

Hyperhomocysteinemia inhibits tibial fracture healing in rats through PI3K/AKT signaling pathway

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Abstract. To explore the influence of hyperhomocysteinemia (hHcys) on the tibial fracture healing in rats and its effect on the phosphatidylinositol 3-hydroxy kinase (PI3K)/protein kinase B (AKT) signaling pathway. A total of 36 Sprague-Dawley rats were randomly divided into sham group (n=12), tibial fracture group (n=12) and hHcys + fracture group (n=12). The rats in tibial fracture group underwent the tibial fracture surgery, while the model of tibial fracture and hHcys was established in hHcys + fracture group. The level of plasma homocysteine (Hcy) in each group was analyzed using the full-automatic biochemical analyzer, the fracture stress biomechanical measurement was performed, and the ultimate bending strength and torque were calculated. Moreover, the protein expressions of PI3K and phosphorylated (p)-AKT in tibial tissues were detected using western blotting, the messenger ribonucleic acid (mRNA) levels of Bcl-2 associated X protein (Bax) and caspase-3 were detected using quantitative polymerase chain reaction (qPCR), the apoptosis was detected via terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling (TUNEL) staining, and the expressions of inflammatory factors were detected via immunohistochemistry. Compared with sham group, tibial fracture group and hHcys + fracture group had a significantly increased level of plasma Hcy, significantly decreased ultimate bending strength and torque, obviously decreased relative protein expressions of PI3K and p-AKT, increased mRNA levels of Bax and caspase-3 and an increased expression of pro-inflammatory factor tumor necrosis factor- α (TNF- α). Compared with tibial fracture group, hHcys + fracture group had a higher level of plasma Hcy, lower ultimate bending strength and torque, lower relative protein expressions of PI3K and p-AKT, higher mRNA levels of Bax and caspase-3,

a higher apoptosis rate and a higher expression of TNF- α . hHcys blocks the downstream apoptotic signal transduction, promotes apoptosis and inflammatory response, and affects fracture healing through affecting the PI3K/AKT signaling pathway.

Introduction

Tibial fracture is a clinically common type of long tubular bone fracture accounting for 13.7% of whole body fractures (1). The tibial fracture healing is a very complex process of injury repair, which can be affected by many factors. Hyperhomocysteinemia (hHcys) is an important risk factor for cardiovascular disease and thrombotic disease (2). The clinical manifestations of hHcys include developmental retardation and skeletal anomalies, and high-level homocysteine (Hcy) will damage the function of osteocytes regulating bone remodeling, thereby resulting in fractures. Hcy, through increasing osteoclast activity, can regulate bone remodeling (3), induce apoptosis of bone marrow stromal cells (4), osteocytes and osteoblasts (5,6), and inhibit osteoblast differentiation (7).

The phosphatidylinositol 3-hydroxy kinase (PI3K)/protein kinase B (AKT) signal transduction pathway plays an important role in post-traumatic repair and healing, which is a regulatory pathway for cell growth. Studies have proved that this pathway is essential for the differentiation of osteoprogenitor cells (8,9). Tetrahydroxystilbene glycoside (TSG) can facilitate the differentiation of MC3T3-E1 cells through activating the PI3K/AKT signal transduction. The downregulation of PTEN, a negative feedback regulator of PI3K/AKT in osteoblasts, can activate the PI3K/AKT signaling pathway, thus affecting osteocyte differentiation (10).

In this study, therefore, the effect of hHcys on the tibial fracture healing in rats was observed, and the role of PI3K/AKT signaling pathway in tibial fracture was explored, hoping to provide a theoretical basis for the diagnosis and treatment of tibial fracture.

Materials and methods

Animal experiments and grouping. A total of 36 specific pathogen-free male Sprague-Dawley rats weighing 200-240 g were adaptively fed in an environment in line with animal

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ethical requirements for 1 week before surgery, and they had free access to food and water. They were deprived of food, but not water, for 12 h before surgery, and divided into sham group (n=12), tibial fracture group (n=12) and hHcys + fracture group (n=12) using a random number table. This study was approved by the Animal Ethics Committee of Soochow University Animal Center (Suzhou, China).

