



OPEN ACCESS

EDITED BY

Danar Praseptianga,
Sebelas Maret University, Indonesia

REVIEWED BY

Qiaozhi Zhang,
Zhejiang Gongshang University, China
Maria Eugenia Jaramillo-Flores,
Instituto Politécnico Nacional (IPN),
Mexico

*CORRESPONDENCE

Zichao Li
✉ zichaoli@qdu.edu.cn

†These authors have contributed
equally to this work and share first
authorship

SPECIALTY SECTION

This article was submitted to
Nutrition and Food Science
Technology,
a section of the journal
Frontiers in Nutrition

RECEIVED 25 October 2022

ACCEPTED 16 December 2022

PUBLISHED 09 January 2023

CITATION

Fu K, Gao X, Hua P, Huang Y, Dong R,
Wang M, Li Q and Li Z (2023)
Anti-obesity effect of *Angelica keiskei*
Jiaosu prepared by yeast
fermentation on high-fat diet-fed
mice.
Front. Nutr. 9:1079784.
doi: 10.3389/fnut.2022.1079784

COPYRIGHT

© 2023 Fu, Gao, Hua, Huang, Dong,
Wang, Li and Li. This is an open-access
article distributed under the terms of
the [Creative Commons Attribution
License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/). The use, distribution
or reproduction in other forums is
permitted, provided the original
author(s) and the copyright owner(s)
are credited and that the original
publication in this journal is cited, in
accordance with accepted academic
practice. No use, distribution or
reproduction is permitted which does
not comply with these terms.

Anti-obesity effect of *Angelica keiskei* Jiaosu prepared by yeast fermentation on high-fat diet-fed mice

Kunli Fu^{1†}, Xiang Gao^{1,2†}, Puyue Hua¹, Yuedi Huang¹,
Ruitao Dong³, Mingji Wang⁴, Qun Li^{3,4} and Zichao Li^{1,4*}

¹College of Life Sciences, Institute of Biomedical Engineering, Qingdao University, Qingdao, China, ²Anqiu Huatao Food Co., Ltd., Weifang, China, ³College of Chemistry and Chemical Engineering, Qingdao University, Qingdao, China, ⁴Joint Institute of *Angelica keiskei* Health Industry Technology, Qingdao Balanson Biotech Co., Ltd., Qingdao, China

In this study, an *Angelica keiskei* (*A. keiskei*) Jiaosu (FAK) was prepared by yeast fermentation to investigate its anti-obesity effect on high-fat diet (HFD)-fed mice. 70 SPF grade male C57BL/6J mice were randomly divided into 7 groups ($n = 10$): blank control group (N), high-fat model group (M), positive control group (Orl), unfermented control group (NF), high-dose intervention group (FH), medium-dose intervention group (FM), and low-dose intervention group (FL). The results showed that FAK intervention significantly reduced the body weight, Lee's index and liver index of HFD-fed mice ($P < 0.05$). Compared with M group, the serum levels of triglyceride (TG), total cholesterol (TC), leptin and glucose (GLU) in FH group were remarkably decreased and that of interleukin-27 (IL-27) were increased ($P < 0.05$). The levels of TG, and TC in the liver of mice were also markedly decreased in the FH group ($P < 0.05$). HE staining results showed that the liver cells in the three intervention groups had less degeneration and fatty vacuoles in the cytoplasm, and the liver cords were orderly arranged compared with that of M group. Furthermore, FAK significantly inhibited epididymal adipose tissue cell expansion induced by HFD. FAK up-regulated the protein expression levels of p-AMPK and PPAR α to promote lipolysis and down-regulated the expression of PPAR γ to reduce lipid synthesis ($P < 0.05$). Additionally, the results of gut microbiota showed that after the intervention, a decrease trend of F/B value and *Deferribacterota* was noticed in the FH group compared with M group. At the genus level, FAK intervention significantly increased that of *Ileobacterium* compared to the M group ($p < 0.05$). A rising trend of norank_f_Muribaculaceae, *Lactobacillus*, and *Bifidobacterium* were also observed in the HF group. Conclusively, these findings demonstrated that FAK intervention can effectively improve obesity

in mice caused by HFD and the potential mechanisms was related to the regulation of serum levels of leptin and IL-27, lipogenesis and lipolysis in adipose tissue and gut microbiota composition.

KEYWORDS

Angelica keiskei, Jiaosu, total flavonoids, anti-obesity, mechanism

1. Introduction

Over the past few decades, the global prevalence of obesity has rapidly increased, resulting in a worldwide issue (1). Obesity is a multifactorial chronic disease with multiple complications, which is associated with a variety of diseases, including metabolic diseases (e.g., type 2 diabetes), hypertension, non-alcoholic fatty liver disease, musculoskeletal diseases, cardiovascular diseases, psychiatric diseases, and more seriously, certain types of cancer. It seriously threatens human life and health, and damages the quality of life (2). At present, the change of the global dietary structure and the lifestyle of more calorie intake and less physical activity have become the main driving factors for the occurrence of obesity. Treatment for obesity now focuses on lifestyle changes (low-calorie diets and increased physical activity), drug therapy, and bariatric surgery (3). However, due to the side effects of drugs and the fact that some individuals are not suitable for surgery, dietary intervention remains the most ideal way to curb obesity.

The word “Jiaosu” was originated in Japan and first became popular in European and American countries. Until the last decade, it has begun to be recognized by the Chinese and thrived in the local market. In December 2018, the Chinese Ministry of Industry and Information Technology issued the “Guidelines for Jiaosu Products Classification” (QB/T 5324-2018), in which Jiaosu was officially defined as: a product containing specific biologically active ingredients obtained by microbial fermentation, using animals, plants, fungi, etc. as raw materials, with or without added auxiliary materials. A large number of experimental studies have proved that Jiaosu cannot only retain the original nutrients in the raw food ingredients, but also its probiotics can hydrolyze macromolecular substances, produce enzymes and increase the content of phenolic compounds, vitamins, minerals and other metabolites, thereby greatly improving bioavailability (4). Meanwhile, it has the advantages of sensory improvement, prolonging the shelf life and etc. (5). Increasing evidence indicated that Jiaosu can effectively lower body weights and alleviate obesity. Verón et al. (6) reported that the cactus pear (*Opuntia ficus-indica*) fruit juice fermented by autochthonous *Lactobacillus plantarum* S-811 can significantly reduce the body weights in the high-fat diet (HFD) mice, suggesting it had the potential to prevent and improve obesity and related diseases. Pan et al. examined the

anti-obesity effect of fermented lemon peel on high-fat diet induced (HFD-induced) obese mice, and the results showed that the supernatant of fermented lemon peel significantly inhibited body weight gain and improved liver and epididymal adipose tissue accumulation in HFD mice lesions (7). Cho et al. found that fermented green tea extract suppressed adipogenesis and lipogenesis in adipocytes by enhancing mRNA expression of fatty acid oxidation-related genes (8).

Angelica keiskei (*A. keiskei*) is a perennial cold-hardy herb of the *Umbelliferae* family. It is native to the large islands of Hachijo and Izu on the Pacific coast of Japan (9). To date, it has been cultivated in Shandong, Yunnan, Guangxi, Jiangsu, and Hainan provinces etc. in China (10), and was officially recognized as a new food ingredient by the National Health Commission in 2019. *A. keiskei* is rich in chalcone, flavonoids, coumarin, and other bioactive substances (11), and shows a variety of potential biological activities, such as antitumor, antioxidant, hypoglycemic, hypotensive, liver protection, antibacterial, and antiviral, etc. (12–14). The aerial parts (stems and leaves) have been eaten for centuries as a healthy vegetable, and the whole plant can be used medicinally. Collectively, *A. keiskei* has been recognized as a valuable, healthy and green functional food source.

To our knowledge, there is no scientific report to prepare Jiaosu with *A. keiskei* as raw material and to study its anti-obesity effect *in vivo*. Herein, the *A. keiskei* Jiaosu was firstly prepared via optimized yeast fermentation, and particularly, its effects on weight loss in HFD induced obese mice were studied. Furthermore, the underlying mechanism involves the regulation of lipid synthesis and lipolysis and intestinal microbiota were investigated. The findings will provide research basis for the in-depth functional food development and high-value utilization of *A. keiskei*.

2. Materials and methods

2.1. Food ingredients and reagents

A. keiskei freeze-dried powder was provided by Shandong *A. keiskei* Biotech Co., Ltd. (Rizhao, China). Orlistat was purchased from Zeen Biotechnology Co., Ltd. (Changsha, China). XTHF60-1 high-fat feed was purchased from

Jiangsu Cooperation Pharmaceutical Bioengineering Co., Ltd. (Nanjing, China).

2.2. The preparation of *A. keiskei* Jiaosu

The process parameters for preparation of *A. keiskei* Jiaosu were optimized and fixed through a series of pretests. Briefly, highly active dry yeast (*Saccharomyces cerevisiae*) and distilled water were mixed and dissolved at a ratio of 1:10 (g/mL), and activated at 37°C for 30 min. Subsequently, *A. keiskei* freeze-dried powder of 17.5 g was added to 200 mL sterile water and mixed well. Then 1.5% (v/v) activated yeast liquid was inoculated into the *A. keiskei* fermentation broth, and fermented at 37°C for 96 h. After that, the supernatant was centrifuged and freeze-dried for 24 h to obtain *A. Keiskei* Jiaosu (FAK) powder, dissolved in dosage and administered intragastrically. In addition, we also set a unfermented *A. keiskei* group, in which the *A. keiskei* freeze-dried powder was not inoculated with yeast, while the rest were the same to the processes of preparing FAK.

2.3. Determination of chemical constituents of FAK

The total flavonoid contents of FAK were determined by NaNO₂-AL(NO₃)₃-NaOH chromogenic method as reported (13). The levels of γ -aminobutyric acid was detected by a colorimetric method (15). The contents of free amino acids was determined by ninhydrin colorimetric method. Total protein was measured by the biuret method. Total phenolic content was determined according to the Folin-Ciocalteu colorimetric method (16). The pH value was measured directly using an acidity meter. Soluble solids were measured using a hand-held food sugar meter.

