

Research Article

Li-Cheng Xi, Hong-Yu Li*, Dong Yin

Long non-coding RNA-2271 promotes osteogenic differentiation in human bone marrow stem cells

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Abstract: Background: Human bone marrow mesenchymal stem cells (BMSCs) are of great significance for bone regeneration and bone formation. Long non-coding RNAs (lncRNAs) may be involved in modulating cell differentiation. This study aimed to investigate the role of lncR-2271 in promoting osteogenic differentiation in human BMSCs. Methods: Human BMSCs were infected using lncR-2271 overexpression (group A) with lentiviral system or transfected with lncR-2271 siRNA (group B). Cells transfected with scrambled plasmids were used as a negative control (group C). Osteogenesis markers were evaluated using alkaline phosphatase (ALP) activity, RUNX2 and osterix (OSX) at protein levels and calcification by Alizarin Red staining. Results: BMSCs from group A showed significantly higher ALP activity compared to BMSCs in group B and control group (group C) at both days 7 and 14 following osteogenic induction; ALP activity was significantly lower in the group B compared to the group C. RUNX2 and OSX protein expressions were significantly higher in group A and significantly lower in group B, compared to those in group C, respectively. At day 21, calcification in human BMSCs in group A was significantly higher compared to groups B and C as shown by Alizarin Red staining; calcification was significantly lower in group B compared to group C. Conclusion: Our data suggested lncR-2271 played a role in promoting osteogenic differentiation in human BMSCs. This study is the first to illustrate the important role of lncR-2271 in bone formation.

Keywords: long non-coding RNA, osteogenesis, human bone marrow stem cells

1 Introduction

Bone formation by osteoblasts is crucial for bone remodeling hemostasis. A defect in bone formation can greatly hamper bone remodeling, bone regeneration, fracture healing, and overall skeletal health [1, 2]. Bone marrow mesenchymal stem cells (BMSCs) are a group of multipotent stem cells that can differentiate into functional osteoblasts [3]. BMSCs have stable self-renewal ability and they are considered promising candidates for bone tissue regeneration in clinical applications [4-6]. Osteogenesis is a complex cell differentiation process modulated by several genetic and epigenetic factors, among which RUNX2 and osterix (OSX) are key transcription factors that play a pivotal role in osteoblast differentiation and bone formation [7, 8]. However, the epigenetic control of osteogenesis is not clearly described [9].

Long non-coding RNA (lncRNA) is a class of non-coding RNA sized between 200 – 100,000 nucleotides, which is found in eukaryotes and plays an important role in epigenetic regulation in mammals [10]. Current findings suggest that lncRNAs are largely involved in multiple important molecular functions such as regulation of gene expression and post-transcriptional RNA modifications, most likely mediated via interactions with relevant transcription factors [10, 11]. lncRNAs are also closely related to many human diseases including cardiometabolic and immunology disorders [12, 13]. More importantly, recent research indicates that lncRNAs effectively regulate mesenchymal stem cell differentiation [14], indicating that lncRNAs may also affect the osteogenic potential of BMSCs. However, relevant data to support this is scarce.

In our previous study, we screened 87 lncRNAs by microarray analysis and demonstrated that several lncRNAs were differentially expressed during

*Corresponding author: Hong-Yu Li, Department of Orthopedics, The People's Hospital of Guangxi Zhuang Autonomous Region, No 6, Taoyuan Road, Qingxiu District, Nanning 530021, Guangxi Zhuang Autonomous Region, China, E-mail: 469459442@qq.com

Li-Cheng Xi, Dong Yin, Department of Orthopedics, The People's Hospital of Guangxi Zhuang Autonomous Region, No 6, Taoyuan Road, Qingxiu District, Nanning 530021, Guangxi Zhuang Autonomous Region, China

osteogenic differentiation, among which lncRNA ENSG00000232271 (named lncR-2271; chromosome 20; start site: 7,050,261; end site: 7,127,303) was the most significantly differentially expressed lncRNA [15]. This suggested that lncR-2271 is possibly involved in the process of BMSC osteogenesis, however, the specific relationship between lncR-2271 and human BMSC osteogenesis is unclear.

In this study, we intended to investigate the role of lncR-2271 in osteogenic differentiation of BMSCs by overexpressing or inhibiting lncR-2271 expression and evaluate its effect on the osteogenic differentiation of human BMSC. We also attempted to explore the potential mechanisms by probing the expressions of critical regulators of osteogenesis including RUNX2 and OSX.

