

## Basic Study

## Maternal inappropriate calcium intake aggravates dietary-induced obesity in male offspring by affecting the differentiation potential of mesenchymal stem cells

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## Abstract

## BACKGROUND

The effects of inappropriate dietary calcium intake in early life on later obesity have not been fully elucidated.

## AIM

To raise the mechanism of maternal calcium intake on the multi-differentiation potential of mesenchymal stem cells among their male offspring.

## METHODS

Four-week-old female C57BL/6N mice were fed by deficient, low, normal and excessive calcium reproductive diets throughout pregnancy and lactation. Bone MSCs (BMSCs) were obtained from 7-day-old male offspring to measure the adipogenic differentiation potential by the Wnt/ $\beta$ -catenin signaling pathway. The other weaning male pups were fed a high-fat diet for 16 wk, along with normal-fat diet as the control. Then the serum was collected for the measurement of biochemical indicators. Meanwhile, the adipose tissues were excised to analyze the adipocyte sizes and inflammatory infiltration. And the target gene expressions on the adipogenic differentiation and Wnt/ $\beta$ -catenin signaling pathway in the adipose tissues and BMSCs were determined by real-time reverse transcription polymerase chain reaction.

## RESULTS

Compared with the control group, maternal deficient, low and excessive calcium intake during pregnancy and lactation aggravated dietary-induced obesity, with larger adipocytes, more serious inflammatory infiltration and higher serum metabolism indicators by interfering with higher expressions of adipogenic differentiation (*PPAR $\gamma$* , *C/EBP $\alpha$* , *Fabp4*, *LPL*, *Adiponectin*, *Resistin* and/or *Leptin*) among their male offspring ( $P < 0.05$ ). And there were significantly different expression of similar specific genes in the BMSCs to successfully polarize adipogenic differentiation and suppress osteogenic differentiation *in vivo* and *in vitro*, respectively ( $P < 0.05$ ). Meanwhile, it was accompanied by more significant disorders on the expressions of Wnt/ $\beta$ -catenin signaling pathway both in BMSCs and adulthood adipose tissues among the offspring from maternal inappropriate dietary calcium intake groups.

## CONCLUSION

Early-life abnormal dietary calcium intake might program the adipogenic differentiation potential of BMSCs from male offspring, with significant expressions on the Wnt/ $\beta$ -catenin signaling pathway to aggravate high-fat-diet-induced obesity in adulthood.

**Key Words:** Calcium; Obesity; Bone mesenchymal stem cells; Wnt/ $\beta$ -catenin signaling pathway; Adipogenic differentiation; Male offspring

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**Core tip:** Maternal inappropriate dietary calcium intake could aggravate high-fat-diet-induced obesity among male offspring, with larger adipocytes and more serious inflammatory infiltration by interfering with the higher expressions of adipogenic genes, which was accompanied by significant expressions of specific genes on the adipogenic and osteogenic differentiation. It was worsened by the disorders of Wnt/ $\beta$ -catenin signaling pathway both in the BMSCs and adipose tissues. So the importance of this study was that the prevention of adulthood obesity could be moved forward to the appropriate calcium intake in the neonatal period, even the formation of maternal germ cells and fertilized egg.

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## INTRODUCTION

Obesity has become a worldwide noncommunicable health crisis with rising prevalence in the past few decades due to excess calorie intake, fat accumulation, and adiposity[1,2]. It can cause severe metabolic disorders such as nonalcoholic steatohepatitis, type 2 diabetes, cardiovascular diseases, and cancer[1-3]. These above pathological complications are characterized by more hypertrophy and hyperplasia of adipocytes to cause dynamic expansion in the adipose tissues, in which hyperplasia is a complicated process including disruption of the commitment of mesenchymal stem cells (MSCs) to form preadipocytes, and terminal differentiation from preadipocytes to mature adipocytes[4-6]. And MSCs (CD29+, CD90+, Sca-1+, CD31-, CD34-, CD45- and CD49d-), as a group of cells with multi-lineage differentiated potential and self-renewal capacity, are the major original sources of mature adipocytes. , in which the key coordinated cascade of transcription factors were mainly included *PPAR $\gamma$* , *C/EBP $\alpha$* , *LPL* and *FABP4*, with the significant secretory molecules such as *Leptin*, *Adiponectin* and *Resistin*[7,8]. In this process, the mechanisms governing the adipogenic differentiation of MSCs can be regulated by the coordination of complex networks in many signaling pathways, such as JAK2/STAT3, SIRT1/SIRT2, ERK1/ERK2, TGF- $\beta$ /BMP, Wnt/ $\beta$ -catenin and RHO-family GTPase[9,10], in which the activation of Wnt/ $\beta$ -catenin signaling can inhibit adipogenic differentiation and promote osteogenic differentiation through endogenous regulatory genes (*CTNNB1*, *Wnt1*, *Wnt10a*, *Wnt10b*, *Wnt5a*, *Gsk3 $\beta$* , *Axin2* and *TGF7L2*)[11,12]. It has been demonstrated that the differentiation potential of MSCs mainly occurs in early life, and the numbers and differentiation potential significantly decline with the age[12], so the nutritional status and exposure to adverse factors at this stage, especially pregnancy and lactation, are important for the differentiation potential of MSCs to affect later metabolic disturbances in adulthood [13-16].

Calcium is an important functional nutrient on the regulation of energy balance and glucose uptake in the battle against obesity[17-19]. However, daily calcium intake is still lower than its recommended nutrient intake among pregnant women[20,21], so the imbalance of calcium intake in early life may have detrimental effects on later health. Both our high-fat-diet (HFD) induced obese mouse model and epidemiological cohorts showed that both dietary insufficient and excessive calcium intake during pregnancy and lactation increased body weight gain by affecting the gut microbiota structure, and abnormal expression of lipolysis and liposynthesis among their male offspring [22-24]. However, the specific mechanisms by which maternal calcium intake modulates body weight and fat and glucose homeostasis of their infants are still not fully understood. Some research had found that  $\text{Ca}^{2+}$  formed in the culture medium had osteo-inductive properties to promote osteogenic differentiation of MSCs[25]. Previous studies also had demonstrated that neonatal calcium deficiency could reduce the osteogenic priming of MSCs by enlarging the subpopulation with adipogenic potential in piglets and mice *in vivo* [26]. Furthermore, it is competing and reciprocal on the balance of adipogenic and osteogenic differentiation of MSCs[27,28]. However, whether maternal inappropriate dietary calcium intake can increase the adipogenic differentiation potential of MSCs among their male offspring is still unclear.

Thus, this study was designed to investigate whether abnormal dietary calcium intake during gestation and lactation affected the multi-differentiation potential of bone MSCs (BMSCs) to aggravate the development of adulthood obesity among their male offspring and explore the possible signaling pathways. This deeper understanding of early-life calcium intake could play a significant role on preventing later obesity.

## MATERIALS AND METHODS

### Animal procedures

Sixty 4-week-old C57BL/6N female mice were obtained from Beijing Vital River Laboratory Animal Technology (License SCXK-Beijing) and housed at the Animal Center in the Academy of Military Medical Sciences under a 12-h light/dark cycle (lights-on 08:00 h) with adequate food and water intake at 22°C and 50% humidity. All mice were randomly divided into four groups ( $n = 15/\text{group}$ ) and fed with the deficient (DC, 0.05%), low (LC, 0.25%), normal (NC, 0.70%) and high-calcium (HC, 1.20%) reproductive diets respectively for 6 wk. Five mice in each group ( $n = 5/\text{group}$ ) were killed to determine the maternal contents of calcium and other metabolic indicators in the serum before mating. Then the remaining mice ( $n = 10/\text{group}$ ) were mated with 10-week-old C57BL/6N male mice from Beijing Vital River Laboratory Animal Technology (2:1/cage), and continued on their own diets throughout gestation and lactation. According to the previous studies[21-23], the male offspring were used to study the development of obesity after the different calcium interventions during pregnancy and lactation. The 7-day-old male offspring ( $n = 9/\text{group}$  from more than three cages) in each group were killed to obtain BMSCs. While at age 21 d, the male offspring in the DC, LC, NC and HC groups ( $n = 10/\text{group}$ ) were respectively weaned onto the HFD (34.9% fat by weight, 60% kcal, No. H10060) for 16 wk (NC-HFD, DC-HFD, LC-HFD and HC-HFD groups); with the normal fat diet (4.3% fat by weight, 10% kcal, No. H10010) as the control (NC-C group). All above diets were prepared by Beijing HFK Bioscience Co. Ltd. (<http://www.hfkbio.com/>) (Table 1). Body weight, food intake and energy intake in the NC-C, NC-HFD, DC-HFD, LC-HFD and HC-HFD groups were recorded weekly. Their blood samples were collected through the eye-drop, then they were anesthetized by the carbon dioxide inhalation. Immediately, their adipose tissues including the epididymal white adipose tissue (eWAT), inguinal white adipose tissue (iWAT), kidney adipose tissue (KAT) and brown adipose tissue (BAT) were freely dissected from the surrounding tissues, in which some were fixed in 10% phosphate-buffered formalin, some were stored in phosphate-buffered saline (PBS) to analyze the percentage of immune cells, and the remainder was frozen in liquid  $\text{N}_2$ . The serum samples were separated at 3000 r/min for 15 min after they were kept for 30 min at room temperature. All biological samples were stored in a -80°C refrigerator until use.

All animal studies were approved and conducted in accordance with the Beijing Academy of Military Medical Sciences Guide for the Care and Usage Committee of Laboratory Animals. The animal protocol was approved by the Ethics of Animal Experiments in the Academy of Military Medical Sciences in China (No. IACUC-DWZX-2019-704).