2.4. Protocols of animal study

All animal experimental protocols and procedures were approved by the Ethics Committee of Qingdao University, and the experimental unit license number was SYXK (LU) 20200009. Seventy specific pathogen free male C57BL/6J mice [License number: SCXK (Beijing) 2019-0010] aged 6–8 weeks were purchased from SPF (Beijing) Biotechnology Co., Ltd. After a 1 week adaptively period, the mice were weighted and 10 mice were randomly selected as blank control group (N), which were kept feeding with common chow (main composition: soybean meal, fish meal, beer yeast powder, vegetable oil, bran, corn, wheat, vitamins, and minerals, and etc.) during the experimental period. The rest of the mice were fed with a HFD (32.5% basal diet, 28% lard, 8% sucrose, 10.8% whole milk powder, 13.5% casein, 3% premix for experimental animals,

and 2% microcrystalline cellulose, 1.8% calcium bicarbonate, 0.4% stone powder) for 8 week for model building. Afterward, the obese model mice were randomly divided into 6 groups according to body weight ($n = 10$): the high-fat model group (M), given 10 mL/kg/d of normal saline; the positive control group (Orl), given 30 mg/kg/d orlistat; unfermented control group (NF), given 400 mg/kg/d unfermented *A. keiskei* freeze-dried powder; the high-dose intervention group (FH), given 400 mg/kg/d FAK; the middle-dose intervention group (FM), given 200 mg/kg/d FAK; the low-dose intervention group (FL), given 100 mg/kg/d FAK. The treatment doses were referenced to previous report (17). The FAK dose of 400 mg/kg/d (400 mg of FAK powder was obtained from 20 mL of Jiaosu solution after freeze-drying) was equivalent to consumption of 113.82 mL Jiaosu solution/day by a 70 kg human according to the K_m factor ratio of 3 and 37 for mice (30 g) and humans (70 kg), respectively. The gavage intervention was performed daily at a fixed time for 8 weeks. The body weights of mice were measured once a week. All the mice were free access to drinking and eating, and kept in the Animal Center of the Medical Department of Qingdao University, under a standard condition.

2.5. Blood and tissue samples acquired

Lastly, the mice in each group were fasted for 8 h, weighed, and their feces were collected and stored in a refrigerator at -80°C . After the mice were sacrificed, the blood was collected and centrifuged at 3,500 r/min for 15 min to obtain the serum. The adipose tissue of the liver and epididymis were removed, washed and weighed. After that, a portion of the liver tissues was fixed in a 4% paraformaldehyde solution for pathological analysis, and the rest of the liver tissues were immediately frozen with liquid nitrogen and stored at -80°C until analysis. The epididymal adipose tissues were fixed with a 10% formalin solution.

2.6. Determination of Lee's index and liver index

The Lee's index and liver index were calculated according to the following equation (18).

$$\text{Lee's index} = \frac{\sqrt[3]{\text{Body weight (g)} * 10^3}}{\text{Mouse length(cm)}}$$

$$\text{Liver index} = \frac{\text{Mice liver wet weight(g)}}{\text{Body weight(g)}} \times 100\%$$

2.7. Biochemical analysis

The serum levels of triglyceride (TG), total cholesterol (TC), high-density lipoprotein cholesterol (HDL-c), low-density lipoprotein cholesterol (LDL-c), aspartate aminotransferase (AST), alanine aminotransferase (ALT), and glucose (GLU) were analyzed by an automatic biochemical analyzer (Chemray 800, Redu life science Co., Ltd., Shenzhen, China). The levels of leptin and Interleukin-27 (IL-27) in serum were detected via an ELISA kit (EK227-02, Lianke Biotechnology Co., Ltd., Hangzhou, China).

2.8. Histological analysis

Part of liver and epididymal tissues were fixed with 10% neutral formaldehyde, routinely embedded in paraffin, sectioned, deparaffinized, stained with hematoxylin and eosin (HE) or Oil-Red O, and sealed with neutral gum. The morphology was observed by an optical microscope (Eclipse E100, Nikon Inc., Tokyo, Japan).

2.9. Analysis of gut microbiota in mice

The microbial DNA in the feces were extracted and the high-throughput sequencing on the V3–V4 hypervariable regions of the bacterial 16S rRNA was performed on the Illumina MiSeq platform (Illumina, San Diego, USA) at Majorbio Bio-Pharm Technology Co., Ltd. (Shanghai, China). The data were analyzed on the free online platform.¹

2.10. Western blotting analysis

Total protein was extracted from epididymal adipose tissue. The cytosolic protein lysates were prepared by lysing the cells with RIPA buffer containing 1% protease inhibitor and 1% PMSF. After centrifuging at 12,000 rpm and 4°C for 10 min, the protein concentration of the cell lysate was tested using a bicinchoninic acid (BCA) assay kit (Wanleibio Co., Ltd., Shenyang, WLA004, China) before solubilization in the loading buffer. Equal amounts of proteins were separated on SDS-PAGE and subsequently transferred to PVDF membranes. Then, the PVDF membrane was blocked in 5% milk solution before incubated with various primary antibodies (Wanleibio Co., Ltd., Shenyang, China): P-AMPK α 1/2 (WL05103), PPAR α (WL00978), PPAR γ (WL01800), β -actin (WL01372). After that, the PVDF membranes were washed in TBST (5 min \times 4 times) and combined with the secondary antibody goat anti-rabbit IgG-HRP (WLA023). The ECL luminescence solution

¹ <https://cloud.majorbio.com>

TABLE 1 Analysis of chemical ingredients composition of FAK.

Index	Control group	FAK
Gamma-aminobutyric acid content/(g/L)	1.74 \pm 0.003 ^a	8.46 \pm 0.09 ^b
Total flavonoids/(g/L)	0.58 \pm 0.08 ^a	1.40 \pm 0.01 ^b
Total chalcone content/(g/L)	0.03 \pm 0.01 ^a	0.05 \pm 0.01 ^b
Total phenolic content/(g/L)	0.03 \pm 0.02 ^a	0.07 \pm 0.02 ^b
Free amino acid content/(g/L)	0.53 \pm 0.04 ^a	10.96 \pm 0.64 ^b
Protein content/(g/L)	2.15 \pm 0.82 ^a	4.11 \pm 0.12 ^b
pH	3.69 \pm 0.10 ^a	5.44 \pm 0.10 ^b
Soluble solids/%	3.65 \pm 0.21 ^a	3.63 \pm 0.29 ^a

^{a,b}The different superscripts means significant difference statistically ($P < 0.05$).

(WLA003) was utilized for luminescence. Finally, the films were scanned and the intensities of bands were analyzed.

2.11. Statistical analyses

The experimental data were presented as mean \pm standard deviation (SD) and the statistical analysis were performed by the Origin 2019 software. Data variability between groups was analyzed by the one-way analysis of variance (ANOVA). A p -value < 0.05 was accepted as statistical significance.

3. Results

3.1. Analysis of chemical ingredients composition of FAK

The optimum fermentation process for FAK was obtained via a series of preliminary experiments. Table 1 displays the chemical composition of both FAK and *A. keiskei* FD powder without fermentation (control group). As seen, comparatively, the content of free amino acids, γ -aminobutyric acid (GABA) and total flavonoids in FAK were raised by 20.68, 4.86, and 2.41 times via yeast fermentation, respectively. Moreover, the content of total chalcone, total phenolic content, protein, and pH value were all markedly increased ($P < 0.05$), while no notable change was observed in the soluble solids, indicating that the yeast fermentation process effectively elevated the active ingredient contents of *A. keiskei*, thus further improved its healthy properties for application.

3.2. Growth parameters of the mice

The body weight change, final body weight, Lee's index and liver index of mice were recorded as Figures 1A–D. All mice showed similar body weight at the start of the experiment. After

8 weeks feeding, the final body weight of the mice in the M group was markedly increased by 43.35% compared with the N group ($P < 0.05$), along with increased Lee's index and liver index ($P < 0.05$). These data suggest that obesity was successfully induced in mice by HFD feeding. Compared with the M group, the body weight of the mice in the Orl group and the FH group was notably decreased by 20.60 and 7.57%, respectively ($P < 0.05$). A dose-effect relationship was observed among the FH, FM, and FL group, while there was no prominent difference between the NF group and the M group ($P < 0.05$), indicating that the body weight of mice on a HFD can be effectively controlled by the intervention of FAK. The Lee's index of the FH group was significantly lower than those of the M group as well ($P < 0.05$). Compared with the M group, the liver index of the Orl, FH, FM, and FL group were drastically declined by 2.97, 5.32, 2.01, and 0.52% ($P < 0.05$), respectively. Nevertheless, there was no notable difference between the N group and the FH group ($P > 0.05$).

As shown in **Figures 1E,F**, both the weight of perirenal and epididymal adipose tissues were significantly increased in the M group over the N group ($P < 0.05$). The weight of perirenal adipose tissue was notably decreased by the intervention of FAK and Orl ($P < 0.05$). The epididymal adipose tissue weights in the Orl, FH, and FM group were markedly declined, in comparison with those of the M group. A clear dose-effect relationship can be noticed among the FH, FM, and FL groups. No difference was found between the NF and M group ($P > 0.05$). It can be deduced that the increase in visceral fat in mice induced by HFD was effectively ameliorated by FAK supplement. The morphology of epididymal fat was further examined by HE staining (**Figure 1G**). The size of epididymal adipocytes in the M group was markedly larger than that in the N group. Combined with **Figure 1H** to measure the cells volume, it can be found that the number of cells per unit area was significantly reduced in the M group, compared with the N group ($P < 0.05$). Compared with the M group, the adipocytes per unit area in the Orl group, FH group, FM group, and FL group were increased by 87.21, 184.88, 138.37, and 54.65%, respectively ($P < 0.05$), which proved that HFD-induced enlargement of epididymal adipose tissue cells in mice was also eminently improved by FAK supplement.

3.3. Effects of FAK on the liver of mice

The morphology of liver was observed by HE and Oil Red-O staining, respectively. It can be observed from **Figure 2A** that the hepatocytes of the mice in the N group were relatively intact, with radial arrangement of hepatic cords and abundant cytoplasm, and their nuclei were located in the center of the cells with clear boundaries. The cytoplasm of liver cells of mice in the M group was filled with a large number of small fat vacuoles, the cell morphology was destroyed, the arrangement of

liver cords was disordered, and the cell boundaries were blurred. Compared with the M group, the degeneration of liver cells in the Orl, NF, FH, FM, and FL group were improved, the fat vacuoles in the cytoplasm were reduced, the liver cords were neatly arranged, and the cell morphology was relatively intact, while the improving effect of the NF group was not ideal.

