C and reprepared once a week. The quality control for ECD concentrated granules has been evaluated by LC-MS/MS determined by Agilent 1290uplc and qtof6550 with a WATERS BEH C18 column (2.1 * 100 mm 1.7μm). The injection volume was 5μL. The electrospray ionization source was set to positive and negative modes.

2.3. Tissue Collection and Biochemical Analysis. The rats were fasted overnight, and trunk blood was then obtained, left at room temperature for 30 min, and then centrifuged at 4,500 × g and 4°C for 15 min. The lipid concentrations and glucose levels of the serum obtained were measured using a Hitachi clinical analyzer. The plasma level of insulin was determined using Rat ELISA kits (Invitrogen Inc). To evaluate the degree of insulin resistance, the homeostasis model assessment (HOMA) was used as an index of IR and calculated by the following formula: insulin (mU/mL) * glucose (mmol/L)/22.5 [18].

2.4. Histological Analysis. Tissue sections (5 mm) were obtained from epididymal visceral adipose and interscapular BAT samples that had been fixed and embedded in paraffin. These were then deparaffinized in xylene, rehydrated, and washed in phosphate-buffered saline, prior to hematoxylin and eosin (HE) staining. For immunohistochemical staining, endogenous peroxidase activity was quenched in hydrogen peroxide for 10–15 min, and the sections were washed four times in PBS buffer. Sections were then incubated with the primary antibodies, at 4°C overnight. And immunostaining was identified using DAB peroxidase substrate solution as the chromogen.

2.5. Real-Time Reverse Transcription-PCR. Real-time PCR was performed as previously described [19]. cDNA was synthesized using a Transcriptor First Strand cDNA Synthesis Kit (Roche, Basel, Switzerland), according to the manufacturer’s instructions. For PGC1-α, the primers were 5′- AACAAATGAGCTGCGAAACA-3′ (forward) and 5′-TGAGGACCGCTAGAAAGTTTG-3′ (reverse), resulting in a 71 bp RT-PCR product. For UCP-1, the primers were 5′-GTGAAGGTCAGAATGCAAGC-3′ (forward) and 5′-AGGGCCCCCTTCATGAGGTC-3′ (reverse), resulting in a 100 bp RT-PCR product. For GAPDH, the primers were 5′-AGACAGCCGCATCTCTTGT-3′ (forward) and 5′-AGACAGCCGCATCTCTTGT-3′ (reverse), resulting in a 100 bp RT-PCR product. For GAPDH, the primers were 5′-AGACAGCCGCATCTCTTGT-3′ (forward) and 5′-
CTTGCCGTGGGTAGAGTCAT-3′ (reverse), resulting in a 207 bp RT-PCR product. RT-PCR reactions were performed using a CFX96™ Real-Time System (Bio-Rad, Hercules, CA, USA) in final volumes of 20 μL, which contained 10 μL of FastStart Universal SYBR Green Master (Rox) (Roche, Basel, Switzerland), 1 μL of each primer (10 μM), 2 μL of cDNA, and 7 μL of PCR-grade water. The RT-PCR products underwent melting point analysis and were quantified using the ΔΔCT method, with GAPDH as the reference gene. The expression in the HF and HF+CL groups was normalized to that of the C group.

### 2.6. Immunoblotting

Immunoblotting was performed as previously described [20]. Briefly, tissue lysates were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and then electrotransferred to membranes. The primary antibodies used were as follows: anti-UCP-1, anti-PGC-1α (PA5-22958; Thermo), anti-AMPK (5831; Cell Signaling Technology), anti-p-AMPK (2535; Cell Signaling Technology), anti-ATGL (2439; Cell Signaling Technology), anti-p-ATGL (135093; Abcam), and anti-glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (5174; Cell Signaling Technology). Membranes were incubated with primary and
secondary antibodies using standard techniques, and the detection of specific protein bands was accomplished using enhanced chemiluminescence. Images of the bands were acquired and analyzed using a quantitative digital imaging system (Quantity One; Bio-Rad), avoiding saturation.