Modeling. i) Establishment of tibial fracture model (11): After intraperitoneal anesthesia with pentobarbital sodium, the left lower limb was depilated, and the fascia and muscle were separated to expose the tibia. Then the fracture model was established in the middle tibia, and fixed intramedullarily using the Kirschner wire (1 mm in diameter). ii) Establishment of hHcys model (12): After establishment of tibial fracture model, the rats in hHcys + fracture group were fed with L-methionine for 4 weeks. iii) In sham group, the tibia was exposed only without establishing the tibial fracture model, and the rats were fed with normal diet.

Main instruments and equipment. AG-1S electronic universal mechanical testing machine (Shimadzu), PI3K, p-AKT, Bax, caspase-3 and tumor necrosis factor- α (TNF- α) primary antibodies (Abcam), terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling (TUNEL) assay kit (Shanghai Beyotime Biotechnology), quantitative polymerase chain reaction (qPCR) kit (Vazyme), immunohistochemistry kit (Maxim), bicinchoninic acid (BCA) protein quantification kit (Shanghai Beyotime Biotechnology), fluorescence qPCR instrument (ABI 7500; Applied Biosystems; Thermo Fisher Scientific, Inc.), Image-Lab image analysis system (Bio-Rad Laboratories), and Leica DM4000B LED microscope (Leica Microsystems GmbH).

Detection of plasma Hcy concentration. The venous blood was drawn, placed in the test tube containing ethylenediaminetetraacetic acid (EDTA) and coagulant, and centrifuged at 2,000 x g at 4°C for 5 min to separate the plasma and serum. Then the plasma Hcy concentration was measured using a full-automatic biochemical analyzer. Plasma (100-150 μ l) was used for each measurement, and each measurement was repeated for 3 times.

Biomechanical measurement of fracture stress. After the rats were sacrificed, the tibial specimens were taken for the biomechanical three-point bending test at a loading rate of 0.01 mm/sec and span of 15 mm. The mechanical properties of fractured tibia were analyzed using a clinical biomechanical three-point bending strain detector, and the load-displacement relation curve and stress-strain relation curve were plotted, based on which the ultimate bending strength and torque were read and calculated.

Determination of protein expressions of PI3K and p-AKT in tibial tissues via western blotting. The tibial tissues were taken, fully lysed with cell lysis buffer in an ultrasonic homogenizer and centrifuged at 3,500 x g at 4°C for 10 min. After the supernatant was discarded, the protein samples were obtained, and the total protein concentration was determined using the BCA protein concentration kit. The protein was separated via gel electrophoresis, transferred onto a polyvinylidene

fluoride (PVDF) membranes (Roche Diagnostics), sealed at room temperature and incubated with PI3K and p-AKT primary antibodies at 37°C for 1 h, with β -actin antibody as an internal reference. After the membrane was washed, the protein was incubated again with horse radish peroxidase (HRP)-labeled secondary antibodies, and the membrane was washed again. Finally, the image was developed using the electrochemiluminescence (ECL) kit in a darkroom.

Determination of messenger ribonucleic acid (mRNA) levels of Bax and caspase-3 via qPCR. The total RNA was extracted from tibial tissues using the TRIzol method (Invitrogen; Thermo Fisher Scientific, Inc.), and its concentration was measured using a spectrophotometer. Then the total RNA was reversely transcribed into complementary deoxyribose nucleic acid (cDNA) and quantitatively amplified using the reverse transcription kit and qPCR kit. The amplification conditions are as follows: pre-denaturation at 95°C for 1 min, denaturation at 95°C for 5 sec, and annealing/extension at 58°C for 15 sec, a total of 40 cycles. Primer sequences of glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (internal reference gene), forward, 5'-GGTGTGAGTATGTCGTGGA-3' and reverse, 5'-TGCTGA CAATCTTGAGGGAG-3'. Primer sequences of caspase-3, forward, 5'-GACCCGGTGCCTCAGGATGC-3' and reverse, 5'-GTGGCATGAGCTCTTGATAATG-3'. Primer sequences of Bax forward, 5'-CAGAGGCGGGGGATGATTG-3' and reverse, 5'-TGTCCAGCCCATGATGGTTC-3'. The relative mRNA expression level was calculated using the $2^{-\Delta C_t}$ formula.