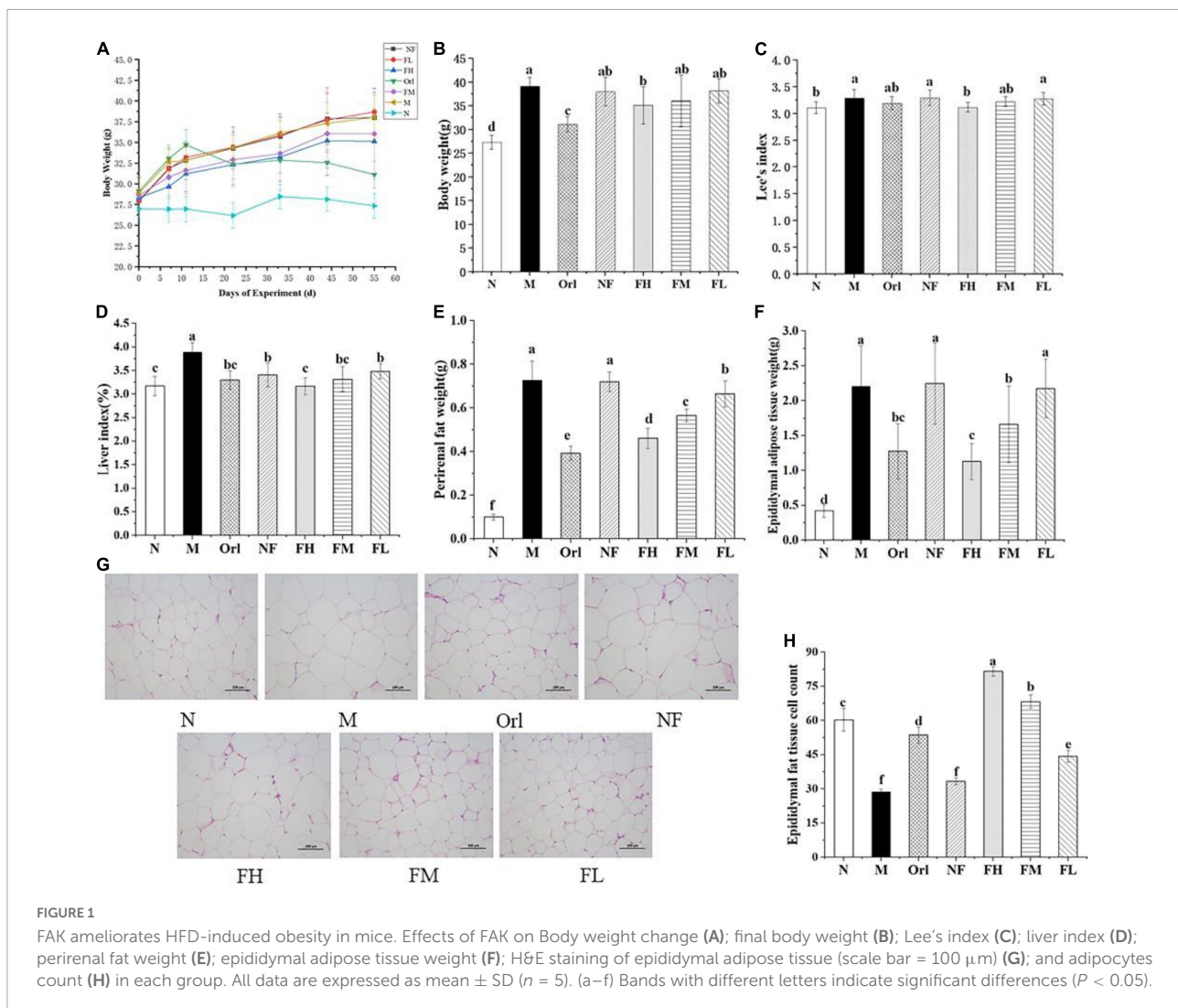
As shown in **Figure 2B**, there were more red areas stained by Oil Red O staining in the M group than the N group. Among all the experimental groups, the intracellular lipid content of the FH group was comparable to that of the N group, of which the improving effect was the best, although the fat accumulations in the liver cells for the Orl, FM, and FL group were also improved.

The levels of TG and TC in liver of mice were also determined, as shown in **Figures 2C,D**. Compared with the N group, the levels of TG and TC in the mice liver for the M group were significantly raised by 248.93 and 48.03% ($P < 0.05$), respectively. Compared with the M group, the TG and TC levels of the other groups were markedly reduced ($P < 0.05$), indicating that the accumulation of hepatic fat caused by HFD in mice was effectively ameliorated by FAK intervention.

3.4. Effects of FAK on serum biochemicals of mice

Figures 3A–D shows the levels for serum biochemicals of mice. The serum TG, TC, and LDL-c of the M group mice were dramatically elevated than those of the N group ($P < 0.05$). Compared with the M group, serum TG levels in the FH and FM group were markedly decreased by 46.51 and 21.55%, respectively ($P < 0.05$). The levels of serum TC in the Orl, NF, FH, FM, and FL group were prominently declined by 14.09, 15.86, 31.92, 22.71, and 19.44%, respectively ($P < 0.05$). The serum HDL-c contents of the M group were increased by 78.45%, compared with those of the N group ($P < 0.05$), while there was no remarkable distinction among the other experimental groups ($P > 0.05$). Moreover, that the serum glucose levels of the mice (**Figure 3E**) in the M group was notably higher than that in the N group ($P < 0.05$). Compared with the M group, the serum glucose levels in the FH group was significantly decreased ($P < 0.05$). There was no conspicuous difference between the other experimental groups and the M group ($P > 0.05$). As seen in **Figures 3E,G**, the serum ALT and AST levels of the M group was markedly elevated by 46.62% ($P < 0.05$) than the N group. Compared with the M group, the Orl and FAK treated groups exhibited significant declined ALT levels ($P < 0.05$). There was no distinction in ALT levels between NF and M groups ($P > 0.05$). The levels of AST in the serum of the Orl, FH, and FM group were eminently declined ($P < 0.05$) by 7.10, 22.79, and 22.45%, respectively.

We also detected serum levels of leptin and IL-27 of the mice. As shown in **Figures 3H,I**, the serum leptin levels of the mice in the M group was 17.59 times that of the N group.



Compared with the M group, the serum leptin levels of the Orl, NF, FH, and FM groups were markedly decreased ($P < 0.05$) by 60.52, 50.28, 80.68, and 69.26%, respectively. The serum levels of IL-27 in each group are shown in **Figure 3F**. Compared with the M group, the serum IL-27 levels of the Orl and FH group were raised by 116.83 and 94.35%, respectively ($P < 0.05$), and a dose-effect relationship was noticed on the FAK treated groups.

3.5. Effects of FAK on the expression of genes related to lipid metabolism in epididymal adipose tissue

To explore the potential mechanisms underlying the anti-obesity effect of FAK, we measured the expression of several proteins related to lipid metabolism in epididymal adipose tissue (**Figures 4A–E**). The results show that, compared with the N group, the relative expressions of p-AMPK/AMPK and PPAR α

in the M group were significantly reduced ($P < 0.05$), and that of PPAR γ was markedly elevated ($P < 0.05$). Compared with the M group, the relative expressions of p-AMPK/AMPK and PPAR α in the FH, FM, and FL groups were remarkably raised ($P < 0.05$), and that of PPAR γ was notably declined ($P < 0.05$). Moreover, a dose-response relationship was found among the FH, FM, and FL groups. Therefore, it can be inferred that FAK improved HFD-induced obesity in mice by regulating the expressions of lipid metabolism-related proteins.

3.6. Effects of FAK on the gut microbiota of mice

The Rank-Abundance curve is mainly used to show the richness of the samples. In the Rank-Abundance curve, the smoother the curve declines, the higher the sample species diversity. As shown in **Figure 5A**, the mice in the N group

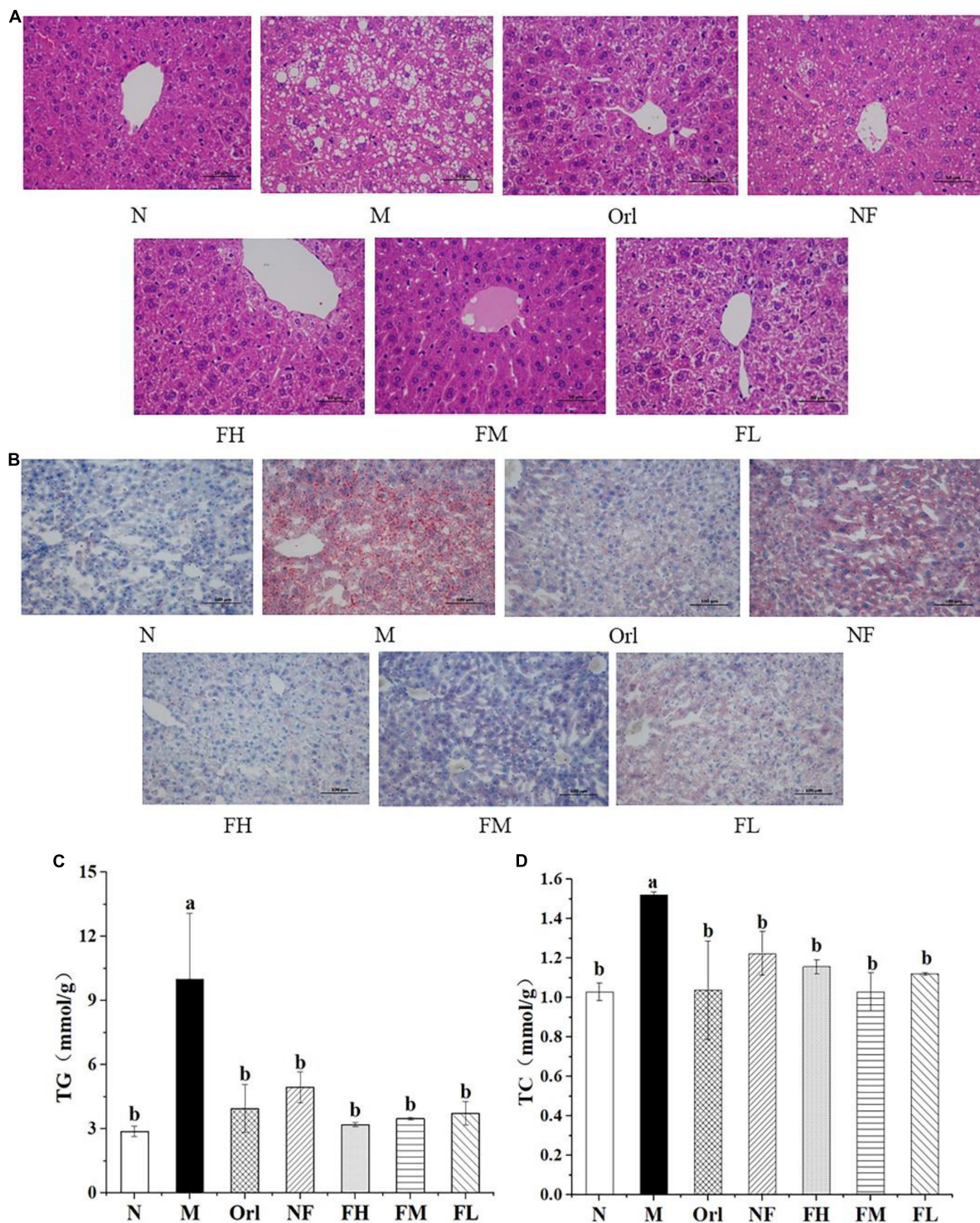
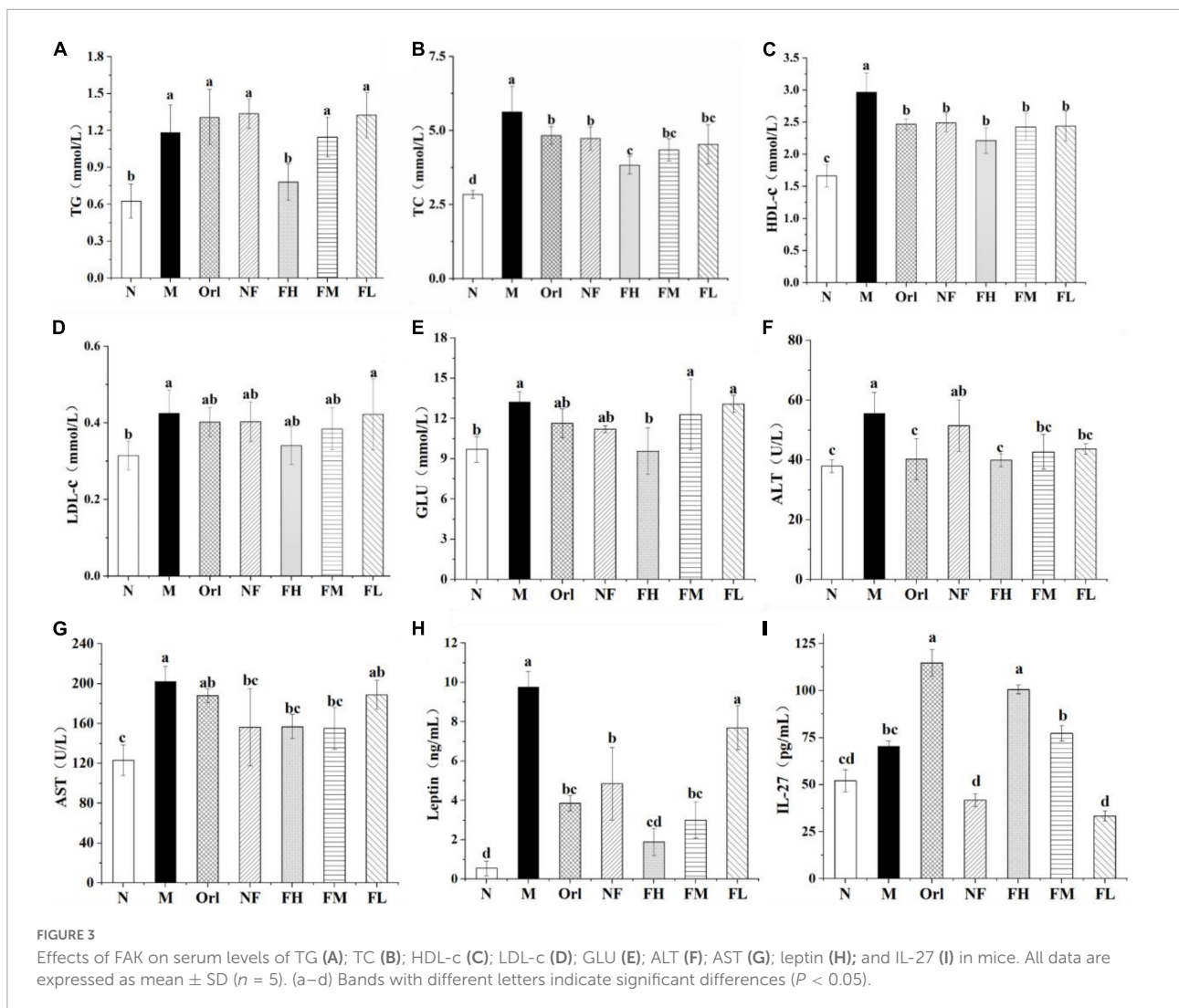