2.7. Statistical Analysis. Data are presented as the mean ± standard deviation (SD) and represent the results of at least three independent experiments. Statistical comparisons were made using Student’s t-test or one-way analysis of variance, followed by Tukey’s test where applicable, to identify significant differences between mean values. \( p < 0.05 \) was considered to represent statistical significance.

3. Results

3.1. LC-MS/MS Analyses of ECD. The total ion chromatogram of ECD is shown in Figure 1. This was matched with the OTCML database. A total of 119 compounds in ECD were identified through chromatogram matching (Supplementary material 1). The concentration of hesperidin in ECD was 91.48 ppm.

3.2. ECD Administration Reduces the Body Mass and Ameliorates the Serum Lipid Abnormalities and Insulin Resistance of High-Fat Diet-Fed Rats. As expected, the body masses of the HF rats were significantly higher than those of the CON rats, but ECD administration reduced the body mass of the HF rats (Figure 2(a)). The HF rats had significantly higher total serum cholesterol, triglyceride, and low-density lipoprotein- (LDL-) cholesterol concentrations than the CON group, whereas the high-density lipoprotein- (HDL-) cholesterol concentration in the HF rats was significantly lower. However, ECD administration improved the lipid profile of the HF rats (Figure 2(b)). The HF rats had significantly higher serum glucose, and insulin levels and ECD administration reduced serum glucose and insulin levels (Figures 2(c) and 2(d)). Consistently, ECD administration reduced HOMA-IR in HF rats (Figure 2(e)).

3.3. ECD Administration Reduces the Epididymal Fat Depot Masses of HFD-Fed Rats. HE staining of tissue sections showed that HF rats had larger lipid droplets in their interscapular brown and epididymal adipocytes (Figure 3(a)). Concurrent ECD administration reduced the weight of epididymal adipose tissue fed with high-fat diet (Figure 3(b)).
3.4. ECD Administration Increases the Expression of UCP-1 and PGC-1α in the Interscapular Brown Adipose Tissue of HF Rats. ECD administration increased UCP-1 and PGC-1α mRNA expression levels in the interscapular brown adipose tissue of HF rats (Figure 4(a)). Consistent with this, immunoblotting and immunohistochemistry assay showed that the protein expression levels of UCP-1 and PGC-1α in the interscapular brown adipose tissue of HF rats were significantly lower than those of CON rats, and ECD administration eliminated these differences (Figures 4(b) and 4(c)).

3.5. ECD Treatment Increases p-AMPK and p-ATGL Expression in the Interscapular Brown Adipose Tissue of HF Rats. Immunoblotting showed that the protein expression levels of p-AMPK and p-ATGL in their interscapular brown adipose tissue of HF rats were significantly lower than those of CON rats, and ECD administration eliminated these differences (Figures 5(a) and 5(b)).

4. Discussion

Obesity is one of the most common metabolic diseases around the world partially due to the consumption of fast food containing high levels of fat and glucose [21, 22]. In the present study, we first found that high-fat diet for 12 weeks increased body weight and serum lipid profiles and exacerbated glucose tolerance and insulin sensitivity. In the theory of TCM, high-fat diet leads to Pi deficiency and increased dampness in the body. We also found that the epididymal visceral adipose tissue had larger lipid droplets which is a sign of increased dampness in the body after high-fat diet feed. ECD is clinically used in China to treat obesity and hyperlipidemia by its dampness-reducing effects [12, 13]. ECD has been also demonstrated to prevent high-fat diet-induced metabolic disorders [23–25]. Herein, we consistently found that ECD treatment was able to reduce body weight and serum lipid profiles in high-fat diet-fed rats. ECD treatment also significantly reduced the
epididymal visceral adipose tissue weight and the diameter of the lipid droplets in the epididymal visceral adipose tissue. These results indicated that ECD was able to remove the phlegm in the form of triglyceride in adipose tissue.