2 Materials and Methods

2.1 Cells

This study used human BMSCs purchased from the Cobioer Biosciences Co., Ltd (Nanjing, China). Routine cell culture was performed using Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS; Invitrogen, California, USA) in a humidified incubator with 5% CO₂ at 37°C. All culture medium was changed every 2 days and routine cell passage was performed when cells reached confluency. Cells were consistently used at passages 3 to 5. After adjusting the density of BMSCs to 1 × 10⁶ ml by cell counting method, the cell culture medium was inoculated into the pore plate.

2.2 Analysis of BMSCs surface antigen markers

The third passage of BMSCs were digested and suspended by trypsin, then centrifuged twice with a rotational speed of 1200 r/min for 5 mins with phosphate-buffered saline (PBS). Then the BMSCs were counted in L-DMEM (Gibco, USA) complete culture medium, and the density of BMSCs was adjusted to 1 × 10⁶ cells per mL, and divided into 5 tubes. Briefly, the cells were grown to 70% confluence, trypsinized, and washed with PBS. One was used as a blank control, and the remaining four were included to each of them by adding mouse anti-human CD34, CD45, CD44 and CD29 / FITC monoclonal antibodies (Cell Signaling Technology, Inc., Santa Cruz, CA, USA) 5 μl respectively, and then placed at room temperature for 20 mins, and then centrifuged each sample tube (rate

1200 rpm/min) 5 mins and discard the supernatant. After washing with 0.5 ml washing solution for two times, the cells were resuspended in 1 × PBS equilibrium salt solution of each tube. Finally, the cell surface antigen was detected and analyzed on a FACS Canto II (Becton Dickinson) using Diva Software (Beckton Dickinson).

2.3 Knockdown or overexpression of lncR-2271

The design and synthesis of lncR-2271, plasmids and lentiviruses were accomplished by GeneChem (Shanghai GeneChem, Shanghai, China). The lncR-2271 sequence was amplified using the indicated primers: lncR-2271 Forward: 5'-AAC CTT TGC TCC AAC TTT CTC C-3' and Reverse: 5'-ACA CCC TGG CAG TGC CTA TT-3'. Full-length lncR-2271 and pGreenFire control plasmids (Cat no. TR000PA-1, System Biosciences, LLC, CA, USA) were cloned into pcDNA3.1 vector (Invitrogen, Carlsbad, CA, USA). And the blank plasmid vector is made as an NC. For lentivirus packaging, 293T cells were transferred with an lncRNA-2271 plasmid vector. Preparation of lentivirus mixture before transfection: adding 15 μL EndoFection transfection reagent (Invitrogen, Carlsbad, CA, USA) into the mixture containing 5 μL Lenti-Pac HIV reagents (Cat no. HPK-LvTR-40, GeneCopoeia, USA) and lentivirus expression plasmids, then resting at room temperature, then adding them to the medium for overnight culture. The harvested lentiviruses were concentrated, purified and preserved at -80°C. BMSCs were cultured at a concentration of 1 × 10⁶ cells per well in 24-well plates. After the cells had grown to 50-60% confluence, they were transfected with lentiviruses in the presence of polybrene at a multiplicity of transfection of 50. Plasmid transfection was conducted using Lipofectamine 2000 (Cat no. 11668-019, Invitrogen) transfection reagent following the manufacturer's protocol. After 48 h of transfection, the cells were collected and used for further investigations. The supernatants were placed on primary human BMSCs and incubated in the presence of polybrene (Nanjing SenBeiJia Biological Technology Co., Ltd., Nanjing, China) for 72 hours.

Transfection was performed by treating BMSCs with lncR-2271 siRNA (Cat no. L1025, Genepharma, Shanghai, China) (group B) using lipofectamine 2000 (Cat no. 11668-019, Invitrogen). Cells transfected with control siRNA (cat no. CW0871, System Biosciences, LLC, CA, USA) were categorized as control (group C). Sequences of siRNAs used in these studies: F: 5'-GGACGAGGAGTTATTCTATTG dTdT-3' and reverse: R: 5'-dTdTCTGCTCGTCAATAAGATAAC-3'.

2.4 Analysis of cell survival and transfection efficiency

After 48 hours of transfection, 10 μ L Cell Counting Kit-8 solution (Cell Proliferation Kit, Dojindo Laboratories, Japan) were added into each 24-well and incubated for 2 hours at 37. Absorbance of each well was measured using a microplate reader (μ Quant, Bio-Tek, USA) at 450 nm.

After 48 hours of transfection, media was removed from cells and rinsed with PBS. GFP intensity was measured using versatile fluorescence micro plate reader at 509 nm (μ Quant, Bio-Tek, USA).