### Measurement of the biochemical indicators

The concentrations of serum calcium, glucose, triglyceride (TG) and total cholesterol (TC) were respectively measured by the coloristic methods using the enzymatic assay kits (Maccura Biotechnology Co. Ltd., Sichuan, China). The male offspring mice in each group were orally gavaged with 20% glucose (weight/volume: 2.0g/kg) after a 10-h overnight fast and blood samples were collected from the tail vein at 15, 30, 60, 90 and 120 min to determine the glucose content (oral glucose tolerance test, OGTT) at 13 wk. The insulin tolerance test (ITT) was performed 1 wk after the OGTT, in which the blood samples were collected from the tail vein for the determination of blood glucose after 2 h fasting.

**Table 1** Details of diet formulations in this study (g/kg diet)

Ingredients (g)	Reproductive diets				Feeding diets (0.70%)	
	Deficient calcium diet (0.05%)	Low calcium diet (0.25%)	Normal calcium diet (0.70%)	High calcium diet (1.20%)	Normal fat diet (H10010)	High fat diet (H10060)
Casein	200.00	200.00	200.00	200.00	189.58	258.45
Cystine	3.00	3.00	3.00	3.00	2.84	3.88
Cornstarch	396.30	391.30	380.00	367.50	298.59	161.53
Maltodextrin	132.00	132.00	132.00	132.00	33.18	88.91
Sucrose	100.00	100.00	100.00	100.00	331.77	---
Fibrin	50.00	50.00	50.00	50.00	47.40	64.61
Soybean oil	70.00	70.00	70.00	70.00	23.70	32.31
Lard oil	---	---	---	---	18.96	316.60
Mineral mixture without calcium (M1004)	35.00	35.00	35.00	35.00	---	---
Mineral mixture (M1002)	---	---	---	---	9.48	12.92
Calcium bicarbonate	---	---	---	---	12.32	16.80
Calcium carbonate (CaCO <sub>3</sub> )	1.25	6.25	17.50	30.00	5.21	7.11
Potassium citrate, H <sub>2</sub> O	---	---	---	---	15.64	21.32
Vitamin mixture (V1002)	10.00	10.00	10.00	10.00	9.48	12.92
Choline Bitartrate	2.50	2.50	2.50	2.50	1.90	2.58
Antioxidant (TBHQ)	0.014	0.014	0.014	0.014	0.047	0.065

### Analysis of the immune cells in the adipose tissues

Stromal vascular fraction cells (SVFs) were extracted from the eWAT and iWAT in PBS. The infiltration and percentages of M1 macrophages (CD45<sup>+</sup>CD64<sup>+</sup>CD11C<sup>+</sup>), M2 macrophages (CD45<sup>+</sup>CD64<sup>+</sup>CD11C<sup>-</sup>) and adipose tissue dendritic cells (ATDCs, CD45<sup>+</sup>CD64<sup>+</sup>CD11C<sup>+</sup>) were determined using the BD FACSCanto II Flow Cytometer (BD Biosciences, USA), and analyzed by FlowJo flow cytometry software (Treestar Inc., Ashland, OH, USA).

### Histological analysis of adipose tissues

eWAT, iWAT, KAT and BAT in the NC-C, DC-HFD, LC-HFD, NC-HFD and HC-HFD groups were embedded in paraffin and cut into 6-μm sections, and stained with hematoxylin and eosin to measure the adipocyte size and inflammatory infiltration under a light microscope at 200× magnification, and analyzed by Image-pro Plus. All above histological experiments were performed by Servicebio (Beijing, China).

### Gene expression related to adipogenic differentiation and Wnt/β-catenin signaling pathway in adipose tissues

Total RNA in eWAT, iWAT, KAT and BAT was extracted using TRIzol Reagent (Invitrogen, Carlsbad, CA, USA), and cDNA was reverse transcribed by Transcript<sup>®</sup>One-Step gDNA Removal and cDNA Synthesis SuperMix (TransGen Biotech, China).

The genes related to adipogenic differentiation (*PPARγ*, *C/EBPα*, *LPL*, *Fabp4*, *Adiponectin*, *Resistin* and *Leptin*) and Wnt/β-catenin signaling pathway (*CTNNB1*, *Wnt1*, *Wnt10a*, *Wnt10b*, *Wnt5a*, *Gsk3β*, *Axin2* and *TGF7L2*) were determined by real-time reverse transcription polymerase chain reaction (RT-PCR) (No. AQ101-03, TransStart<sup>®</sup>Green qPCR SuperMix, TransGen Biotech, China) (CFX-96; Bio-Rad, USA), and 36B4 was the invariant internal gene (Supplementary Table 1). Gene expression was normalized using the 2<sup>-CT</sup> method.

### BMSC derivation and maintenance from male offspring

BMSCs from 7-day-old male offspring in the NC, DC, LC and HC groups were isolated and cultured as follows. The tibia and fibula from three pups with different mothers were isolated after washing with PBS to remove the residual muscle and blood under sterile conditions. They were shredded into small pieces of 2 mm<sup>3</sup> and digested in 0.1% type II collagenase (Gibco) at 37°C for 40 min, and transferred into α



-minimal essential medium ( $\alpha$ -MEM) (Gibco) containing 10% fetal bovine serum (FBS; Gibco), 100 U/mL penicillin, and 100 mg/mL streptomycin. The medium with FBS was changed every 3 d. When the adherent BMSCs reached 80%–90% confluence, they were collected in 0.25% trypsin (Gibco) and subcultured at a ratio of 1:3 for the further expansion and identification until the P3 generation, and used for subsequent experiments.

### **Detection of cell cycle and surface antibodies of BMSCs by flow cytometry**

P3 generation BMSCs ( $1 \times 10^6$ ) from the DC, LC, NC and HC groups were phenotypically fixed, stained and characterized by the antibody permeabilization process. Mouse phycoerythrin-conjugated monoclonal antibodies Sca-1 (AB\_2539218, MA5-17834), CD90 (AB\_469640, 25-0900-82) and CD31 (AB\_657735, 17-0311-82) (eBioscience, Waltham, MA, USA), and fluorescein-isothiocyanate-conjugated antibodies including CD29 (AB\_2572449, 11-0291-82), CD34 (AB\_465021, 11-0341-82) and CD45 (AB\_465050, 11-0451-82) and CD49d (AB\_465083, 11-0492-82) (eBioscience) were used to detect the purity of BMSCs. The Cell Cycle and Apoptosis Analysis kit (Beyotime) was obtained to measure the cell cycle of BMSCs. All above signals were recorded by flow cytometry with the FACScalibur system (Becton Dickinson) and analyzed using FlowJo software (Supplementary Figure 1 and Table 2).

### **Differentiation potential of BMSCs**

To identify the adipogenic differentiation potential of BMSCs in the NC, DC, LC and HC groups, P3 BMSCs ( $8 \times 10^4$ ) were cultured with  $\alpha$ -MEM containing 10% FBS and the related adipogenic inducer ( $10^3$  mM dexamethasone, 0.5 mM isobutyl methylxanthine, 0.2 mM indomethacin, and 10  $\mu$ g/mL insulin) (Sigma, Germany) for 7 d, in which the medium was changed every 3 d. Self-differentiated BMSCs without the above inducers ( $3 \times 10^4$ ) were as the controls. The induced and self-differentiated BMSCs were stained with Oil Red O (Sigma, Germany) and measured the gene expression related to adipogenic differentiation[29].

The osteogenic differentiation capacity of BMSCs was assessed by incubating the cells ( $3 \times 10^4$ ) with  $\alpha$ -MEM containing 10% FBS and osteogenic inducer ( $10^7$  mM dexamethasone, 0.5 mM ascorbic acid, and 10 mM  $\beta$ -glycerol phosphate) (Sigma, Germany) for 10 d, while the self-differentiated BMSCs without the above inducers ( $3 \times 10^4$ ) were as the controls. To demonstrate the osteogenic differentiation capacity of BMSCs, they were identified by immunocytochemical staining with alkaline phosphatase[29]. Expression of genes related to osteogenic differentiation were determined by RT-PCR.

### **Quantitative RT-PCR of BMSCs**

Total RNA was extracted from the P3 BMSCs, adipogenic and osteogenic differentiated BMSCs and their related self-differentiated BMSCs using TRIzol Reagent (Invitrogen), and their cDNA samples were reverse transcribed by Transcript<sup>®</sup>One-Step gDNA Removal and cDNA Synthesis SuperMix (TransGen Biotech, China). The genes involved in adipogenic differentiation (*PPAR $\gamma$* , *C/EBP $\alpha$* , *LPL*, *Fabp4*, *Adiponectin*, *Resistin* and *Leptin*), osteogenic differentiation (*Runx2*, *ALP*, *COL1A1*, *Osteocalcin* and *Osteopontin*) and Wnt/ $\beta$ -catenin signaling pathway (*Wnt1*, *Wnt10a*, *Wnt10b*, *Wnt5a*, *CTNNB1*, *Gsk3 $\beta$* , *Axin2* and *TGF7L2*) (Supplementary Table 1) were determined by the RT-PCR (No. AQ101-03, TransStart<sup>®</sup>Green qPCR SuperMix, TransGen Biotech, China), and 36B4 was the invariant internal control. The assays were performed in triplicate and normalized to the internal standard mRNA levels using the 2<sup>CT</sup> method.

### **Statistical analysis**

All statistical analyses were conducted using SPSS 21.0, with an  $\alpha$  level of 0.05 and effect coefficient of 0.90. All values were expressed as mean  $\pm$  standard deviation (or standard error), in which the Percent-Percent plot was chosen to determine data normality. The differences among all groups were tested and analyzed for the repeated measurement data based on whether the data were normally distributed (normal distribution: *t* test and ANOVA for continuous variables and  $\chi^2$  test for categorical variables; non-normal distribution: Kruskal Wallis H test).  $P < 0.05$  was considered to be statistically significant.