FIGURE 2

Effects of FAK on the mice liver. (A) H&E staining of liver (scale bar = 50 μ m). (B) Oil-Red-O staining of liver (scale bar = 100 μ m). (C) TG levels in liver. (D) TC levels in liver. All data are expressed as mean \pm SD ($n = 5$). (a,b) Bands with different letters indicate significant differences ($P < 0.05$).

possessed the most abundant gut microbiota species, while the Orl group had the least species of bacteria. The species of gut microbiota in the M group was lower than that in the N

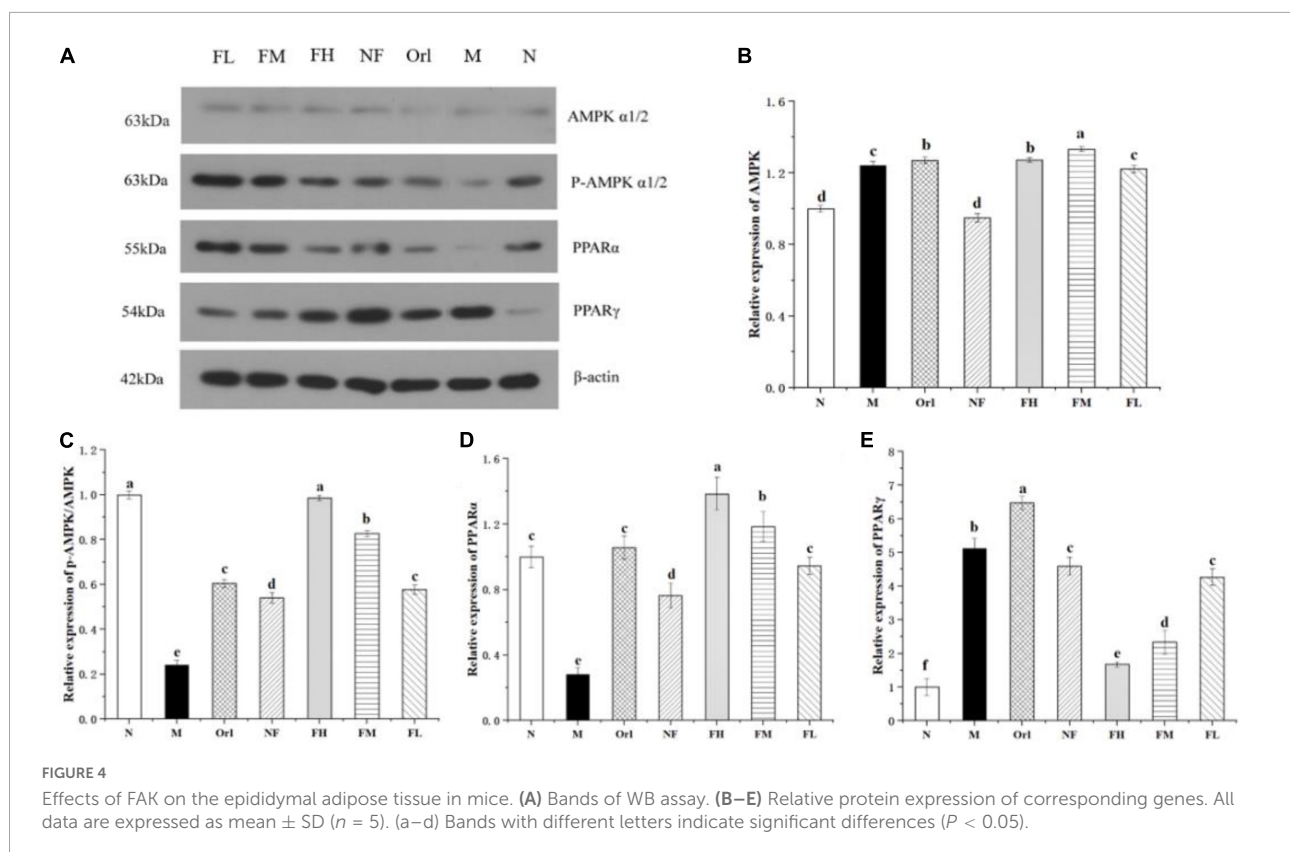
group, indicating that HFD can decrease the abundance of gut microbiota. The results of Alpha diversity analysis are listed in **Figures 5B–F**. Among them, Sobs, Ace, and Chao indices reflect



community richness, while Shannon and Simpson indices show community diversity. The greater the Shannon value, the higher the community diversity. Conversely, the smaller the Simpson Index, the higher the community diversity. Compared with the M group, the Sobs, Shannon, Ace, and Chao values of the FH group were not changed, while these indexes of the Orl group were significantly decreased. FH, NF, and Orl groups exhibited higher Simpson index than the M group ($p < 0.05$). The results indicate that FAK intervention imposed no side effects on the richness and abundance of gut microbiota in the HFD mice. To observe the differences in the overall structure of gut microbiota, principal coordinate analysis (PCoA) was performed. As shown in **Figures 5G,H**, the HFD fed mice were significantly separated from those fed a normal diet, suggesting differences in the number of gut microbes between the M and N groups after the HFD intervention. There was also a trend of isolation between the FH group and M group, indicating that the FAK intervention markedly changed the overall structure of the gut microbiota in

the HFD mice, reflecting that the intervention had a significant effect on the structure of the gut microbiota.

The composition of the gut microbiota at the phylum level was displayed in **Figures 6A,C–E**. It can be observed that the gut microbiota of mice was mainly composed of *Firmicutes*, *Bacteroidota*, and *Actinobacteriota*. The ratio of *Firmicutes/Bacteroidota* (F/B) in the M group was higher than the N group. FH and Orl administration decreased the F/B value, compared with the M group. While, NF group showed a higher F/B value than the M group. HFD feeding led to an elevated trend of *Actinobacteriota*, *Desulfobacterota*, and *Deferribacterota* than the N group, while no significant difference was observed. FAK intervention significantly increased the relative levels of *Actinobacteriota* ($p < 0.05$) and led to a notable declining trend of *Deferribacterota*, compared to the M group. Orl group displayed higher relative abundance of *Actinobacteriota* than the M group ($p < 0.05$). The relative abundance of gut microbiota at the genus levels