2.5 Osteoblastic differentiation induction

Following transfection, BMSC osteoblastic differentiation was induced using osteoblastic culture media containing DMEM with 5% FBS and 200 ng/ml recombinant human bone morphogenic protein 2 (BMP2; PeproTech, Rocky Hill, USA) for 21 days.

2.6 Alkaline phosphatase activity assay

Alkaline phosphatase (ALP) activity was evaluated at days 7 and 14 after differentiation induction using a commercial kit according to the manufacturer's protocol (Labtest Diagnostica SA, Lagoa Santa, MG, Brazil). Briefly, cells were lysed by five cycles of thermal shock (-20°C for 20 min and 37°C for 15 min). Fifty μ L of preheated (37°C for 2 min) thymolphthalein monophosphate containing 0.3 M diethanolamine buffer (pH 10.1) was then added to the cell lysate and the mixture was kept at 37°C for 10 min. The reaction was stopped by adding 2 mL of a sodium carbonate (0.09 mmol/mL) and sodium hydroxide (0.25 mmol/mL) solution. Optical density was measured at 590 nm (μ Quant, Bio-Tek, USA) using distilled water as a blank control. ALP activity was averaged from quintuplicate ($n = 5$) samples and normalized to total protein content [16].

2.7 Alizarin red staining

Calcium deposits in differentiated BMSCs at day 21 were detected by alizarin red staining. Cells were fixed with 4% paraformaldehyde after washing with phosphate buffered saline (PBS) for 30 min. Alizarin red solution was then added to the cell culture for 3 to 5 min. Cells were then observed under a microscope (SZL6745 Zoom

Stereo Microscope, Yuyao Dagong Instrument Co., Ltd., Hangzhou, China) after washing with PBS again, and images were taken. Image-Pro Plus 6.0 analysis software (Media Cybernetics, Bethesda, MD, USA) was used to evaluate the staining density of the images.

2.8 Real-time quantitative PCR

Total RNA was extracted from BMSCs at days 7 and 14 after differentiation using the TRIzol reagent (Life Technologies, USA), and RNA concentrations were evaluated using NanoDrop 1000 (Thermo Fisher Scientific, MA, USA). cDNA was first synthesized using the Moloney Murine Leukemia Virus Reverse Transcriptase kit (Life Technologies, USA) using random primers (Takara Bio, Otsu, Japan) at 55°C for 30 min. Real-time PCR was then performed by combining the cDNA product with 2 \times Taq PCR Master mix (Tiangen Biotech Co., Beijing, China) containing 10 pmol/l of primer pairs (Takara Bio, Japan). The qPCR procedure was performed on a LightCycler 480 instrument (Roche Diagnostics, Mannheim, Germany) and the cycle was as follows: 95°C for 15 min, 40 cycles of 94°C for 15 sec, 55°C for 30 sec, and 72°C for 30 sec. The fluorescence was measured and the relative expression was calculated by the $2^{-\Delta\Delta\text{Ct}}$ method using glyceraldehyde phosphate dehydrogenase (GAPDH) for normalization [17]. The primer sequences were as follows: IncR-2271 Forward: 5'-AAC CTT TGC TCC AAC TTT CTC C-3' and Reverse: 5'-ACA CCC TGG CAG TGC CTA TT-3'; GAPDH forward: 5'-ACA GCC TCA AGA TCA TCA GCA-3' and reverse: 5'-CAT CAC GCC ACA GTT TCC C-3'.

2.9 Western blot

Total protein was extracted from BMSCs using a lysis buffer (Beyotime, China) and further separated by 10% SDS-PAGE. Separated proteins were then transferred to a nitrocellulose filter membrane. The membrane was incubated with primary antibodies for either RUNX2 (1:500, ab76956, Abcam) or OSX (1:1,000, ab187158, Abcam), and tubulin (1:1,000, ab59680, Abcam) as a loading control. The membranes were then incubated with a corresponding horseradish peroxidase (HRP)-conjugated secondary antibody (anti-mouse IgG, HRP, 1:5,000, ab6728, or anti-rabbit IgG, HRP, 1:5,000, ab6721, Abcam). Protein band signals were visualized using an enhanced chemiluminescence (ECL) kit (Pierce, IL, USA), and band intensity was further analyzed by Image-J software (version 1.42, NIH, USA).

2.10 Statistical analysis

All statistical analyses were performed with SPSS18.0 statistical software (SPSS Inc., USA). Comparisons of lncR-2271 transfection efficacy, ALP activity, Osteoblastic marker expression and Calcification expression between two groups were performed by independent t-test. Data are presented as mean \pm standard deviation (SD). A P value less than 0.05 was considered statistically significant. All experiments were replicated at least three times.