It was previously thought that BAT was regressed in adulthood [26]. However, a large quantity of bioactive brown/beige adipocytes was recently found in the neck and shoulder regions of adults by 18F-FDG-PET/CT detection [27, 28]. This has sparked a resurgence in the targeting of BAT to treat obesity and type 2 diabetes. Many traditional Chinese medicine drugs have been shown to activate BAT to prevent metabolic disorders in high-fat diet-fed mice and rats [29–31]. According to the theory of TCM, high-fat diet-induced Pi deficiency leads to skeletal muscle dysfunction, while ECD increases Pi and has beneficial metabolic effects on high-fat diet-fed mice partially due to the upregulation of the protein levels of lipoprotein lipase in skeletal muscle. BAT and skeletal muscle both originated from myogenic factor 5 positive cells [16, 17]. Thus, we hypothesized that ECD might activate the thermogenesis function of BAT and found ECD was able to increase the protein and mRNA levels of UCP-1 in BAT [33, 34]. Next, we consistently found that ECD also upregulated the protein and mRNA levels of PGC-1α in BAT in high-fat diet-fed rats.

Lipolysis is a fundamental biochemical process which occurs essentially in WAT and BAT. The catabolism of triglyceride molecules is through the activation of the major TAG hydrolases, such as adipose triglyceride lipase (ATGL) and hormone-sensitive lipase (HSL) [35]. Lipolytic activity of ATGL affects BAT function, and adipocyte-specific ATGL knockout (KO) mice show impaired thermogenic functions in response [36]. Furthermore, activation of ATGL in BAT was found to improve metabolic parameters in high-fat diet-fed mice [37]. Thus, we next detected the expression levels of p-ATGL in BAT and found that ECD was able to activate ATGL in BAT in high-fat diet-fed rats. AMP-activated protein kinase (AMPK) is a widely expressed multisubstrate serine/threonine kinase and a well-known sensor of the intracellular energy state that responds to metabolic stresses and other regulatory signals [38]. By sensing the cellular energy state, AMPK activation induces FA oxidation partially by phosphorylating ATGL to increase its TAG hydrolase activity [39]. Our results showed that ECD upregulated the expression levels of p-AMPK in BAT in high-fat diet-fed rats. These indicated that ECD might activate lipolysis in BAT partially through the AMPK-ATGL pathway.

Figure 5: ECD activated AMPK/ATGL signaling. (a) ECD increased p-AMPK protein expression levels in interscapular brown adipose tissue in HFD-fed rats. p-AMPK densitometry values were adjusted to AMPK intensity, then normalized to expression from the control sample. Expression in the CON group was normalized to 1, n = 3. (b) ECD increased p-ATGL protein expression levels in interscapular brown adipose tissue in HFD-fed rats. p-ATGL densitometry values were adjusted to ATGL intensity, then normalized to expression from the control sample. Expression in the CON group was normalized to 1, n = 3. ECD: Erchen decoction; CON: control group; HF: high-fat diet group; ECDL: high-fat diet and low dose of ECD treatment group; ECDH: high-fat diet and high dose of ECD treatment group. *p < 0.05 versus CON; †p < 0.05 versus HF.
5. Conclusions

In conclusion, the present study demonstrated that Erchen decoction improves metabolic parameters of high-fat diet-fed rats via activating lipolysis and thermogenesis in brown adipose tissue.

Data Availability

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

Ethical Approval

All applicable international, national, and institutional guidelines for the care and use of animals were followed. The animal experiments were approved by the Animal Research Center of Guangzhou University of Chinese Medicine. The ethical approval certificate number is 2016139.

Conflicts of Interest

The authors declare that they have no competing interests.

Authors’ Contributions

Conception and design were performed by Jin-Wen Xu, Li Guan, Hai-Mei Liu, and Ya-Xing Zhang; experiments were performed by Ya Cheng, Lu-Yao Xu, Ning Zhang, and Jun-Hua Yang; manuscript writing was performed by Jin-Wen Xu and Run-Mei Li; final approval of manuscript was performed by all authors. Ya Cheng and Lu-Yao Xu contributed equally to this work and are the first authors.

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Supplementary Materials

Supplementary material 1: a total of 119 compounds in ECD were identified through chromatogram matching. (Supplementary Materials)

References

[1] World Health Organization, Noncommunicable Diseases, World health Organization, 2017, http://www.who.int/mediacentre/factsheets/fs355/en/.

[2] Q. Zeng, N. Li, X. F. Pan, L. Chen, and A. Pan, "Clinical management and treatment of obesity in China," The Lancet Diabetes and Endocrinology, vol. 9, no. 6, pp. 393–405, 2021.