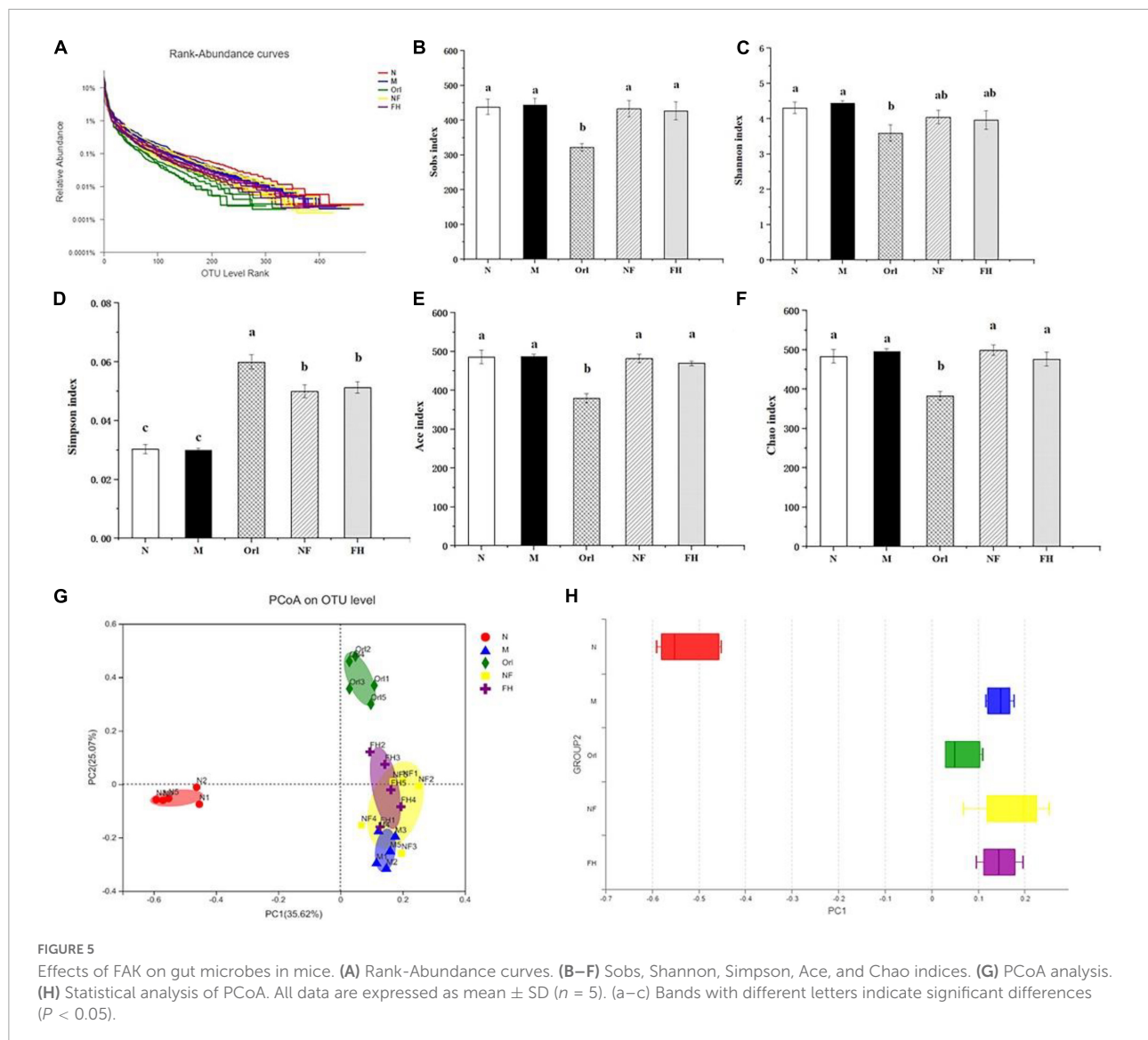


are shown in **Figures 6B,E,G**. As seen, the relative abundance of *norank_f_Muribaculaceae* in the M group was markedly decreased ($p < 0.05$), while *unclassified_f_Lachnospiraceae* and *Ileobacterium* were increased notably, compared with the N group ($p < 0.05$). A rising trend of *Lactobacillus* and decrease of *Bifidobacterium* were also noticed, compared to the N group. FAK intervention remarkably raised that of *Ileobacterium*, compared to the M group ($p < 0.05$). Moreover, decline of *unclassified_f_Lachnospiraceae* and increase of *norank_f_Muribaculaceae*, *Lactobacillus* and *Bifidobacterium* were observed in the HF group as well. Meanwhile, Orl treatment enhanced the relative levels of *norank_f_Muribaculaceae*, *Ileobacterium*, *Lactobacillus*, and decreased that of *unclassified_f_Lachnospiraceae*, compared to the M group ($p < 0.05$). However, there was a prominent increase of *Lactobacillus* in the NF group, comparing with the M group. In addition to α and β diversity analysis of the gut microbiota (19), to identify specific bacteria in the M and FH groups, gut microbiota-diverse species were analyzed using linear discriminant analysis (LDA) effect size (LEfSe) based on the non-parametric factorial KW and rank tests (20). As shown in **Figure 7**, by LEfSe analysis (LDA threshold of 2), *c_Bacilli*, *f_Erysipelotrichaceae*, *o_Erysipelotrichales*, and *g_Ileobacterium* were significantly enriched in the FH group compared with those in the M group, while *f_Lachnospiraceae*,

o_Lachnospirales, and *c_Clostridia* were enriched in the M group.

4. Discussion

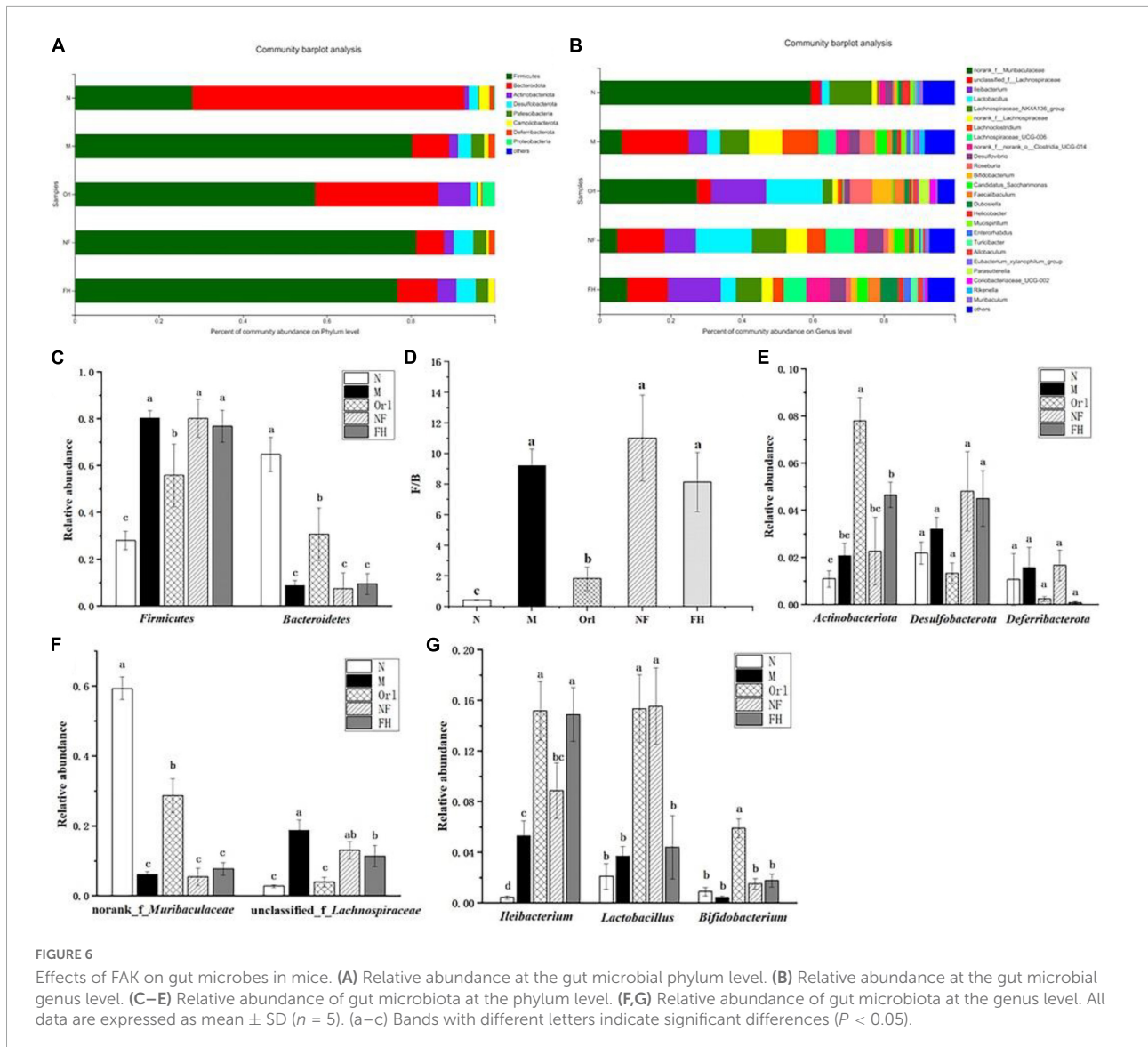
Obesity is a chronic metabolic disease that is characterized by excess accumulation of body fat in the body and has a serious negative impact on human health. In this study, *A. keiskei* was utilized as raw material for the first time to prepare Jiaosu via yeast fermentation and its anti-obesity effect was evaluated. The results showed that the contents of total flavonoid, γ -aminobutyric acid, chalcone, total phenol, free amino acids and protein of FAK were all dramatically enriched than the unfermented *A. keiskei*, indicating that the fermentation of yeast has a notably positive effect on the preparation of FAK, which may be attributed to the synergistic action of functional enzymes such as protease, cellulase, and pectinase secreted by yeast during the fermentation process. These enzymes decomposed substances such as macromolecular polysaccharides, proteins, and lipids that were difficult to be digested in plant cells into small molecules that can be easily absorbed by animals (21). Moreover, the microbial activity in the fermentation broth can also break the plant cells, thus accelerate the dissolution of phenols, flavonoids and other substances in the cells, and increase the yield. Moreover, in a high-fat induce



obese mouse model, FAK effectively reduced the body weights gain, blood lipid levels, damage, and fat accumulation in liver. We also proposed the underlying mechanism in the inhibition of the expressions for the obesity-related genes and regulation of gut microbiota.

Body weight and Lee's index can reflect the body shape of mice and are important indicators of the obesity degree in mice, and increase in liver index is often accompanied by elevation in hepatic steatosis (22, 23). Long-term intake of HFD will aggravate the expansion of adipose tissue, cause endocrine dysfunction, interfere with normal adipokines secretion and may also remodel adipose tissue by increasing the size of adipocytes. These disturbances are positively associated with metabolic disorders such as obesity, type 2 diabetes, and non-alcoholic fatty liver disease (24). In this study, both FAK and orlistat effectively reduced the weight of visceral (perirenal and