Detection of apoptosis using TUNEL staining. The tissue sections were deparaffinized with xylene, dehydrated with gradient alcohol, and subjected to citrate antigen retrieval, followed by staining using the TUNEL staining kit. Then the sections were sealed with anti-fluorescence quenching blocking buffer and observed under a fluorescence microscope, and the number of positive cells and apoptosis rate were calculated based on apoptosis rate = number of positive cells/total number of cells x 100%.

Detection of expressions of inflammatory factors through immunohistochemistry. The tibial tissues were taken in each group, washed with phosphate-buffered saline (PBS), and decalcified with 10% EDTA decalcifying solution until softening of tissues, followed by paraffin embedding into 5 μ m-thick coronal sections. Then the sections were deparaffinized with xylene, dehydrated with gradient alcohol, and subjected to citrate antigen retrieval, followed by staining using the immunohistochemistry kit. The sections were incubated with 3% H₂O₂ at room temperature for 10 min, washed, sealed with 5% goat serum at room temperature for 10 min, and dropwise added with TNF- α primary antibody in a wet box at 4°C overnight. After rewarming at 37°C, the sections were incubated again with biotinylated secondary antibody at room temperature for 10 min, followed by color development using the diaminobenzidine (DAB) developer and counterstaining with hematoxylin. Finally, the absorbance was analyzed using the Motic Med 6.0 pathologic image analysis system.

Statistical analysis. Statistical Product and Service Solutions (SPSS) 24.0 (IBM Corp.) software was used for the data

Table I. Comparison of fracture biomechanics in each group (mean \pm SD).

Group	n	Ultimate bending strength (N)	Torque (N·mm)
Sham group	6	116.78 \pm 12.65	357.49 \pm 7.65
Tibial fracture group	6	77.23 \pm 6.77 ^a	216.82 \pm 17.71 ^a
hHcys + fracture group	6	45.84 \pm 4.32 ^{a,b}	124.15 \pm 11.84 ^{a,b}

^aP<0.05 vs. sham group; ^bP<0.05 vs. tibial fracture group. hHcys, hyperhomocysteinemia.

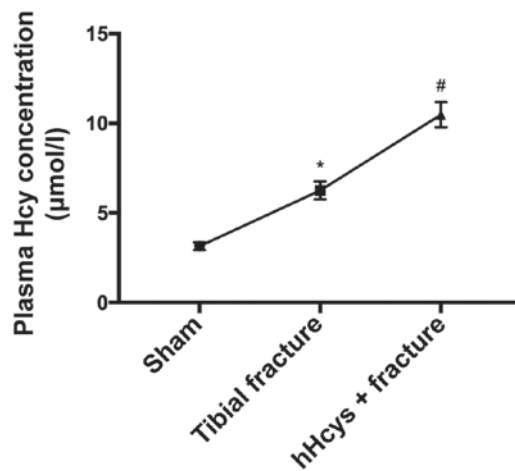


Figure 1. Comparison of plasma Hcy concentration in each group. *P<0.05 vs. sham group; #P<0.05 vs. tibial fracture group. Hcy, homocysteine.

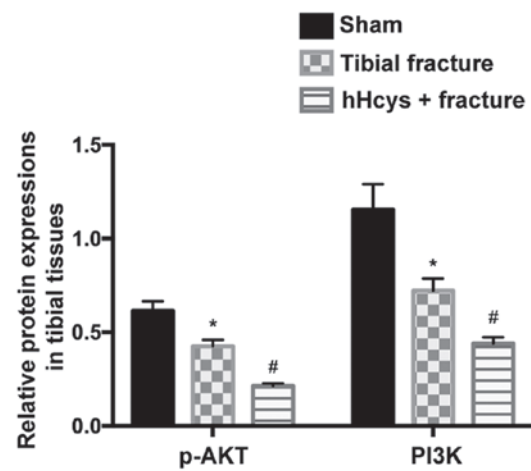


Figure 3. Comparison of relative protein expressions in tibial tissues in each group. *P<0.05 vs. sham group; #P<0.05 vs. tibial fracture group.