## **RESULTS**

### **Abnormal dietary calcium intake during pregnancy and lactation aggravated development of obesity among male offspring**

As shown in Supplementary Figure 2, there were no significant differences in body weight, daily dietary intake, related indexes of glucose (OGTT and ITT) and lipid (TC and TG), and bone Ca/P among the maternal DC, LC, NC and HC groups ( $P > 0.05$ ), with lower bone calcium and phosphorus in the DC and LC groups than in the NC group and higher levels in the HC group ( $P < 0.05$ ), which all proved that the animal model was successful.

**Table 2 Effects of maternal different calcium intake on cell cycle and purity of bone marrow mesenchymal stem cells among male offspring**

Indicators (%)	NC group (n = 9)	DC group (n = 9)	LC group (n = 9)	HC group (n = 9)	$\chi^2$	P value
<b>Cell cycle</b>						
G0G1 Phase	91.27 ± 3.14	94.26 ± 2.09	95.08 ± 2.54	92.46 ± 5.01	1.774	0.939
G2M Phase	0.85 ± 0.0091	0.94 ± 0.011	1.05 ± 0.013	0.79 ± 0.0082		
S Phase	7.89 ± 0.81	4.81 ± 0.56	3.87 ± 0.75	6.75 ± 0.94		
<b>Flow cytometry</b>						
Sca-1 (+)	95.03 ± 4.85	98.72 ± 6.12	99.31 ± 5.48	95.49 ± 3.84	4.788	0.188
CD90 (+)	98.12 ± 7.15	99.15 ± 2.04	99.07 ± 8.12	99.31 ± 6.74	0.608	0.895
CD29 (+)	99.13 ± 6.48	99.90 ± 3.46	97.02 ± 8.07	99.62 ± 7.01	0.549	0.908
CD34 (-)	99.08 ± 7.42	96.67 ± 2.93	97.94 ± 1.48	97.29 ± 1.92	0.364	0.546
CD31 (-)	99.46 ± 6.82	99.42 ± 4.12	98.94 ± 1.38	99.26 ± 1.57	0.159	0.690
CD45 (-)	99.79 ± 1.08	99.83 ± 2.54	98.48 ± 1.12	98.65 ± 1.75	0.287	0.963
CD49d (-)	99.13 ± 6.21	98.16 ± 2.32	99.06 ± 2.98	98.11 ± 1.09	5.087	0.166

Four-week-old C57BL/6J female mice were respectively fed with deficient (DC, 0.05%), low (LC, 0.25%), normal (NC, 0.70%) and high (HC, 1.20%) calcium reproductive diets for 6 wk before mating and continued their diets throughout gestation and lactation. After weaning, male offspring at 7-day-old from DC, LC, NC and HC groups were chosen to determine the cell cycle and purity of bone marrow mesenchymal stem cells ( $1 \times 10^6$ ) by flow cytometry. All values were shown as mean  $\pm$  standard deviation ( $n = 9$ /group), and comparisons were made by  $\chi^2$  test. NC: Normal-calcium reproductive diet; DC: Deficient-calcium reproductive diet; LC: Low-calcium reproductive diet; HC: High-calcium reproductive diet.

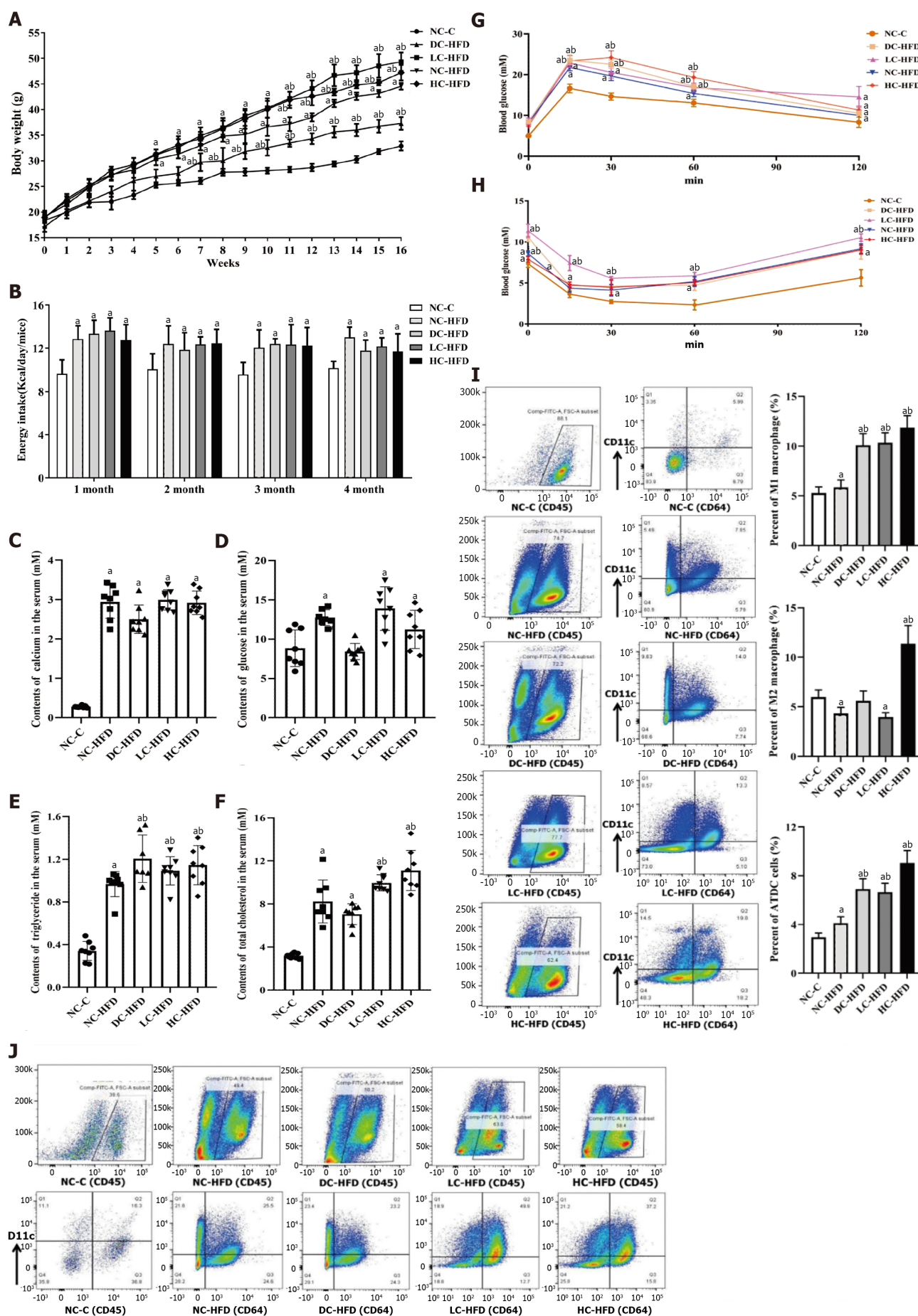
Among the male offspring, the body weight and mental state at weaning were not significantly different among the NC-C, DC-HFD, LC-HFD, NC-HFD and HC-HFD groups ( $P > 0.05$ , **Figure 1A**). During the HFD-induced adulthood, body weight (**Figure 1A**), energy intake (**Figure 1B**) and concentrations of serum metabolism-related indicators (TG, TC and glucose; **Figure 1C-F**) were all higher in the HFD groups (DC-HFD, LC-HFD, NC-HFD and HC-HFD) than in the NC-C group ( $P < 0.05$ ). Likewise, the circulating glucose responses to the glucose load, as indicated by OGTT and ITT (**Figure 1G and H**), showed that there were higher glucose levels in the HFD groups after intraperitoneal glucose administration ( $P < 0.05$ ). Compared with those in the NC-HFD group, maternal low (LC-HFD) and excess (HC-HFD) dietary calcium intake aggravated development of obesity, with significantly higher TC and TG ( $P < 0.05$ ). In contrast, body weight in the DC-HFD group was lower. However, the content of TG was higher than in the NC-HFD group ( $P < 0.05$ ).

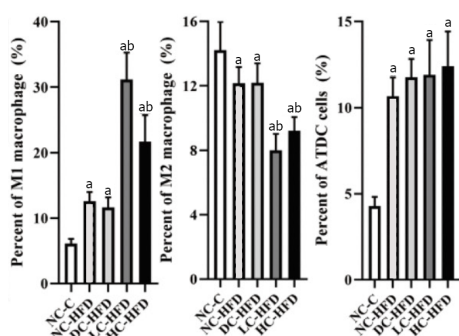
#### **Abnormal dietary calcium intake during pregnancy and lactation caused disorder of infiltration and percentages of immune cells in adipose tissues among male offspring**

The percentages of M1 macrophages, M2 macrophages and ATDC cells were demonstrated in eWAT (**Figure 1I**) and iWAT (**Figure 1J**). The percentages of M1 macrophages and ATDCs in eWAT were higher, and M2 macrophages were lower in the obese (DC-HFD, LC-HFD, NC-HFD and/or HC-HFD) groups than in the NC-C group ( $P < 0.05$ ). Further comparison among all the HFD groups showed that the percentages of M1 macrophages and ATDCs were increased in the DC-HFD, LC-HFD and HC-HFD groups when compared with the NC-HFD group ( $P < 0.05$ ). In iWAT, the percentages of M1 macrophages and ATDCs were significant higher, and M2 macrophages were lower in the HFD groups than in the NC-C group ( $P < 0.05$ ). Compared with the NC-HFD group, maternal low (LC-HFD) and high (HC-HFD) dietary calcium intake aggravated disorder of M1 and M2 macrophages ( $P < 0.05$ ), which was not found in the DC-HFD group ( $P > 0.05$ ).