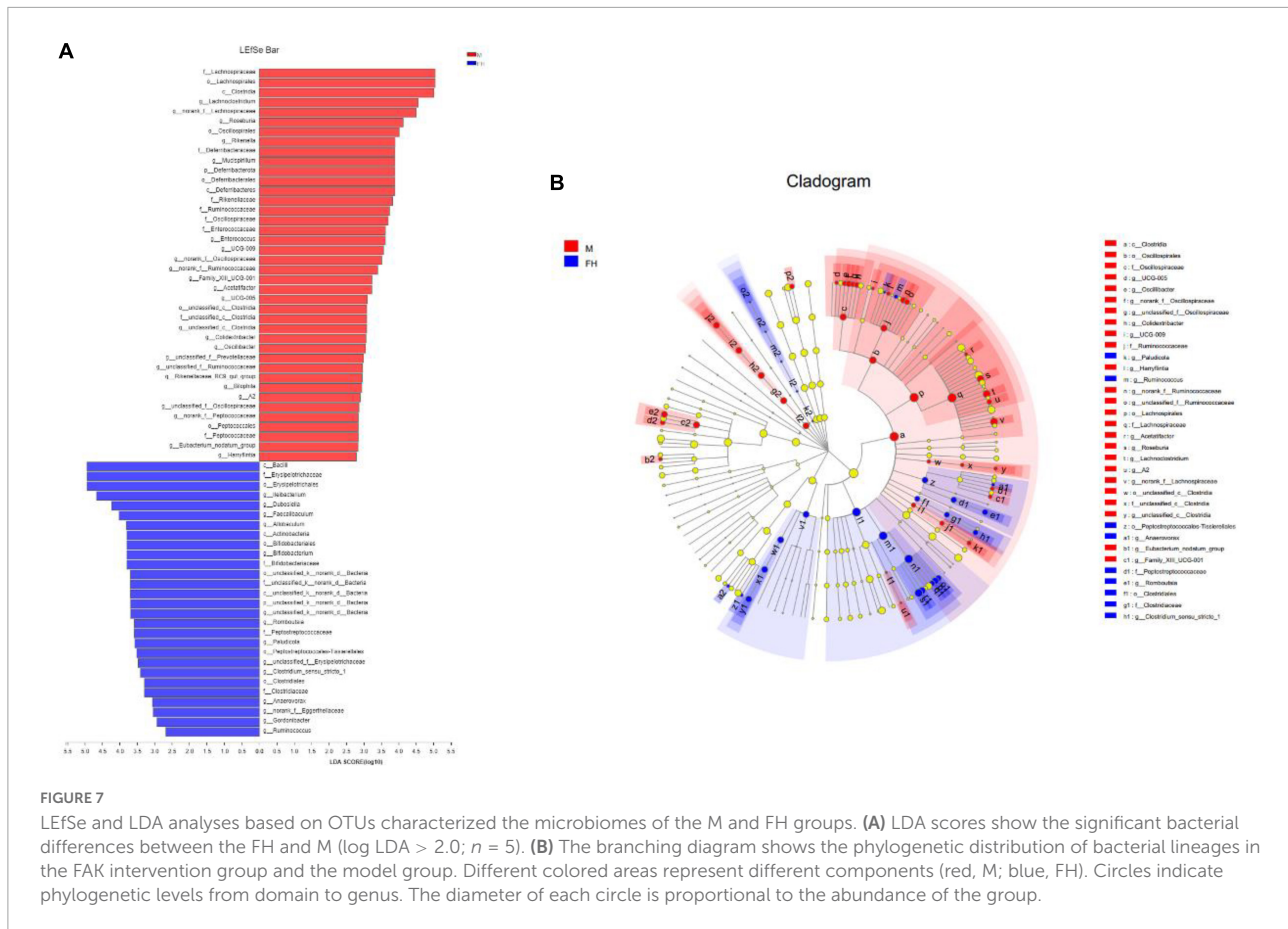
epididymal) adipose tissue in mice, and the effect of FAK was proportional to the intervention dose. These results suggested that FAK had a good weight loss effect in obese mice. Moreover, FAK but not NF significantly inhibited the hypertrophy of adipocytes suggesting the fermentation process enhanced the anti-obesity effect of *A. keiskei*. It has been reported that prophylactic removal of visceral adipose tissue can prominently improve obesity-induced insulin resistance and hepatic steatosis (25). Therefore, we speculate that FAK can improve the obesity-related metabolic disorders in mice by reducing visceral adipose tissue. The components of FAK are complex and the main ingredients are total flavonoids and GABA. Zhou et al. (26) reported that GABA-rich rice bran eminently decreased the body weight and blood lipid levels of HFD-fed rats. Zhang et al. demonstrated that two major flavonoids, 4-hydroxyderricin (4-HD) and xanthoangelol (XAG) from *A. keiskei* inhibit adipocyte



differentiation through the AMPK and MAPK pathways (27). The findings of Zhang et al. suggested that ingestion of flavonoid enriched fresh *A. keiskei* juice, which can prevent HFD-induced obesity and metabolic disorders by improving the intestinal microbiota composition (17). Our findings are consistent with these reports. In addition, the fermentation process also produces free amino acids, active peptides, polysaccharides, and other substances. Wang et al. employed the HPLC-MS method to prove that the components of fermented chili powder (FMPs) were transformed during fermentation, and FMPs had the higher quantity of free amino acids (28). The results of Liu et al. indicated that fermentation enhanced the anti-obesity effect of pepper on the HFD mice (29). The enriched ingredients during the fermentation process may explain the superior anti-obesity effect of FAK than NF. The liver is important for fat lipid metabolism, and the imbalance between lipid synthesis

and decomposition can lead to excessive fat deposition in liver, namely fatty liver (30). The serum ALT and AST levels are potential markers for the function of liver (31). Our study showed that FAK intervention effectively improved liver function, and reduced fat accumulation in liver tissue. Lee et al. (32) demonstrated that isopsoralen chalcone isolated from *A. keiskei* reduced hepatic fat deposition caused by high-fat cholesterol diet and alleviate hepatic steatosis. Several other studies have found that GABA could effectively protect ethanol-induced and dietary-induced hepatocyte injury (33, 34). The above studies have supported that FAK enriched substances can ameliorate hepatic steatosis and protect liver form dysfunction.

The precisely mechanism of obesity is largely unknown. Previous studies indicated that various hormones involved in its process. Leptin is a main signal of energy storage and has a negative feedback effect on energy intake (35, 36). However,

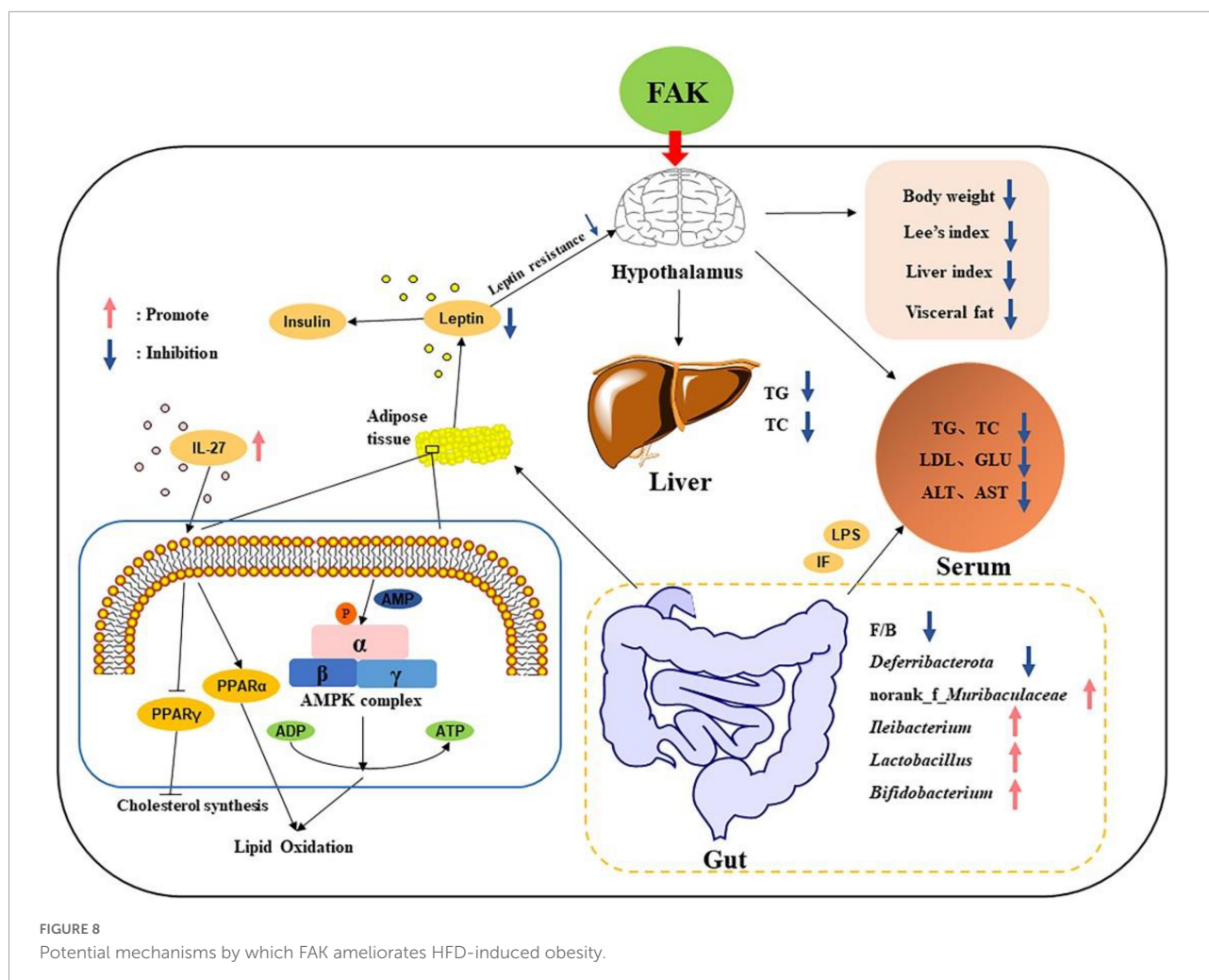


HFD can induce leptin resistance and lead to obesity resulting in increased blood leptin levels (37, 38). In our study, FAK treatment effectively reduced serum levels of leptin, indicated the improved leptin resistance after FAK intervention. IL-27 is a heterodimeric cytokine that regulates the immune system, promotes adipocyte browning and thermogenesis to reduce obesity symptoms (39). IL-27 could directly target adipocytes to activate p38 MAPK-PGC-1 α signaling and stimulate UCP1 production, which in turn upregulates PPAR α expression and promotes fat oxidation (39, 40). Herein, FAK dramatically increased the serum level of IL-27 in HFD-fed mice, suggesting the potential of promoting the burning of adipose tissue and fat oxidation in mice to ameliorate obesity. Taken together, the results indicated that the protective effect of FAK on obesity may be related to the regulation of Leptin and IL-27.

Obesity or excessive expansion of fat mass in the host mainly attribute to the unbalance between lipogenesis and lipolysis in adipose tissue. Lipogenesis involves in the synthesis of TGs and the terminal differentiation of preadipocytes into mature adipocytes. This process is tightly regulated by several key lipogenic transcription factors, such as PPAR γ and SREBP-1C, that regulate a variety of signaling molecules, like FAS, ACC, etc. (41). PPAR α is a key ligand-activated transcription factor that

mediated lipolysis (42). AMPK is a central regulator of energy balance. When energy intake is excessive, AMPK is activated by phosphorylation that promote fat and glucose metabolism to maintain homeostasis (43). Upon AMPK activation, it controls its downstream signals, such as PPAR α , UCP1 to promote lipolysis and fat oxidation (44, 45). In this study, we found that FAK up-regulated the expression of p-AMPK and PPAR α protein and decreased PPAR γ in adipose tissue of HFD mice, indicated the activation of lipolysis and inhibition of lipogenesis. Zhang et al. found that *A. keiskei* extract prevented obesity by regulating phosphorylation of AMPK in adipose tissue and liver (46). Another study confirmed that the natural flavonoid pre-treatment had an alleviative effect on alcoholic liver injury in mice, and the mechanism might be related to the regulation of PPAR α -mediated gene expression (47). Our data agree well with these results. Therefore, we suppose that FAK protected the mice from obesity through inhibiting fat synthesis and promoting lipolysis in adipose tissue.