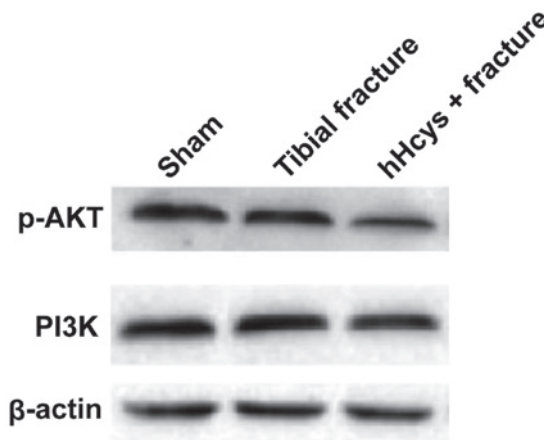


Figure 2. Relative protein expressions in tibial tissues in each group.

processing. The t-test was used for analyzing measurement data. Differences between two groups were analyzed by using the Student's t-test. Comparison between multiple groups was done using One-way ANOVA test followed by Post Hoc Test (Least Significant Difference). P<0.05 suggested that the difference was statistically significant.

Results

Comparison of plasma Hcy concentration in each group. The plasma Hcy concentration was significantly increased in tibial

fracture group and hHcys + fracture group compared with that in sham group (P<0.05), while it was also significantly increased in hHcys + fracture group compared with that in tibial fracture group (P<0.05), showing statistically significant differences (Fig. 1).

Comparison of fracture biomechanics in each group. The ultimate bending strength and torque were obviously decreased in tibial fracture group and hHcys + fracture group compared with those in sham group (P<0.05), while they declined in hHcys + fracture group compared with those in tibial fracture group (P<0.05), showing statistically significant differences (Table I).

Protein expressions of PI3K and p-AKT in tibial tissues detected via western blotting. The relative protein expressions of PI3K and p-AKT evidently declined in tibial fracture group and hHcys + fracture group compared with those in sham group (P<0.05), while they also declined in hHcys + fracture group compared with those in tibial fracture group (P<0.05), displaying statistically significant differences (Figs. 2 and 3).

Comparison of mRNA levels of Bax and caspase-3. Compared with sham group, tibial fracture group and hHcys + fracture group had increased mRNA levels of Bax and caspase-3 (P<0.05). The mRNA levels of Bax and caspase-3 were higher in hHcys + fracture group than those in hHcys + fracture group (P<0.05), and the differences were statistically significant (Fig. 4).

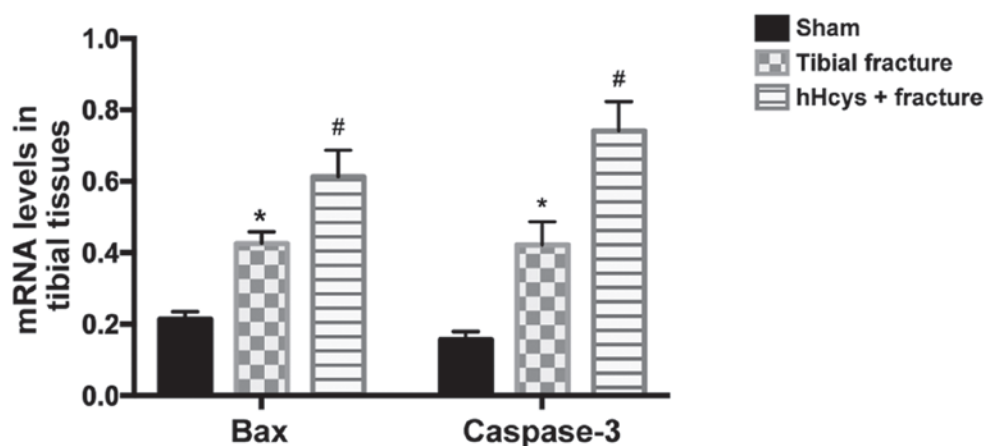


Figure 4. Comparison of mRNA levels of Bax and caspase-3. * $P < 0.05$ vs. sham group; # $P < 0.05$ vs. tibial fracture group.