Informed consent: Informed consent has been obtained from all individuals included in this study.

Ethical approval: The research related to human use has been complied with all the relevant national regulations, institutional policies and in accordance the tenets of the Helsinki Declaration, and has been approved by the authors' institutional review board or equivalent committee.

3 Results

3.1 Characterization of human BMSCs surface marker antigen (CD molecule)

The human BMSCs cultured in this experiment was characterized by surface antigen relative specificity molecular identification. The results showed that 97% of

the cells were CD44+ and 89% of the cells were CD29+, 6% of the cells were CD45- and 8% of the cells were CD34-. The results of CD antigen molecules on the surface of the above cells were consistent with the identification characteristics of hBMSCs cell surface markers and purity was acceptable.

3.2 lncR-2271 transfection efficacy

Cell survival rate after transfection was higher than 90% and transfection efficiency was more than 50%. Successful manipulations of lncR-2271 expression were demonstrated by RT-PCR in BMSCs 48 hours after transfection (Figure 1). Overexpression of lncR-2271 in group A was about 7-fold greater than the control group C ($P < 0.01$), the relative expression of lncR-2271 in group B was decreased to 30% compared to group C ($P < 0.01$). The endogenous level of lncR-2271 was comparable to the control (group C). Our data suggested that lncR-2271 was successfully overexpressed in group A, and knocked down in the group B.

3.3 Alkaline phosphatase activity

The ALP activity in human BMSCs were not significantly different among group A, group B and group C before transfection. The activity of ALP showed the most obvious difference after 7 days of BMSCs transfection, so we chose to assess it at 7 and 14 days after differentiation. ALP

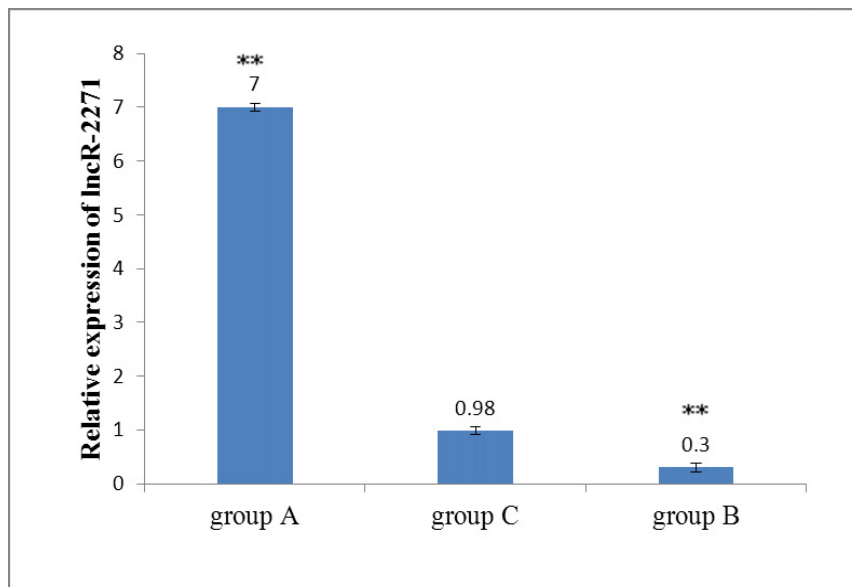


Figure 1. lncR-2271 was overexpressed and knocked down in human BMSCs. Group A: lncR-2271 was overexpressed human BMSCs; Group B: lncR-2271 was knocked down in human BMSCs by transfection with siRNA-lncRNA-2271; Group C: human BMSCs were transfected with scramble plasmids - control; Data are presented as mean \pm SD; ** $P < 0.01$ compared to group C

activity was assessed at days 7 and 14 after differentiation (Figure 2). Our data demonstrate that ALP activity in group A was significantly higher compared to groups B and C at both days (day 7, A: 119.6 ± 3.2 vs. B: 39.0 ± 1.0 ; A vs. C: 69.4 ± 1.9 ; day 14, A: 340.4 ± 5.9 vs. B: 97.8 ± 2.7 ; A vs. C: 159.0 ± 3.8 ; all $P < 0.01$). ALP activity in group B was also significantly lower compared to group C (day 7, B vs. C, $P < 0.01$; day 14, B vs. C, $P < 0.01$). This part of the experiment shows that lncR-2271 can enhance the ALP activity in bone differentiation of BMSCs, while ALP activity is a sign of early osteogenic differentiation, so the experiment further shows that lncR-2271 can promote the osteogenic differentiation of BMSCs.