#### **Abnormal dietary calcium intake during pregnancy and lactation affected weight and morphology of adipose tissues among male offspring**

The weights of eWAT, iWAT, KAT and BAT (**Figure 2A**) and adipose tissue weight/body weight (**Figure 2B**) were all higher in the four HFD groups than in the NC-C group ( $P < 0.05$ ). Compared the NC-HFD group, maternal deficient (DC-HFD, eWAT and eWAT/body weight), low (LC-HFD, eWAT, KAT, BAT, eWAT/body weight, KAT/body weight and BAT/body weight) and excess (HC-HFD, BAT and BAT/body weight) dietary calcium intake groups showed increased weights of eWAT, iWAT, KAT and/or BAT, and related adipose tissue weight/body weight ( $P < 0.05$ ). Compared with the NC-C group, the differentiation of adipocytes (number and diameter) in eWAT (**Figure 2C and D**), iWAT





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**Figure 1** Abnormal dietary calcium intake during pregnancy and lactation aggravated development of obesity among male offspring. A: Body weight; B: Energy intake; C: Contents of serum calcium; D: Contents of serum glucose; E: Contents of serum triglyceride; F: Contents of serum total cholesterol; G: Oral glucose tolerance test; H: Insulin glucose tolerance test; I and J: The infiltration and percentages of M1 macrophages, M2 macrophages and ATDC cells in the epididymal white adipose tissue and inguinal white adipose tissue. All pooled data was represented as mean  $\pm$  standard error ( $n = 10/\text{group}$ ). One-way analysis of variance was performed to compare the differences among the above four groups, and then Student–Newman–Keuls test was used to determine the differences between each two groups. Compared to the NC-C group, <sup>a</sup> $P < 0.05$ . Compared to the NC-HFD group, <sup>b</sup> $P < 0.05$ . NC-C: Normal-calcium reproductive diet and normal-fat diet after weaning; NC-HFD: Normal-calcium reproductive diet and high-fat-diet (HFD) after weaning; DC-HFD: Deficient-calcium reproductive diet and HFD after weaning; LC-HFD: Low-calcium reproductive diet and HFD after weaning; HC-HFD: High-calcium diet and HFD after weaning; eWAT: Epididymal white adipose tissue; iWAT: Inguinal white adipose tissue.

(Figure 2C and E), KAT (Figure 2C and F) and BAT (Figure 2C and G), was more prominent in the DC-HFD, LC-HFD, NC-HFD and HC-HFD groups ( $P < 0.05$ ). Compared with the NC-HFD group, maternal abnormal dietary calcium intake (DC-HFD, LC-HFD and HC-HFD groups) aggravated disorder of proliferation and differentiation of adipocytes in eWAT, iWAT and BAT among male offspring, with larger adipocytes ( $P < 0.05$ ).

#### Abnormal dietary calcium intake during pregnancy and lactation regulated target gene expression in adipose tissues among male offspring

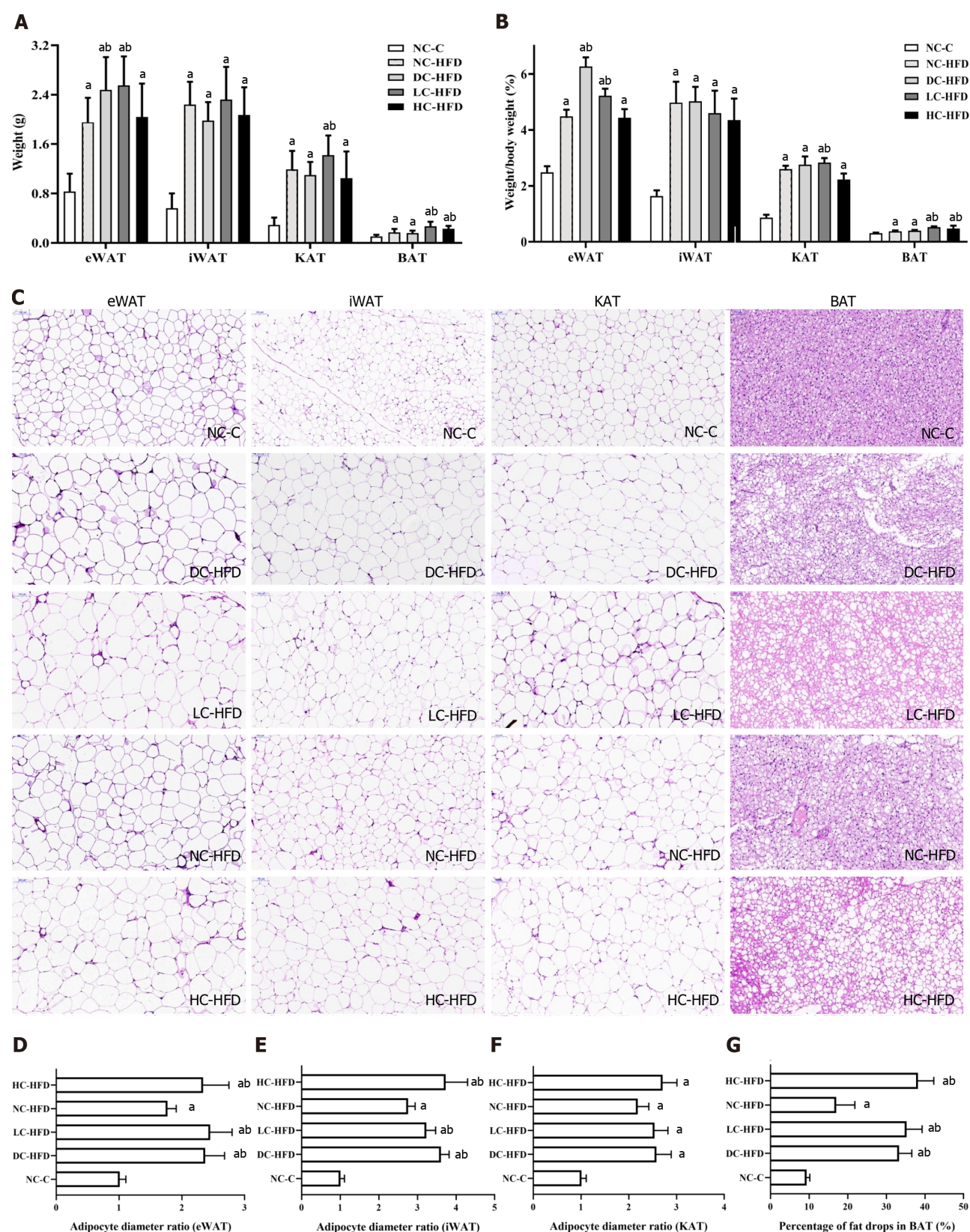
As shown in Figure 3, compared with the NC-HFD group, gene expression related to adipogenic differentiation (*PPAR $\gamma$* , *C/EBP $\alpha$* , *LPL*, *Fabp4*, *Adiponectin*, *Resistin* and *Leptin*) and Wnt/ $\beta$ -catenin signaling pathway (*Wnt1*, *Wnt10a*, *Wnt10b*, *Wnt5a*, *CTNNB1*, *Gsk3 $\beta$* , *Axin2* and *TGF7L2*) in eWAT, iWAT, KAT and BAT among the DC-HFD, LC-HFD and HC-HFD groups was more disordered.

Exactly, in eWAT (Figure 3A and B), compared with the NC-HFD group, there were higher expressions of *PPAR $\gamma$* , *Adiponectin* and *Wnt5a*, and lower expressions of *C/EBP $\alpha$* , *CTNNB1* and *TCF7L2* in the DC-HFD group ( $P < 0.05$ ), and higher expressions of *C/EBP $\alpha$* , *Fabp4*, *LPL*, *Adiponectin* and *Leptin*, and lower expressions of *Resist*, *CTNNB1* and *TCF7L2* in the LC-HFD group ( $P < 0.05$ ). Higher expressions of *PPAR $\gamma$* , *C/EBP $\alpha$* , *LPL*, *Fabp4*, *Adiponectin*, *Leptin* and *Wnt5a*, and lower expressions of *Wnt1*, *CTNNB1* and *TCF7L2* were demonstrated in the HC-HFD group ( $P < 0.05$ ). In iWAT (Figure 3C and D), expressions of *PPAR $\gamma$* , *Adiponectin*, *Resistin* and *Leptin* were higher in the DC-HFD, LC-HFD, and HC-HFD groups (with higher *Fabp4* in the HC-HFD) than in the NC-HFD group ( $P < 0.05$ ), with significantly lower expressions of *C/EBP $\alpha$* , *Wnt1*, *Wnt10a*, *Wnt10b*, *Wnt5a*, *CTNNB1* and *Gsk3 $\beta$*  in the DC-HFD group ( $P < 0.05$ ), lower expressions of *C/EBP $\alpha$* , *Wnt1*, *Wnt10a*, *Axin2* and *TCF7L2* in the LC-HFD group ( $P < 0.05$ ), and lower expressions of *CTNNB1* and *TCF7L2* in the HC-HFD group ( $P < 0.05$ ). As shown in KAT among the four HFD groups (Figure 3E and F), expressions of *PPAR $\gamma$* , *LPL*, *Wnt10b* and *Gsk3 $\beta$*  were higher, with the significantly low levels of *Wnt5a*, *CTNNB1* and *Axin2* in the DC-HFD (accompanied with higher expressions of *Adiponectin* and *Resistin*), LC-HFD (higher expressions of *Adiponectin*, with lower expressions of *C/EBP $\alpha$*  and *Resistin*), and HC-HFD (lower expressions of *Wnt10a* and *Resistin*) groups than in the NC-HFD group ( $P < 0.05$ ). In BAT (Figure 3G and H), expressions of *C/EBP $\alpha$* , *LPL*, *Adiponectin*, *Resistin*, *Leptin* and *TGF7L2* were higher, with the significantly lower expressions of *Wnt10a*, *Wnt5a* and *CTNNB1* in the DC-HFD (lower expressions of *Wnt1*), LC-HFD (higher *Wnt10b* and lower *Gsk3 $\beta$*  expressions), and HC-HFD groups (lower expressions of *Wnt1* and *Gsk3 $\beta$* ) than in the NC-HFD group ( $P < 0.05$ ).