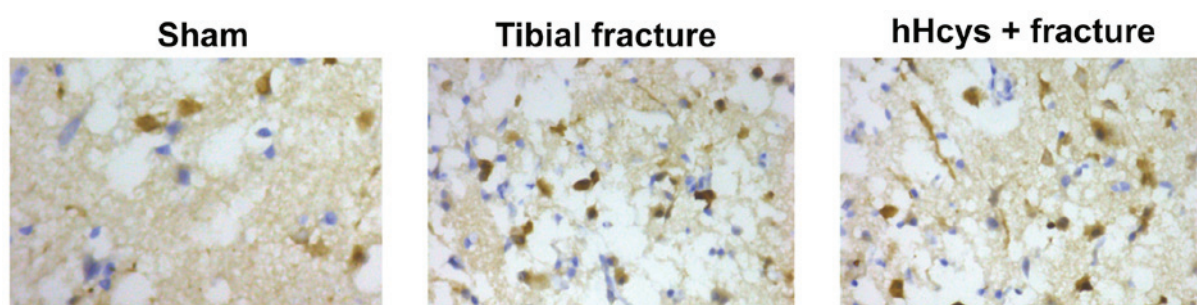


Figure 5. Apoptotic cells in tibial tissues in each group (TUNEL, x400).

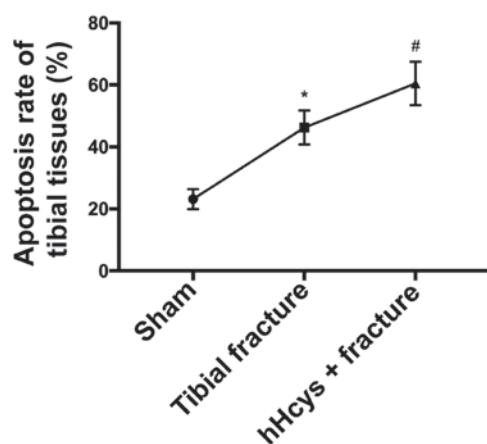


Figure 6. Comparison of apoptosis rate of tibial tissues in each group. * $P < 0.05$ vs. sham group; # $P < 0.05$ vs. tibial fracture group.

Comparison of apoptosis rate. Tibial fracture group and hHcys + fracture group had a higher apoptosis rate than sham group ($P < 0.05$), while hHcys + fracture group also had a higher apoptosis rate than tibial fracture group ($P < 0.05$), and there were statistically significant differences (Figs. 5 and 6).

Comparison of expression of pro-inflammatory factor. The expression of TNF- α was increased in tibial fracture group and hHcys + fracture group compared with that in sham group ($P < 0.05$), while it was also increased in hHcys + fracture group

compared with that in tibial fracture group ($P < 0.05$), and the differences were statistically significant (Figs. 7 and 8).

Discussion

The bone possesses a strong self-repairing ability, but ~5-10% of fracture patients suffer from inadequate union, delayed union or nonunion. In this study, the plasma Hcy level rose after tibial fracture. Hcy is a kind of non-protein amino acid synthesized by methionine and recycled into methionine or converted into cysteine with the help of B vitamins. The increased level of plasma Hcy is associated with the increase in incidence rate of fracture, and the high-level plasma Hcy may affect bone health, which will lead to bone resorption disorders through stimulating osteoclast activity, and interfere with collagen cross-linking. The *in vitro* studies have shown that the elevated concentration of Hcy has an inhibitory effect on bone formation, which induces apoptosis through ROS-mediated mitochondrial pathway and NF- κ B activation in human bone marrow mesenchymal stem cells (hBMSCs), and Hcy can promote the development of osteoporosis by reducing bone formation (13). Moreover, high-concentration Hcy inhibits the activity of lysyl oxidase (an enzyme involved in collagen cross-linking), and the interference with cross-linking will alter the bone matrix and result in fragile bones. Excessive Hcy also leads to loss of bone substance. Type I collagen, the main organic component in bone, is composed of two non-helical telopeptides and a triple helix region at the amino terminal (N) and carboxy terminal (C) of the molecule, which stabilizes