Increasing evidence pointed that the gut microbiota plays primary roles in modulating host metabolism and health. Diet is considered to be the most important external factor affecting the homeostasis of gut microbiota distribution (48). Previous studies have shown that obesity is tightly associated with the gut



microbiota composition and the microbial diversity. Our data show that both FAK and orlistat significantly alleviated obesity and improved the related biochemical indicators. However, the diversity of intestinal flora in the Orl group was lower than that in the FH group. The reason may be that orlistat inhibited the activity of pancreatic lipase in the intestine and also affected the gut microbiota of mice (49), while FAK showed no harmful effect on the diversity of gut microbiota. At the phylum level, *Firmicutes* and *Bacteroidetes* are important categories in the intestinal flora, thus F/B (*Firmicutes/Bacteroidetes*, F/B) value is recognized as one of the prominent markers to evaluate obesity (50). Studies have shown that various specific gut bacteria can regulate intestinal permeability and inflammatory status of host to alleviate obesity-related metabolic syndrome (51). *Firmicutes* and *Bacteroidetes* are the most abundant phyla of humans and mice, and an elevated F/B value are considered as a hallmark of obesity (52). *Bacteroidetes* fermented dietary fibers to produce short-chain fatty acid acids (SCFAs) that is beneficial for health (53). *Actinobacteriota* was reported to be enriched in the feces of obese rodents (54). *Desulfobacterota* contributes

to obesity and its complications through the production of endotoxin lipopolysaccharide (55). *Deferribacterota* in the gut is associated with inflammation and development of obesity of the host (55–57). Herein, FAK intervention led to a decrease trend of *Firmicutes*, *Deferribacterota*, and the F/B values in HFD mice, which were complied with the amelioration of obesity. At the genus level, *norank_f_Muribaculaceae* is SCFAs producing bacteria and was significantly reduced in HFD-fed mice (58). *Lactobacillus* and *Bifidobacterium* are probiotics and are effective in reducing obesity and controlling metabolic disorders (59, 60). *Bifidobacterium* can also reduce metabolic endotoxemia by participating in the inhibition of pathogenic adhesion to the intestinal mucosa (61), stimulate bile acid biotransformation, promote fat digestion and influence lipids Metabolism (62). Lefse with LDA was employed to estimate the effect of each species abundance on the variance. The bacteria affecting the FH group were diverse and distributed in different taxa, such as *c_Bacilli* at the phylum level and *f_Erysipelotrichaceae*, *o_Erysipelotrichales*, and *g_Ileibacterium* at the genus level. Microbial diversity

makes the gut microbiota more resilient (20). In our study, the relative abundances of norank_f_Muribaculaceae, *Ileibacterium*, *Lactobacillus*, and *Bifidobacterium* were increased after FAK intervention, indicating the beneficial effect of FAK on gut microbiota. Collectively, it can be speculated that FAK might attenuate obesity by protecting the gut microbiota diversity and improving the gut microbiota composition.

5. Conclusion

In conclusion, our study shows that FAK can effectively ameliorate HFD-induced obesity, hyperlipemia, hepatic lipids accumulation in mice. The underlying mechanism may be related to the regulation of serum levels of leptin and IL-27, lipogenesis and lipolysis in adipose tissue and gut microbiota composition (Figure 8). This work provides key evidence that FAK prepared by yeast fermentation shows promising anti-obesity effects which can be expected to serve as a potential anti-obesity functional food.

Data availability statement

The data presented in the study have been deposited in the SRA repository, accession number PRJNA914722 (www.ncbi.nlm.nih.gov/sra/PRJNA914722).

Ethics statement

The animal study was reviewed and approved by the Ethics Committee of Qingdao University [SYXK (LU) 20200009].

Author contributions

KF and XG: investigation, data curation, formal analysis, methodology, software, visualization, and writing—original draft. XG: data curation, resources, validation, and writing—review and editing. PH: investigation, formal analysis, and

methodology. YH: formal analysis and visualization. RD: formal analysis. MW: funding acquisition. QL: funding acquisition and validation. ZL: conceptualization, funding acquisition, methodology, project administration, resources, supervision, validation, and writing—review and editing. All authors contributed to the article and approved the submitted version.

Funding

This research was funded by the Shandong Provincial Natural Science Foundation (ZR2019PH020) and Joint Institute of *A. keiskei* Health Industry Technology.

Acknowledgments

We wish to thank Shandong *A. keiskei* Biotech Co., Ltd., for supplying sufficient *A. keiskei* freeze-dried powders.

Conflict of interest

XG was employed by the Anqiu Huatao Food Co., Ltd. MW, QL, and ZL were employed by the Qingdao Balanson Biotech Co., Ltd.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

References

- Patel JJ, Rosenthal MD, Miller KR, Codner P, Kiraly L, Martindale RG. The critical care obesity paradox and implications for nutrition support. *Curr Gastroenterol Rep.* (2016) 18:45. doi: 10.1007/s11894-016-0519-8
- Blüher M. Obesity: global epidemiology and pathogenesis. *Nat Rev Endocrinol.* (2019) 15:288–98. doi: 10.1038/s41574-019-0176-8
- Ruban A, Stoenchev K, Ashrafian H, Teare J. Current treatments for obesity. *Clin Med.* (2019) 19:205–12. doi: 10.7861/clinmedicine.19-3-205
- Leonard W, Zhang P, Ying D, Adhikari B, Fang Z. Fermentation transforms the phenolic profiles and bioactivities of plant-based foods. *Biotechnol Adv.* (2021) 49:107763. doi: 10.1016/j.biotechadv.2021.107763
- Septembre-Malaterre A, Remize F, Poucheret P. Fruits and vegetables, as a source of nutritional compounds and phytochemicals: changes in bioactive compounds during lactic fermentation. *Food Res Int.* (2018) 104:86–99. doi: 10.1016/j.foodres.2017.09.031