#### Effects of dietary calcium intake during pregnancy and lactation on adipogenic and osteogenic differentiation potential of BMSCs

The morphology of BMSCs at P0 (Figure 4A) and P3 generations (Figure 4B) was similar in the DC, LC, NC and HC groups, with no significant differences in the pluripotent stem cells (G0/G1 phase) and purity of BMSCs (Sca-1+, CD90+, CD29+, CD34, CD31, CD45 and CD49d) in the P3 generation ( $P > 0.05$ , Table 2 and Supplementary Figure 1).



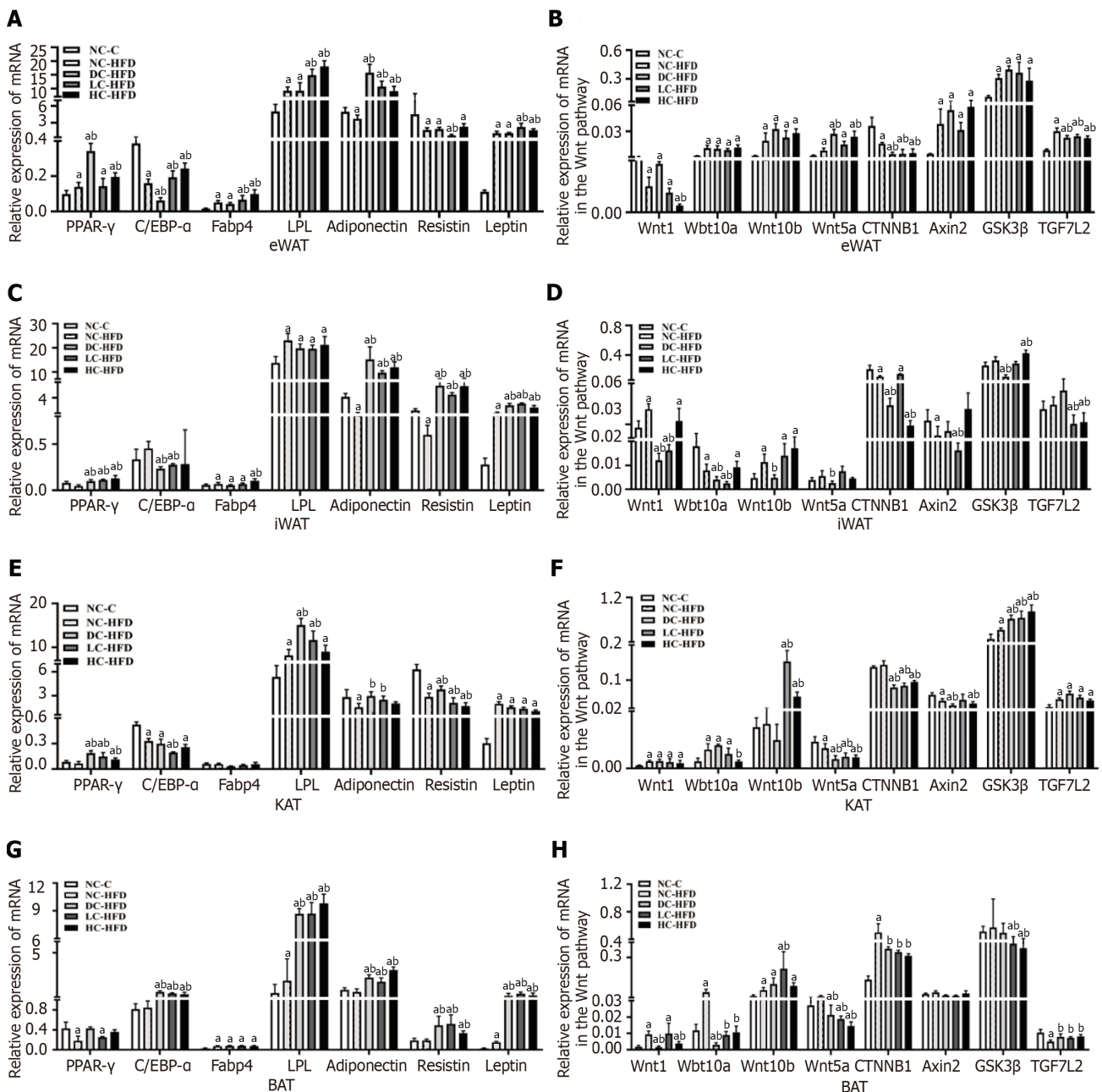


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**Figure 2** Abnormal dietary calcium intake during pregnancy and lactation affected the weight and morphology of adipose tissues among male offspring. A: Weights of the eWAT, iWAT, KAT and BAT; B: Percentage of eWAT, iWAT, KAT and BAT in body weight; C–G: Morphology of adipocytes in eWAT, iWAT, KAT and BAT by hematoxylin and eosin staining. All pooled data was represented as mean  $\pm$  standard error ( $n = 10/\text{group}$ ). One-way analysis of variance (ANOVA) was performed to compare the differences among the above four groups, and then Student–Newman–Keuls was involved to determine the differences between each two groups. Compared to the NC-C group, <sup>a</sup> $P < 0.05$ . Compared to the NC-HFD group, <sup>b</sup> $P < 0.05$ . NC-C: Normal-calcium reproductive diet and normal-fat diet after weaning; NC-HFD: Normal-calcium reproductive diet and high-fat-diet (HFD) after weaning; DC-HFD: Deficient-calcium reproductive diet and HFD after weaning; LC-HFD: Low-calcium reproductive diet and HFD after weaning; HC-HFD: High-calcium reproductive diet and HFD after weaning; eWAT:



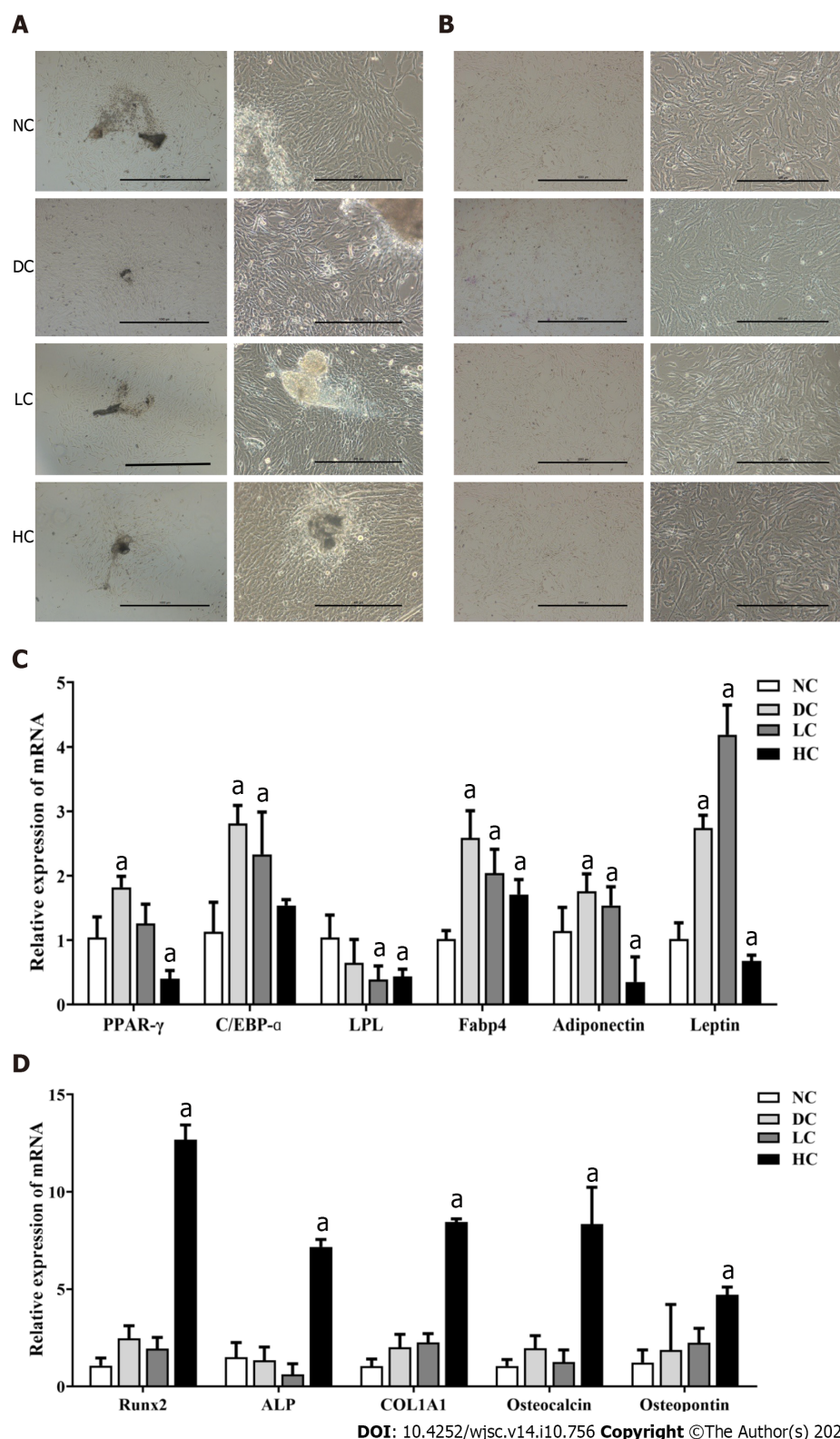
Epididymal white adipose tissue; iWAT: Inguinal white adipose tissue; KAT: Kidney adipose tissue; BAT: Brown adipose tissue.