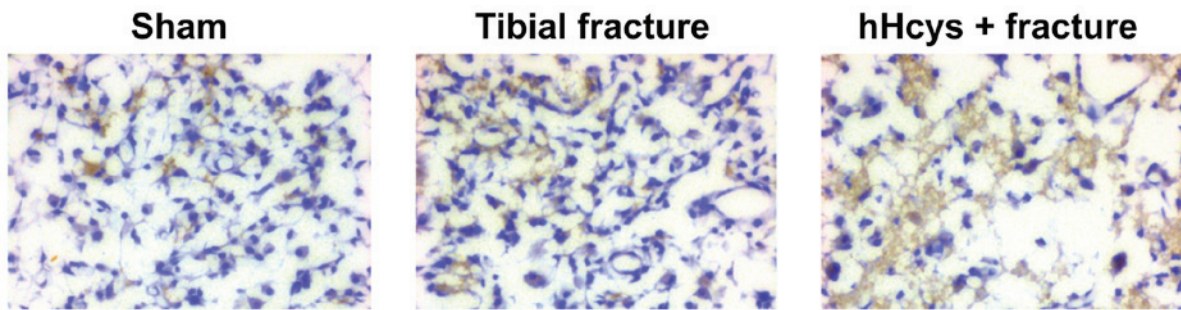


Figure 7. Expression of pro-inflammatory factor in tibial tissues in each group (immunohistochemistry, x400).

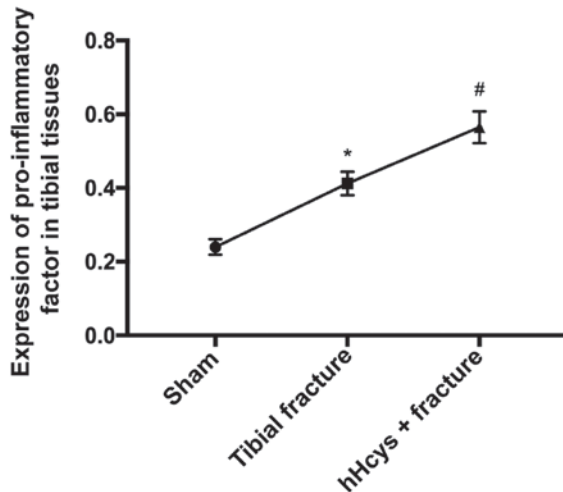


Figure 8. Comparison of expression of pro-inflammatory factor in tibial tissues in each group. *P<0.05 vs. sham group; #P<0.05 vs. tibial fracture group.

the newly secreted collagen molecules through forming cross-linking between adjacent collagen molecules, thus affecting the tensile strength and toughness of bone (14,15). In this study, it was found via biomechanical measurement that both ultimate bending strength and torque declined in hHcys + fracture group compared with those in tibial fracture group, indicating that excessive Hcy affects the bone quality in tibial fracture.

The bone is a kind of dynamic tissue that constantly renews the osteoclast activity through muscles, and absorbs mineralized bones and osteoblasts, forming the new bone matrix. During the whole process, some osteoblasts are embedded in new bones and differentiate into osteoblasts (16). The PI3K/AKT signaling pathway is an important pathway in stress fracture repair, and it is activated in stress fracture callus tissues, which is important for osteoblast differentiation and function (17,18). The AKT signal regulates the osteogenesis after injury through producing osteoprotegerin, and inhibiting the AKT signal transduction in the rat model of fracture nonunion can reduce the transplantation and differentiation of MSCs into fracture callus tissues (19). In addition, studies have demonstrated that PI3K can regulate periosteal thickening in the early stage of fracture repair (20). Therefore, PI3K/AKT is an important signal transduction pathway for bone regeneration. In this study, the PI3K/AKT signaling pathway was damaged in rats after tibial fracture,

and the expressions of PI3K and p-AKT obviously declined in hHcys + fracture group compared with those in tibial fracture group. Besides, the PI3K/AKT signaling pathway mediated apoptosis and inflammatory response, and affected osteocyte differentiation.

In conclusion, in the present study, the role of the PI3K/AKT signaling pathway in fracture repair was explored using the rat model of hHcys and tibial fracture. The results demonstrate that hHcys blocks the downstream apoptotic signal transduction, promotes apoptosis and inflammatory response, and affects fracture healing through affecting the PI3K/AKT signaling pathway.

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Availability of data and materials

All data generated or analyzed during this study are included in this published article.

Authors' contributions

SL and JG designed the study and performed the experiments, SL and YH established the animal models, JG and ST collected the data, WZ and YX analyzed the data, SL and JG prepared the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

This study was approved by the Animal Ethics Committee of Soochow University Animal Center (Suzhou, China).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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