6. Verón HE, Gauffin Cano P, Fabersani E, Sanz Y, Isla MI, Espinar MT, et al. Cactus pear (*Opuntia ficus-indica*) juice fermented with autochthonous *Lactobacillus plantarum* S-811. *Food Funct.* (2019) 10:1085–97. doi: 10.1039/c8fo01591k
7. Pan Y, Tan J, Long X, Yi R, Zhao X, Park KY. Anti-obesity effect of fermented lemon peel on high-fat diet-induced obese mice by modulating the inflammatory response. *J Food Biochem.* (2022) 46:e14200. doi: 10.1111/jfbc.14200
8. Cho D, Jeong HW, Kim JK, Kim AY, Hong YD, Lee JH, et al. Gallic catechin gallate-containing fermented green tea extract ameliorates obesity and hypertriglyceridemia through the modulation of lipid metabolism in adipocytes and myocytes. *J Med Food.* (2019) 22:779–88. doi: 10.1089/jmf.2018.4327
9. Kil YS, Pham ST, Seo EK, Jafari M. *Angelica keiskei*, an emerging medicinal herb with various bioactive constituents and biological activities. *Arch Pharm Res.* (2017) 40:655–75. doi: 10.1007/s12272-017-0892-3
10. Wu X, Jiang Q, Zhang R. Research progress on the health care function of ashitaba and its product development. *Food Sci.* (2016) 37:286–91.
11. Akihisa T, Tokuda H, Ukiya M, Iizuka M, Schneider S, Ogasawara K, et al. Chalcones, coumarins, and flavanones from the exudate of *Angelica keiskei* and their chemopreventive effects. *Cancer Lett.* (2003) 201:133–7. doi: 10.1016/s0304-3835(03)00466-x
12. Pang XN, Gao X, Liu F, Jiang Y, et al. Xanthoangelol modulates caspase-1-dependent pyroptotic death among hepatocellular carcinoma cells with high expression of GSDMD. *J Funct Foods.* (2021) 84:104577. doi: 10.1016/j.jff.2021.104577
13. Zhang L, Jiang Y, Pang X, Hua P, Gao X, Li Q, et al. Simultaneous optimization of ultrasound-assisted extraction for flavonoids and antioxidant activity of *Angelica keiskei* using response surface methodology (RSM). *Molecules.* (2019) 24:3461. doi: 10.3390/molecules24193461
14. Kim A, Lim JW, Kim H, Kim H. Supplementation with *Angelica keiskei* inhibits expression of inflammatory mediators in the gastric mucosa of *Helicobacter pylori*-infected mice. *Nutr Res.* (2016) 36:488–97. doi: 10.1016/j.nutres.2015.12.017
15. Hu S, Wang X, Dai Z, Lan X, Li X. Screening of γ -aminobutyric acid monascus strain and its fermentation conditions. *Food Ferment Technol.* (2011) 47:64–7.
16. Khaleghnezhad V, Yousefi AR, Tavakoli A, Farajmand B. Interactive effects of abscisic acid and temperature on rosmarinic acid, total phenolic compounds, anthocyanin, carotenoid and flavonoid content of dragonhead (*Dracocephalum moldavica* L.). *Sci Hortic.* (2019) 250:302–9. doi: 10.1016/j.scienta.2019.02.057
17. Zhang C, Wu W, Li X, Xin X, Liu D. Daily supplementation with Fresh *Angelica keiskei* juice alleviates high-fat diet-induced obesity in mice by modulating gut microbiota composition. *Mol Nutr Food Res.* (2019) 63:e1900248. doi: 10.1002/mnfr.201900248
18. Lin Y, Ren N, Li S, Chen M, Pu P. Novel anti-obesity effect of scutellarein and potential underlying mechanism of actions. *Biomed Pharmacother.* (2019) 117:109042. doi: 10.1016/j.biopha.2019.109042
19. Liu G, Huang Y, Reis FS, Song D, Ni H. Impact of nutritional and environmental factors on inflammation, oxidative stress, and the microbiome 2019. *Biomed Res Int.* (2019) 2019:5716241. doi: 10.1155/2019/5716241
20. Huhe, Chen X, Hou F, Wu Y, Cheng Y. Bacterial and fungal community structures in loess plateau grasslands with different grazing intensities. *Front Microbiol.* (2017) 8:606. doi: 10.3389/fmicb.2017.00606
21. Shi H, Zhang M, Wang WQ, Devahastine S. Solid-state fermentation with probiotics and mixed yeast on properties of okara. *Food Biosci.* (2020) 36:100610.
22. Feng K, Zhu X, Chen T. Effects of tangerine peel on high-fat diet-induced lipid-lowering weight loss and prevention of fatty liver formation in rats. *Food Sci.* (2018) 39:213–20.
23. Chen P, Wang X, Song W. Weight-reducing effect of pigment from Pu-erh tea on diet-induced obese rats. *Food Res Dev.* (2011) 32:169–71.
24. Hirata BKS, Cruz MM, de Sá RDCC, Farias TSM, Machado MMF, Bueno AA, et al. Potential anti-obesogenic effects of ginkgo biloba observed in epididymal white adipose tissue of obese rats. *Front Endocrinol.* (2019) 10:284. doi: 10.3389/fendo.2019.00284
25. Franczyk MP, He M, Yoshino J. Removal of epididymal visceral adipose tissue prevents obesity-induced multi-organ insulin resistance in male mice. *J Endocr Soc.* (2021) 5:bvab024. doi: 10.1210/10.1210/jendso/bvab024
26. Zhou Z, Zhang H, Liu Z. Evaluation of the effects of GABA rice bran on glucose and lipid metabolism in rats fed a high-fat diet. *Food Sci Technol.* (2017) 42:188–91. doi: 10.13684/j.cnki.spkj.2017.01.040
27. Zhang T, Sawada K, Yamamoto N, Ashida H. 4-Hydroxyderricin and xanthoangelol from ashitaba (*Angelica keiskei*) suppress differentiation of preadipocytes to adipocytes via AMPK and MAPK pathways. *Mol Nutr Food Res.* (2013) 57:1729–40. doi: 10.1002/mnfr.201300020
28. Wang Y, Zhou H, Ding S, Ye M, Jiang L, Wang R. Changes in free amino acids of fermented minced peppers during natural and inoculated fermentation process based on HPLC-MS/MS. *J Food Sci.* (2020) 85:2803–11. doi: 10.1111/1750-3841.15363
29. Liu L, Ding C, Tian M, Yi D, Wang J, Zhao J, et al. Fermentation improves the potentiality of capsaicin in decreasing high-fat diet-induced obesity in C57BL/6 mice by modulating lipid metabolism and hormone response. *Food Res Int.* (2019) 124:49–60. doi: 10.1016/j.foodres.2018.10.015
30. Chung YH, Bang JS, Kang CM, Goh JW, Lee HS, Hong SM, et al. Aqueous extract of humulus japonicus attenuates hyperlipidemia and fatty liver in obese mice. *J Med Food.* (2018) 21:999–1008. doi: 10.1089/jmf.2017.4135
31. Hakkak R, Gauss CH, Bell A, Korourian S. Short-term soy protein isolate feeding prevents liver steatosis and reduces serum ALT and AST Levels in obese female zucker rats. *Biomedicine.* (2018) 6:55. doi: 10.3390/biomedicine620055
32. Lee H, Li H, Kweon M, Choi Y, Kim MJ, Ryu JH. Isobavachalcone from *Angelica keiskei* inhibits adipogenesis and prevents lipid accumulation. *Int J Mol Sci.* (2018) 19:1693. doi: 10.3390/ijms19061693
33. Wu A, Li L, Zhang S, Lin Q, Liu J. Optimization of the hongqu starter preparation process for the manufacturing of red mold rice with high gamma-aminobutyric acid production by solid-state fermentation. *Biotechnol Appl Biochem.* (2022). [Epub ahead of print]. doi: 10.1002/bab.2370
34. Norikura T, Kojima-Yuasa A, Opare Kennedy D, Matsui-Yuasa I. Protective effect of gamma-aminobutyric acid (GABA) against cytotoxicity of ethanol in isolated rat hepatocytes involves modulations in cellular polyamine levels. *Amino Acids.* (2007) 32:419–23. doi: 10.1007/s00726-006-0381-3
35. Li D, Wu H, Dou H. Weight loss effect of sweet orange essential oil microcapsules on obese SD rats induced by high-fat diet. *Biosci Biotechnol Biochem.* (2019) 83:923–32. doi: 10.1080/09168451.2019.1578640
36. Pan H, Guo J, Su Z. Advances in understanding the interrelations between leptin resistance and obesity. *Physiol Behav.* (2014) 130:157–69. doi: 10.1016/j.physbeh.2014.04.003
37. Han H, Zhou W. Leptin and its derivatives: a potential target for autoimmune diseases. *Curr Drug Targets.* (2019) 20:1563–71. doi: 10.2174/1389450120666190729120557
38. Lee HS, Nam Y, Chung YH, Kim HR, Park ES, Chung SJ, et al. Beneficial effects of phosphatidylcholine on high-fat diet-induced obesity, hyperlipidemia and fatty liver in mice. *Life Sci.* (2014) 118:7–14. doi: 10.1016/j.lfs.2014.09.027
39. Wang Q, Li D, Cao G, Shi Q, Zhu J, Zhang M, et al. IL-27 signalling promotes adipocyte thermogenesis and energy expenditure. *Nature.* (2021) 600:314–8. doi: 10.1038/s41586-021-04127-5
40. Zhang H, Li Q, Teng Y, Lin Y, Li S, Qin T, et al. Interleukin-27 decreases ghrelin production through signal transducer and activator of transcription 3-mechanistic target of rapamycin signaling. *Acta Pharm Sin B.* (2020) 10:837–49. doi: 10.1016/j.apsb.2019.12.018
41. Zhang F, Ye J, Zhu X, Wang L, Gao P, Shu G, et al. Anti-obesity effects of dietary calcium: the evidence and possible mechanisms. *Int J Mol Sci.* (2019) 20:3072. doi: 10.3390/ijms20123072
42. Rigano D, Sirignano C, Tagliatalata-Scafati O. The potential of natural products for targeting PPAR α . *Acta Pharm Sin B.* (2017) 7:427–38. doi: 10.1016/j.apsb.2017.05.005
43. Meeks KAC, Bentley AR, Adeyemo AA, Rotimi CN. Evolutionary forces in diabetes and hypertension pathogenesis in Africans. *Hum Mol Genet.* (2021) 30:R110–8. doi: 10.1093/hmg/ddaa238
44. López M. Hypothalamic AMPK and energy balance. *Eur J Clin Invest.* (2018) 48:e12996. doi: 10.1111/eci.12996
45. Li Z, Li J, Miao X, Cui W, Miao L, Cai L. A minireview: role of AMP-activated protein kinase (AMPK) signaling in obesity-related renal injury. *Life Sci.* (2021) 265:118828. doi: 10.1016/j.lfs.2020.118828
46. Zhang T, Yamashita Y, Yasuda M, Yamamoto N, Ashida H. Ashitaba (*Angelica keiskei*) extract prevents adiposity in high-fat diet-fed C57BL/6 mice. *Food Funct.* (2015) 6:135–45. doi: 10.1039/c4fo00525b
47. Wang F, Liu JC, Zhou RJ, Zhao X, Liu M, Ye H, et al. Apigenin protects against alcohol-induced liver injury in mice by regulating hepatic CYP2E1-mediated oxidative stress and PPAR α -mediated lipogenic gene expression. *Chem Biol Interact.* (2017) 275:171–7. doi: 10.1016/j.cbi.2017.08.006
48. Peng C, Xu X, Li Y, Li X, Yang X, Chen H, et al. Sex-specific association between the gut microbiome and high-fat diet-induced metabolic disorders in mice. *Biol Sex Differ.* (2020) 11:5. doi: 10.1186/s13293-020-0281-3

49. Ke J, An Y, Cao B, Lang J, Wu N, Zhao D. Orlistat-induced gut microbiota modification in obese mice. *Evid Based Complement Alternat Med.* (2020) 2020:9818349. doi: 10.1155/2020/9818349
50. Gao X, Du L, Randell E, Zhang H, Li K, Li D. Effect of different phosphatidylcholines on high fat diet-induced insulin resistance in mice. *Food Funct.* (2021) 12:1516–28. doi: 10.1039/d0fo02632h
51. Carmody RN, Gerber GK, Luevano JM Jr, Gatti DM, Somes L, Svenson KL, et al. Diet dominates host genotype in shaping the murine gut microbiota. *Cell Host Microbe.* (2015) 17:72–84. doi: 10.1016/j.chom.2014.11.010
52. Li X, Wang Y, Zhou J, Wang Z, Wang Y, Zheng J, et al. Mixed nuts with high nutrient density improve insulin resistance in mice by gut microbiota remodeling. *Food Funct.* (2022) 13:9904–17. doi: 10.1039/d2fo01479c
53. Chavoya-Guardado MA, Vasquez-Garibay EM, Ruiz-Quezada SL, Ramírez-Cordero MI, Larrosa-Haro A, Castro-Albarran J. *Firmicutes*, *Bacteroidetes* and *Actinobacteria* in human milk and maternal adiposity. *Nutrients.* (2022) 14:2887. doi: 10.3390/nu14142887
54. Gao X, Zhang H, Li K, Shi Y, Guo X, Wang L. Sandalwood seed oil improves insulin sensitivity in high-fat/high-sucrose diet-fed rats associated with altered intestinal microbiota and its metabolites. *Food Funct.* (2021) 12:9739–49. doi: 10.1039/d1fo02239c
55. Hu S, Xu Y, Gao X, Li S, Jiang W, Liu Y, et al. Long-chain bases from sea cucumber alleviate obesity by modulating gut microbiota. *Mar Drugs.* (2019) 17:455. doi: 10.3390/md17080455
56. Chung YW, Gwak HJ, Moon S, Rho M, Ryu JH. Functional dynamics of bacterial species in the mouse gut microbiome revealed by metagenomic and metatranscriptomic analyses. *PLoS One.* (2020) 15:e0227886. doi: 10.1371/journal.pone.0227886
57. Zeng H, Larson KJ, Cheng WH, Bukowski MR, Safratowich BD, Liu Z, et al. Advanced liver steatosis accompanies an increase in hepatic inflammation, colonic, secondary bile acids and *Lactobacillaceae/Lachnospiraceae* bacteria in C57BL/6 mice fed a high-fat diet. *J Nutr Biochem.* (2020) 78:108336. doi: 10.1016/j.jnutbio.2019.108336
58. Lagkouvardos I, Lesker TR, Hitch TCA, Gálvez EJC, Smit N, Neuhaus K, et al. Sequence and cultivation study of *Muribaculaceae* reveals novel species, host preference, and functional potential of this yet undescribed family. *Microbiome.* (2019) 7:28. doi: 10.1186/s40168-019-0637-2
59. Xu Z, Jiang W, Huang W, Lin Y, Chan FKL, Ng SC. Gut microbiota in patients with obesity and metabolic disorders—A systematic review. *Genes Nutr.* (2022) 17:2. doi: 10.1186/s12263-021-00703-6
60. Borgeraas H, Johnson LK, Skattebu J, Hertel JK, Hjelmestaeth J. Effects of probiotics on body weight, body mass index, fat mass and fat percentage in subjects with overweight or obesity: a systematic review and meta-analysis of randomized controlled trials. *Obes Rev.* (2018) 19:219–32. doi: 10.1111/obr.12626
61. Sroka-Oleksiak A, Młodzińska A, Bulanda M, Salamon D, Major P, Stanek M, et al. Metagenomic analysis of duodenal microbiota reveals a potential biomarker of dysbiosis in the course of obesity and type 2 diabetes: a pilot study. *J Clin Med.* (2020) 9:369. doi: 10.3390/jcm9020369
62. Gao R, Zhu C, Li H, Yin M, Pan C, Huang L, et al. Dysbiosis signatures of gut microbiota along the sequence from healthy, young patients to those with overweight and obesity. *Obesity.* (2018) 26:351–61. doi: 10.1002/oby.22088