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**Figure 3** Effects of different dietary calcium intake during pregnancy and lactation on expression of target genes for adipogenic differentiation and Wnt/β-catenin signaling pathway in adipose tissues among male offspring. A, C, E and G: Expression of genes related to adipogenic differentiation in eWAT, iWAT, KAT and BAT; B, D, F and H: Expression of genes related to the Wnt/β-catenin signaling pathway in eWAT, iWAT, KAT and BAT. All data presented as mean ± standard error ( $n = 10/\text{group}$ ). One-way analysis of variance was performed to compare the differences among the above four groups, and then Student-Newman-Keuls test was used to determine the differences between each two groups. Compared to the NC-C group, <sup>a</sup> $P < 0.05$ . Compared to the NC-HFD group, <sup>b</sup> $P < 0.05$ . NC-C: Normal-calcium reproductive diet and normal-fat diet after weaning; NC-HFD: Normal-calcium reproductive diet and high-fat-diet (HFD) after weaning; DC-HFD: Deficient-calcium reproductive diet and HFD after weaning; LC-HFD: Low-calcium reproductive diet and HFD after weaning; HC-HFD: High-calcium reproductive diet and HFD after weaning; eWAT: Epididymal white adipose tissue; iWAT: Inguinal white adipose tissue; KAT: Kidney adipose tissue; BAT: Brown adipose tissue.

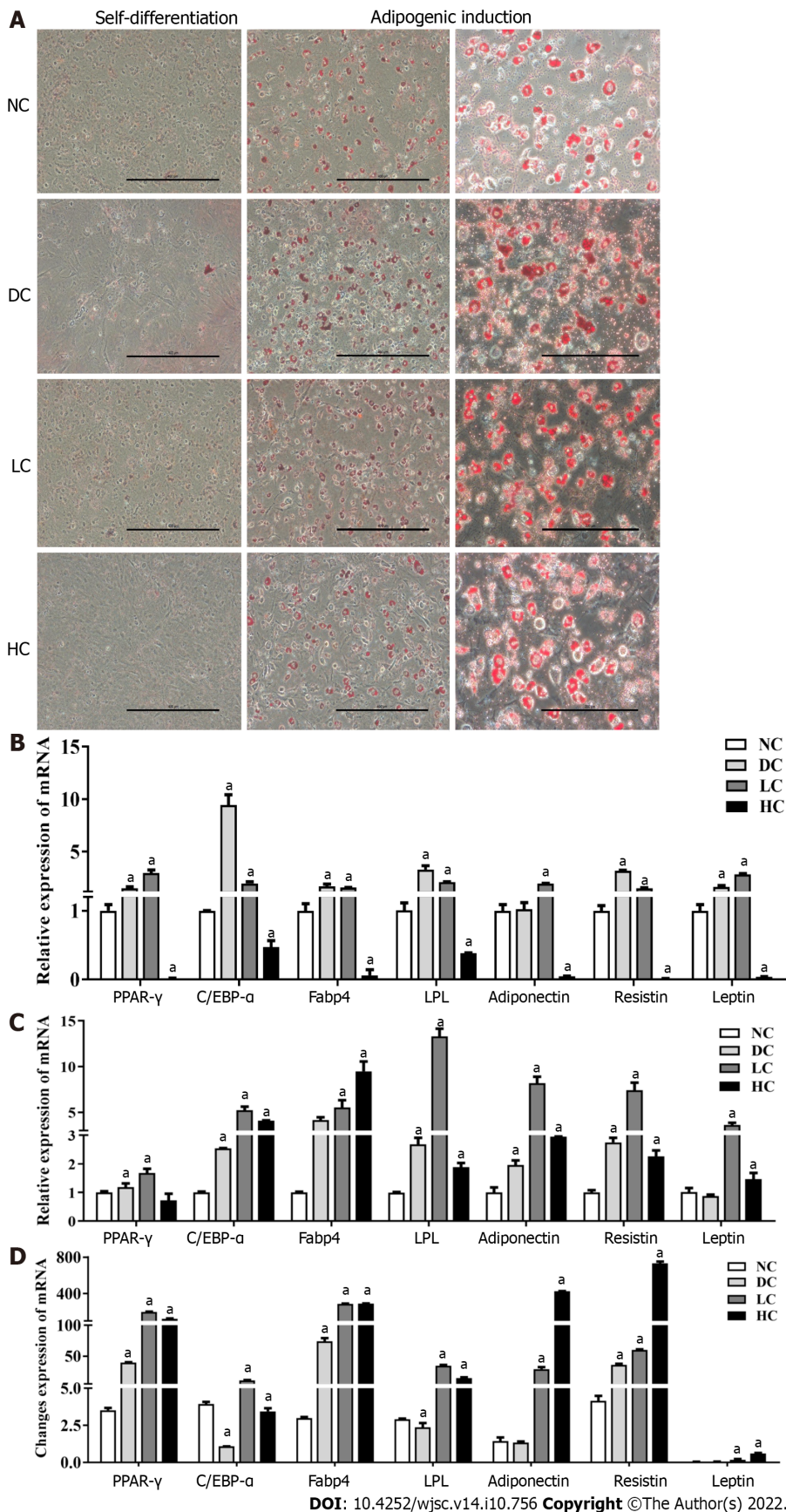
In the P3 generation of BMSCs, compared with the NC group, maternal deficient (DC group) and low (LC group) dietary calcium intake promoted adipogenic differentiation potential of BMSCs, with higher levels of *PPARγ*, *C/EBPα*, *Fabp4*, *Adiponectin* and *Leptin* (Figure 4C,  $P < 0.05$ ). Maternal excess dietary calcium intake (HC group) induced osteogenic differentiation and inhibited adipogenic differentiation of BMSCs, with higher levels of *Runx2*, *ALP*, *COL1A1*, *Osteocalcin* and *Osteopontin*, and lower levels of



**Figure 4 Effects of maternal dietary calcium intake during pregnancy and lactation on multi-differentiation potential of bone marrow mesenchymal stem cells among male offspring.** A: Morphology of P0 BMSCs; B: Morphology of P3 BMSCs; C: Expression of genes related to adipogenic differentiation; D: Expression of genes related to osteogenic differentiation. All data presented as mean  $\pm$  standard error ( $n = 9$ /group). One-way analysis of variance was performed to compare the differences among the above four groups, and then Student–Newman–Keuls test was used to determine the differences between each two groups. Compared to the NC group, <sup>a</sup> $P < 0.05$ . NC: Normal-calcium reproductive diet; DC: Deficient-calcium reproductive diet; LC: Low-calcium reproductive diet; HC: High-calcium reproductive diet; BMSCs: Bone marrow mesenchymal stem cells.

PPAR $\gamma$ , LPL, Adiponectin and Leptin (Figure 4C and D,  $P < 0.05$ ).

Under the adipogenic induction (Figure 5), compared with the NC group, maternal abnormal dietary calcium intake (DC, LC and HC groups) promoted adipogenic differentiation of BMSCs, with more lipid



**Figure 5** Effects of maternal dietary calcium intake during pregnancy and lactation on adipogenic differentiation potential of bone marrow mesenchymal stem cells among male offspring under agent induction. A: Morphology of BMSCs under adipogenic differentiation; B and C: Expression of target genes related to adipogenic differentiation under adipogenic induction and self-differentiation status; D: Ratio of gene expressions between adipogenic differentiation and self-differentiation. All data presented as mean  $\pm$  standard error ( $n = 9/\text{group}$ ). One-way analysis of variance was performed to compare



the differences among the above four groups, and then Student–Newman–Keuls test was used to determine the differences between each two groups. Compared to the NC group, <sup>a</sup> $P < 0.05$ . NC: Normal-calcium reproductive diet; DC: Deficient-calcium reproductive diet; LC: Low-calcium reproductive diet; HC: High-calcium reproductive diet; BMSCs: Bone marrow mesenchymal stem cells.

drops (Figure 5A) and higher expressions of *PPAR $\gamma$* , *C/EBP $\alpha$* , *Fabp4*, *LPL*, *Adiponectin*, *Resistin* and *Leptin* (Figure 5C and D). As with expressions of genes related to adipogenic differentiation in the P3 generation (Figure 4C), compared with that in the NC group, expressions of *PPAR $\gamma$* , *C/EBP $\alpha$* , *Fabp4*, *LPL*, *Adiponectin*, *Resistin* and *Leptin* in the DC and LC groups were higher under the self-differentiation status (Figure 5B), with lower expressions in the HC group. Under osteogenic induction (Figure 6), compared with the NC group, maternal excess dietary calcium intake (HC group) promoted osteogenic differentiation of BMSCs, with more calcium nodules (Figure 6A) and higher expressions of *Runx2*, *ALP*, *COL1A1* and *Osteocalcin* (Figure 6C and D), which was similar to that under the self-differentiation status (Figure 6B). Furthermore, compared with the NC group, the osteogenic differentiation potential of BMSCs was weaker in the DC and LC groups under osteogenic induction and self-differentiation status (Figure 6A, C and D), with lower expressions of *Runx2*, *ALP*, *COL1A1* and *Osteocalcin*.

### **Abnormal dietary calcium intake during pregnancy and lactation regulated gene expressions of BMSCs in the Wnt/ $\beta$ -catenin signaling pathway under different interventions**

In the P3 generation of BMSCs without induction, compared with the NC group, maternal deficient (DC group) and low (LC group) dietary calcium intake inhibited expressions of *Wnt1*, *Wnt10a*, *CTNNB1* and *Axin2* ( $P < 0.05$ ) (Figure 7), while there were significantly higher expressions of *Wnt10b*, *CTNNB1*, *Gsk3 $\beta$*  and *TGF7L2*, with lower expressions of *Wnt10a* and *Wnt5a* in the HC group ( $P < 0.05$ ) (Figure 7A).

Under adipogenic induction, compared with the NC group, maternal abnormal dietary calcium intake (DC, LC and HC groups) decreased expressions of *Wnt10a*, *Wnt10b*, *CTNNB1*, *Gsk3 $\beta$*  and *TGF7L2* to promote the adipogenic differentiation potential of BMSCs (Figure 7D,  $P < 0.05$ ), while under the self-differentiation status, maternal deficient (DC group) and low (LC group) dietary calcium intake inhibited expressions of *Wnt1*, *CTNNB1* and *Axin2* ( $P < 0.05$ ), while there were significantly higher levels of *Wnt1*, *Wnt10a*, *Wnt10b*, *Wnt5a*, *CTNNB1*, *Axin2*, *Gsk3 $\beta$*  and *TGF7L2* in the HC group compared with the NC group (Figure 7B,  $P < 0.05$ ), which was consist with the P3 BMSCs.