[3] M. Blüher, "Obesity: global epidemiology and pathogenesis," Nature Reviews. Endocrinology, vol. 15, no. 5, pp. 288–298, 2019.

[4] S. Yusuf, S. Hawken, S. Ounpuu et al., "Obesity and the risk of myocardial infarction in 27 000 participants from 52 countries: a case-control study," Lancet, vol. 366, no. 9497, pp. 1640–1649, 2005.

[5] V. S. Malik and F. B. Hu, "The role of sugar-sweetened beverages in the global epidemics of obesity and chronic diseases," Nature Reviews Endocrinology, vol. 18, no. 4, pp. 205–218, 2022.

[6] M. Hancková and T. Betáková, "Pandemics of the 21st century: the risk factor for obese people," Viruses, vol. 14, no. 1, p. 25, 2022.

[7] C. J. Murray, A. Y. Aravkin, P. Zheng et al., "Global burden of 87 risk factors in 204 countries and territories, 1990–2019: a systematic analysis for the Global Burden of Disease Study 2019," Lancet, vol. 396, no. 10258, pp. 1223–1249, 2020.

[8] S. N. Shapiro and P. Seale, "Transcriptional control of brown and beige fat development and function," Obesity (Silver Spring), vol. 27, no. 1, pp. 13–21, 2019.

[9] L. Ma, Z. Zhao, X. Guo et al., "Tanshinone IIA and its derivative activate thermogenesis in adipocytes and induce "beiging" of white adipose tissue," Molecular and Cellular Endocrinology, vol. 544, article 111557, 2022.

[10] Q. Chen, D. Wang, Y. Gu, Z. Jiang, and Z. Zhou, "Tangeretin prevents obesity by modulating systemic inflammation, fat browning, and gut microbiota in high-fat diet-induced obese C57BL/6 mice," The Journal of Nutritional Biochemistry, vol. 101, article 108943, 2022.

[11] C. T. Herz, O. C. Kulterer, M. Prager et al., "Active brown adipose tissue is associated with a healthier metabolic phenotype in obesity," Diabetes, vol. 71, no. 1, pp. 93–103, 2022.

[12] H. Liu, J. Xu, H. Li, L. Zhang, and P. Xu, "Network pharmacology-based investigation to explore the effect and mechanism of Erchen decoction against the nonalcoholic fatty liver disease," The Anatomical Record, vol. 304, no. 11, pp. 2605–2619, 2021.

[13] T. Zhao, L. Zhan, W. Zhou et al., "The effects of Erchen decoction on gut microbiota and lipid metabolism disorders in Zucker diabetic fatty rats," Frontiers in Pharmacology, vol. 12, article 647529, 2021.

[14] Q. W. Zhang, G. Y. Li, Y. P. Ren, and Y. F. Gao, "Effect of two Pi deficiency syndrome models on the configuration and function of the skeletal muscle in mice," Zhongguo Zhong Xi Yi Jie He Za Zhi, vol. 35, no. 1, pp. 71–75, 2015.

[15] H. Xia, B. Zhang, D. Yang et al., "Yi-Qi-Jian-Pi-Xiao-Yu-Xie-Zhuo formula improves muscle atrophy via modulating the IGF-1/PI3K/Akt signaling pathway in 5/6 nephrectomized rats," Frontiers in Pharmacology, vol. 12, no. 12, article 624303, 2021.

[16] P. Seale, B. Bjork, W. Yang et al., "PRDM16 controls a brown fat/skeletal muscle switch," Nature, vol. 454, no. 7207, pp. 961–967, 2008.

[17] J. Sanchez-Gurmaches, C.-M. Hung, C. A. Sparks, Y. Tang, H. Li, and D. A. Guertin, ""_PTEN_ loss in the Myf5 lineage redistributes body fat and reveals subsets of white adipocytes that arise from Myf5 precursors," Cell Metabolism, vol. 16, no. 3, pp. 348–362, 2012.