Under osteogenic induction, compared with the NC group, expressions of *Wnt1*, *Wnt5a* and *TGF7L2* were lower in the DC group ( $P < 0.05$ ), *Wnt1* expressions was lower in the LC group ( $P < 0.05$ ), and there were significantly higher levels of *Wnt10a*, *Wnt10b*, *CTNNB1*, *Axin2*, *Gsk3 $\beta$*  and *TGF7L2* in the HC group (Figure 7E,  $P < 0.05$ ). Under the self-differentiation status, maternal DC and LC intake inhibited expressions of *Wnt1*, *Wnt10a*, *Wnt5a*, *CTNNB1* and *TGF7L2* ( $P < 0.05$ ), while there were significantly higher expressions of *Wnt1*, *Wnt10a*, *Wnt10b* and *Axin2* in the HC group than in the NC group (Figure 7C,  $P < 0.05$ ).

## **DISCUSSION**

The correlation between the inappropriate consumption of nutrient and occurrence of obesity presents a greatest global public health problem, which needs more novel therapies[30]. A lot of researches have demonstrated that chronically deficient and excessive calcium exposure is as an important contributing factor to the development of obesity by controlling the *de novo* lipogenesis and lipolytic signals through regulating related gene expressions[31–33]. There is also compelling evidence that maternal calcium dysfunction directly affects fat synthesis and metabolism of their offspring[34–37]. In agreement with our findings using a mouse model that maternal inappropriate dietary calcium intake during pregnancy and lactation aggravated development of obesity by elevating cytosolic calcium, with more and larger adipocytes, and disorders of immune cells (M1 macrophages, M2 macrophages and ATDC cells) in the adipose tissues.

It is agreed that the development of obesity is driven by hypertrophy and hyperplasia of the adipocytes in the process of adipogenic differentiation to cause the expansion of fat depots[38]. Lineage-tracing models have shown that the numbers of adipocytes are primarily determined in early life and mostly stable through to adulthood for the remarkable hypertrophic potential of differentiated adipocytes with HFD induction[39–42]. The modulation of cytosolic calcium can regulate the early stage of adipocyte differentiation and thermogenic capacity of BAT in mice[43]. The propensity of adipogenesis to generate new adipocytes in different adipose tissues (eWAT, iWAT, KAT and BAT) highlights the unique characteristics of fat depots. Thus, we should discuss the roles of abnormal dietary calcium intake during pregnancy and lactation on the adipogenic differentiation potential in different adipose tissues among male offspring. This was consistent with our findings that the imbalance of dietary calcium intake in early life could affect the proliferation and differentiation of eWAT, iWAT and BAT, with higher weight of adipose tissue. However, the mechanism remains to be elucidated.

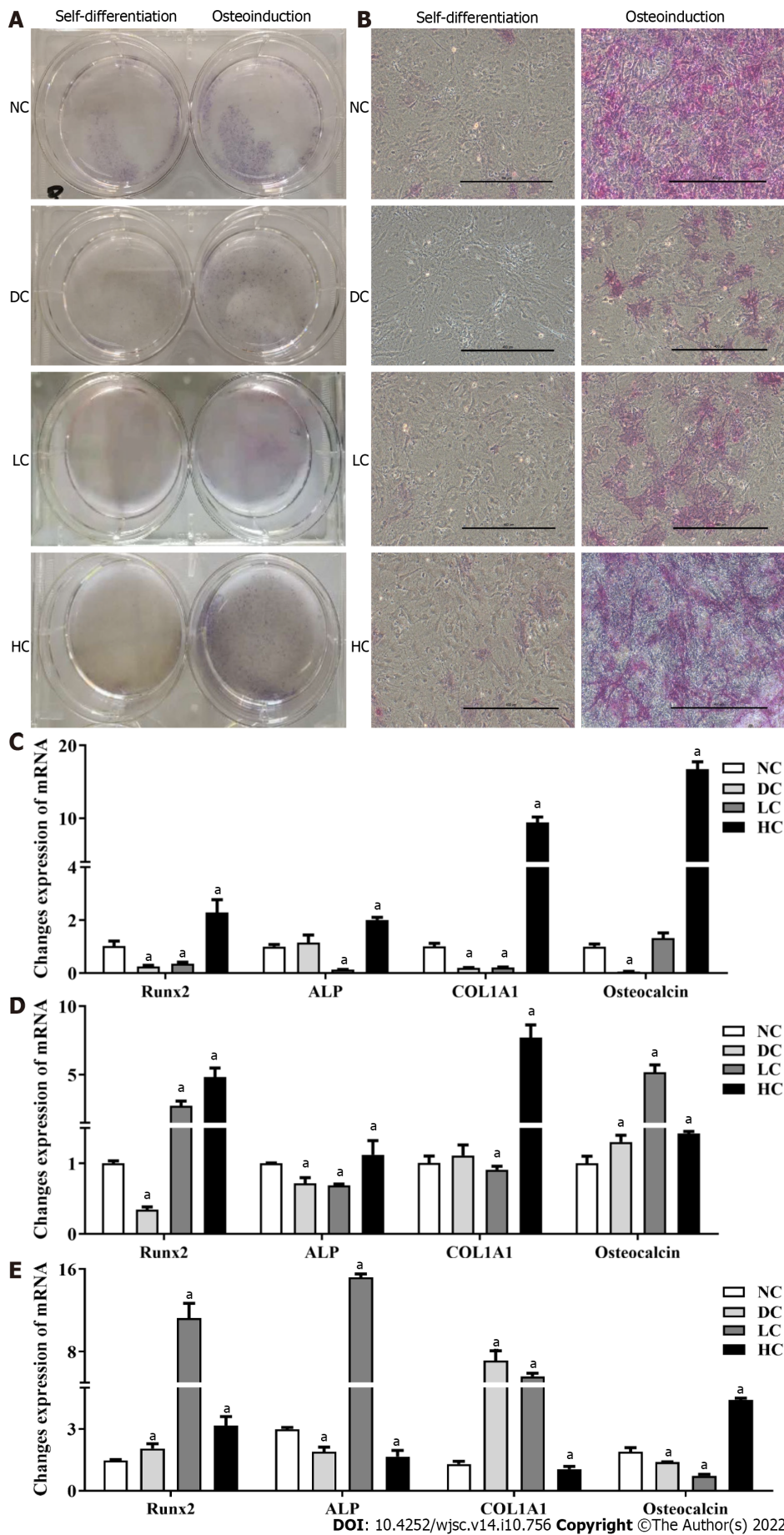


Figure 6 Effects of maternal dietary calcium intake during pregnancy and lactation on osteogenic differentiation potential of bone



**marrow mesenchymal stem cells among male offspring under agent induction.** A and B: Morphology of BMSCs under osteogenic differentiation; C and D: Expression of target genes on osteogenic differentiation under osteogenic induction and self-differentiation status; E: Ratio of gene expressions between osteogenic differentiation and self-differentiation. All data presented as mean  $\pm$  standard error ( $n = 9/\text{group}$ ). One-way analysis of variance was performed to compare the differences among the above four groups, and then Student–Newman–Keuls test was used to determine the differences between each two groups. Compared to the NC group,  $^aP < 0.05$ . NC: Normal-calcium reproductive diet; DC: Deficient-calcium reproductive diet; LC: Low-calcium reproductive diet; HC: High-calcium reproductive diet; BMSCs: Bone marrow mesenchymal stem cells.