[18] A. M. Yaryari, M. Mazdeh, M. Mohammadi, A. R. Haghj, M. Ghiasian, and M. Mehrpooya, "Evaluation of serum levels of asprosin and other metabolic profiles in patients with idiopathic tonic-clonic generalized epilepsy on treatment with
valproic acid,” European Journal of Clinical Pharmacology, vol. 78, no. 3, pp. 393–403, 2022.

[19] S. Q. Chen, L. N. Ding, N. X. Zeng et al., “Icarin induces irisin/FNDC5 expression in C2C12 cells via the AMPK pathway,” Biomedicine & Pharmacotherapy, vol. 115, article 108930, 2019.

[20] L. N. Ding, Y. Cheng, L. Y. Xu et al., “The β3 adrenergic receptor agonist CL316243 ameliorates the metabolic abnormalities of high-fat diet-fed rats by activating AMPK/PGC-1α signaling in skeletal muscle,” Diabetes, metabolic syndrome and obesity: targets and therapy, vol. Volume 14, pp. 1233–1241, 2021.

[21] J. Lu, Y. Bi, and G. Ning, “Curbing the obesity epidemic in China,” The Lancet Diabetes and Endocrinology, vol. 4, no. 6, pp. 470–471, 2016.

[22] S. Vandejijvere, C. C. Chow, K. D. Hall, E. Umali, and B. A. Swinburn, “Increased food energy supply as a major driver of the obesity epidemic: a global analysis,” Bulletin of the World Health Organization, vol. 93, no. 7, pp. 446–456, 2015.

[23] M. Zhang, Y. Shao, B. Gao et al., “Erchen decoction mitigates lipid metabolism disorder by the regulation of PPARγ and LPL gene in a high-fat diet C57BL/6 mice model,” Evidence-based Complementary and Alternative Medicine, vol. 2020, Article ID 9102475, 8 pages, 2020.

[24] B. Z. Gao, J. C. Chen, L. H. Liao, J. Q. Xu, X. F. Lin, and S. S. Ding, “Erchen decoction prevents high-fat diet induced metabolic disorders in C57BL/6 mice,” Evidence-based Complementary and Alternative Medicine, vol. 2015, Article ID 501272, 9 pages, 2015.

[25] S. Ding, J. Kang, L. Tong, Y. Lin, L. Liao, and B. Gao, “Erchen decoction ameliorates lipid metabolism by the regulation of the protein CAV-1 and the receptors VLDLR, LDLR, ABCA1, and SRB1 in a high-fat diet rat model,” Evidence-based Complementary and Alternative Medicine, vol. 2018, Article ID 5309490, 12 pages, 2018.

[26] J. Yang, H. Zhang, K. Parhat et al., “Molecular imaging of brown adipose tissue mass,” International Journal of Molecular Sciences, vol. 22, no. 17, p. 9436, 2021.

[27] A. M. Cypess, S. Lehman, G. Williams et al., “Identification and importance of brown adipose tissue in adult humans,” The New England Journal of Medicine, vol. 360, no. 15, pp. 1509–1517, 2009.

[28] K. A. Virtanen, M. E. Lidell, J. Orava et al., “Functional brown adipose tissue in healthy adults,” The New England Journal of Medicine, vol. 360, no. 15, pp. 1518–1525, 2009.

[29] H. Zhang, Y. Hao, C. Wei et al., “Chinese medicine Jinlida granules improve high-fat-diet induced metabolic disorders via activation of brown adipose tissue in mice,” Biomedicine & Pharmacotherapy, vol. 114, article 108781, 2019.

[30] L. Cheng, S. Zhang, F. Shang et al., “Emodin improves glucose and lipid metabolism disorders in obese mice via activating brown adipose tissue and inducing browning of white adipose tissue,” Frontiers in Endocrinology, vol. 12, article 618037, 2021.

[31] B. Sun, M. Hayashi, M. Kudo et al., “Madecassoside inhibits body weight gain via modulating SIRT1-AMPK signaling pathway and activating genes related to thermogenesis,” Frontiers in Endocrinology, vol. 12, article 627950, 2021.

[32] P. Puigserver and B. M. Spiegelman, “Peroxisome proliferator-activated receptor-γ coactivator 1α (PGC-1α): transcriptional coactivator and metabolic regulator,” Endocrine Reviews, vol. 24, no. 1, pp. 78–90, 2003.