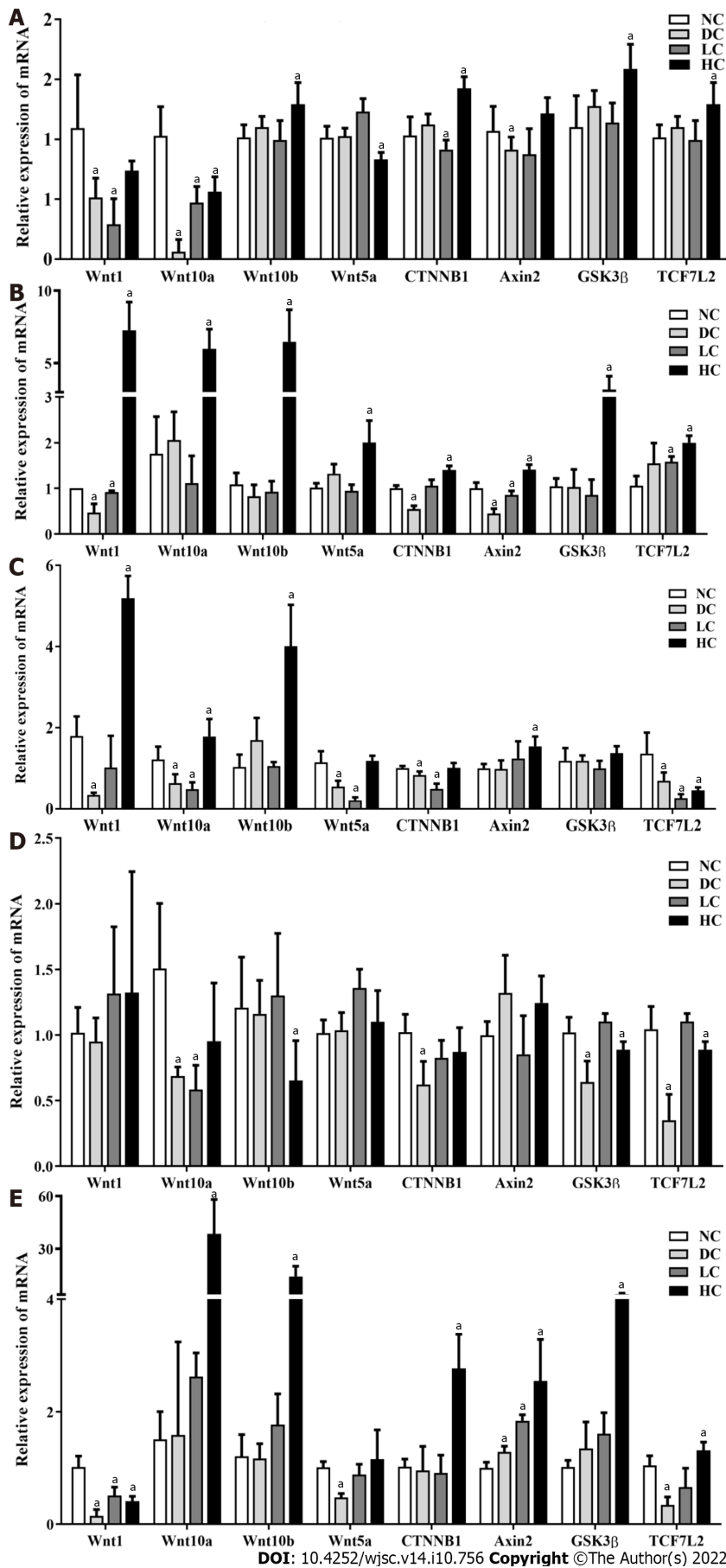
Lineage-tracing studies in animal models have suggested that there are two-step phases in the adipogenic differentiation, including specific preadipocyte formation (MSCs to preadipocytes) and terminal adipocyte maturation (preadipocytes to mature adipocytes), in which the committed preadipocytes from the pluripotent MSCs are activated by a number of critical transcription factors (*PPAR $\gamma$* , *C/EBP $\alpha$* , *C/EBP $\beta$*  and *FABP4*) and related extracellular signals. At the differentiation of mature adipocytes (second stage), they express all the biomarkers of early adipocyte differentiation as well as the peptide hormones, such as *Adiponectin*, *Resistin*, *Leptin*, *ATGL*, *LPL* and *Perilipin 1*[44,45]. All the above transcription factors are involved in the specific Wnt signaling pathway to affect adipogenic differentiation[46–48]. In our study, abnormal dietary calcium intake during pregnancy and lactation (DC-HFD, LC-HFD and HC-HFD) aggravated expressions of genes related to proliferation and differentiation of adipocytes and Wnt/ $\beta$ -catenin signaling pathway in eWAT, iWAT, KAT or BAT in the adulthood of their male offspring, which could more clearly explain the possible causes for the development of obesity. In the early stage, MSCs, as multipotential progenitor cells, are delicately balanced for their terminal adipo-osteogenic differentiation commitment[49–51]. It has also been reported that this decision process of MSCs is competing and reciprocal, and is precisely achieved by a variety of critical and external cues, including phytocannabinoids, conjugated linoleic acid, calcium, and chemical, physical and biological factors[52–58]. Many investigations *in vitro* had demonstrated that deficient calcium exposure inhibited osteogenesis[59–61]. Conversely, little was known about the effects of inappropriate dietary calcium intake during pregnancy and lactation on adipogenic differentiation, to aggravate the development of obesity in adulthood under HFD induction[55,62]. The major novel finding of our study was that maternal deficient and low dietary calcium intake aggravated the potential adipogenic differentiation and suppressed osteogenic differentiation of BMSCs. Maternal excess dietary calcium intake could play an opposing differentiation role without the exogenous stimuli. In response to reagent induction, both maternal deficient, low and excessive dietary calcium intake could polarize adipogenic differentiation and suppress osteogenic differentiation. All the above results were consistent with the results in the adult offspring with HFD induction.

Terminal differentiation of BMSCs is achieved through a coordinated and highly orchestrated program of triggering different signaling pathways, and activated by various transcription factors that guide the programming alterations of BMSCs to commit the lineage to cause the pathophysiological processes of obesity[63–66]. Thus, it is necessary that our research for screening out the roles of different calcium exposure in early life on the expressions of related transcription factors and signaling pathways could regulate both osteogenic and adipogenic differentiation of BMSCs. This proves that the imbalance of terminal adipo-osteogenic differentiation by abnormal calcium exposure in early life results from the above disorders of gene expressions and Wnt/ $\beta$ -catenin signaling pathway on the differentiation of BMSCs among male offspring.

There were some limitations to this study. Firstly, we required more complex and explicit procedures, including western blotting, to explore the related mechanisms more clearly. Secondly, our conclusions need to be verified in other MSCs and animal models to ensure their feasibility and effectiveness. Finally, the inconsistent results of maternal deficient calcium intake still need to be further discussed.

## CONCLUSION

Our results suggest that abnormal dietary calcium intake during gestation and lactation aggravates the development of obesity by programming the adipogenic differentiation potential of BMSCs among male offspring, which is related to the significantly different expressions of target genes for adipogenic differentiation on the Wnt/ $\beta$ -catenin signaling pathway to aggravate dietary-induced obesity in the adulthood. Maternal deficient calcium exposure can inhibit the osteogenic differentiation to cause low body weight. So the importance of this study is that the prevention of adulthood obesity could be moved forward to the appropriate calcium intake in the neonatal period, even the formation of maternal germ cells and fertilized egg.



**Figure 7** Effects of maternal different dietary calcium intake on gene expressions of bone marrow mesenchymal stem cells at the Wnt/β-catenin signaling pathway under different interventions among male offspring. A: P3 BMSCs; B and D: Adipogenic induction and self-differentiation

status of BMSCs; C and E: Osteogenic induction and self-differentiation status of BMSCs. All data presented as mean  $\pm$  standard error ( $n = 9/\text{group}$ ). One-way analysis of variance was performed to compare the differences among the above four groups, and then Student–Newman–Keuls test was used to determine the differences between each two groups. Compared to the NC group,  $^aP < 0.05$ . NC: Normal-calcium reproductive diet; DC: Deficient-calcium reproductive diet; LC: Low-calcium reproductive diet; HC: High-calcium reproductive diet; BMSCs: Bone marrow mesenchymal stem cells.

## ARTICLE HIGHLIGHTS

### Research background

Obesity is characterized by the hypertrophy and hyperplasia of adipocytes, in which the commitment from bone mesenchymal stem cells (BMSCs) to preadipocytes is the important process for their hyperplasia. Our previous study showed that dietary insufficient and excessive calcium intake during pregnancy and lactation increased the body weight of offspring, using a high-fat-diet-induced obese mouse model and epidemiological cohorts. However, whether maternal inappropriate dietary calcium intake could affect the adipogenic differentiation potential of MSCs is still unclear.

### Research motivation

This study was designed to investigate the effects of abnormal dietary calcium intake during gestation and lactation on the multi-differentiation potential of BMSCs among male offspring, and explore the possible role of the Wnt/ $\beta$ -catenin signaling pathway, which might aggravate the development of obesity, with more excessive lipid accumulation in adulthood.

### Research objectives

We presented the possibility that abnormal dietary calcium intake during pregnancy and lactation could derive hyperplastic adipogenesis from BMSCs by regulating target gene expressions profiles through the fetus to adulthood among their male offspring.

### Research methods

Four-week-old female C57BL/6N mice were fed by deficient, low, normal and excessive calcium reproductive diets throughout pregnancy and lactation. The BMSCs were obtained from 7-day-old male offspring to measure their adipogenic differentiation potential through the Wnt/ $\beta$ -catenin signaling pathway. The other weaning male pups were fed a high-fat diet for 16 wk along with a normal-fat diet as the control. Serum was collected for biochemical analysis. Adipose tissues were excised for histological examination, immunohistochemistry, determining the proportions of immune cells by flow cytometry, and gene expressions related to adipogenic differentiation and Wnt/ $\beta$ -catenin signaling pathway by real-time reverse transcription polymerase chain reaction.

### Research results

Maternal deficient, low and excess dietary calcium intake aggravated dietary-induced obesity with more/larger adipocytes and higher serum metabolism indicators, along with disordered expressions of genes related to adipogenic differentiation (*PPAR $\gamma$* , *C/EBP $\alpha$* , *Fabp4*, *LPL*, *Adiponectin*, *Resistin* and *Leptin*) in the adipose tissues among the male offspring. We also showed significantly different expressions of similarly specific genes in BMSCs to successfully polarize adipogenic differentiation and suppress osteogenic differentiation *in vivo* and *in vitro*, respectively. The related mechanistic insights were gained to worsen this adipogenic differentiation through the Wnt/ $\beta$ -catenin signaling pathway in the BMSCs and adult adipose tissues.

### Research conclusions

Abnormal dietary calcium intake during pregnancy and lactation might program the adipogenic differentiation potential of BMSCs among male offspring, which was related to the significantly different expressions of target genes in the Wnt/ $\beta$ -catenin signaling pathway to preserve more adipocytes to aggravate dietary-induced obesity in adulthood.

### Research perspectives

The importance of this study is that the prevention of adulthood obesity could be moved forward to the appropriate calcium intake in the neonatal period, even the formation of maternal germ cells and fertilized egg.

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## FOOTNOTES

**Author contributions:** Li P and Wang Y designed the study, performed the data analysis and wrote the manuscript; Li P, Chen XY and Tang TT were responsible for all the animal procedures and experiments; Liu YL and Liu WJ collected the bone mesenchymal stem cells; Qi KM and Zhang Y supervised the final manuscript.

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**Institutional animal care and use committee statement:** All animal studies were approved and conducted in accordance with the Beijing Academy of Military Medical Sciences Guide for the Care and Usage Committee of Laboratory Animals. Meanwhile, the animal care and use committee statement used in this study was approved on the Ethics of Animal Experiments of Academy of Military Medical Sciences in China, No. IACUC-DWZX-2019-704.

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**Data sharing statement:** The data and materials that support the findings of this study are available from the corresponding author upon the reasonable requests.

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