3.4 Osteogenic marker expression

RUNX2 and OSX are assessed to test effect of lncR-2271 on osteogenic differentiation of BMSC. The expressions of RUNX2 and OSX in human BMSCs were no differences among group A, group B and group C before transfection. The expression of RUNX2 and OSX showed the most obvious difference after 7 days of BMSCs transfection; thus, they were assessed at 7 and 14 days after differentiation. Protein expression of osteoblastic markers RUNX2 and OSX were analyzed at days 7 and 14 after BMSC differentiation (Figure 3). Our data demonstrate that group A had significantly higher RUNX2 and OSX protein expression compared to groups B and group C after transfection at day 7 (RUNX2, A: 0.83 ± 0.05 vs. B: 0.23 ± 0.02 , $P < 0.05$; A vs. C: 0.56 ± 0.04 ,

$P < 0.01$; OSX, A: 0.83 ± 0.05 vs. B: 0.27 ± 0.02 ; $P < 0.05$; A vs. C: 0.53 ± 0.04 , $P < 0.01$) (Figure 3A). A similar trend was also demonstrated at day 14 (RUNX2, A: 1.47 ± 0.12 vs. B: 0.48 ± 0.03 , $P < 0.01$; A vs. C: 0.87 ± 0.05 , $P < 0.01$; OSX: A: 1.47 ± 0.11 vs. B: 0.57 ± 0.03 , $P < 0.01$; A vs. C: 1.00 ± 0.06 , $P < 0.01$) (Figure 3B). RUNX2 and OSX expression in group B was also significantly lower compared to group C (day 7, RUNX2: B: 0.23 ± 0.02 vs. C: 0.56 ± 0.04 , $P < 0.01$; OSX: B: 0.27 ± 0.02 vs. C: 0.53 ± 0.04 , $P < 0.01$; day 14, RUNX2: B: 0.48 ± 0.03 vs. C: 0.87 ± 0.05 , $P < 0.01$; OSX: B: 0.57 ± 0.03 vs. C: 1.00 ± 0.06 , $P < 0.01$) (Figure 3C). Our results indicated that overexpression of lncR-2271 increases the expressions of RUNX2 and OSX in human BMSCs, while the inhibition of lncR-2271 expression reduces the expressions of RUNX2 and OSX. RUNX2 is significant transcription factor in osteogenic differentiation, and OSX is important for bone formation. Therefore, our data suggested that lncR-2271 can promote osteogenic differentiation of human BMSCs.

3.5 Calcification

The calcification in human BMSCs were no differences among group A, group B and group C before transfection. Calcification level are assessed to test effect of lncR-2271 on osteogenic differentiation of BMSC. Calcium deposition is later several days than RUNX2 and OSX. Calcification was evaluated at day 21 after differentiation using Alizarin red staining. BMSCs in group A exhibited the most calcium nodules; whereas BMSC in group B exhibited the least

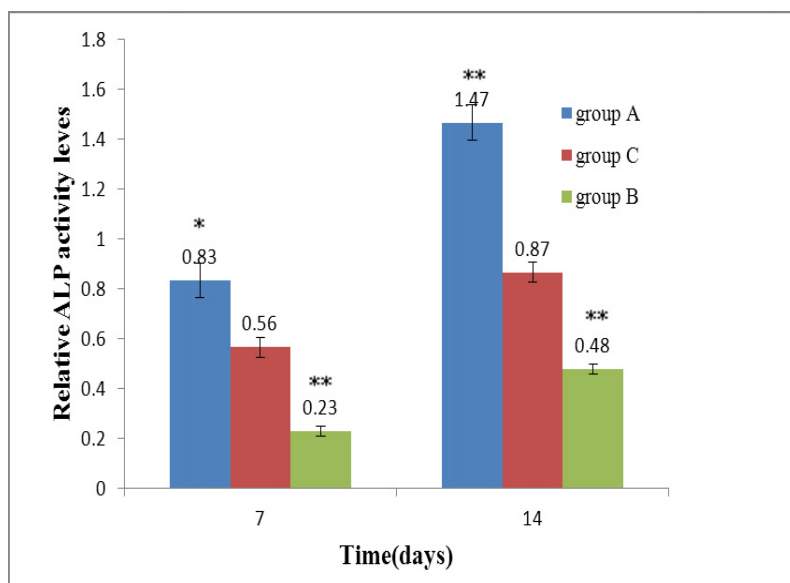


Figure 2. ALP activity was analyzed in human BMSCs at days 7 and 14 after osteoblastic differentiation. Group A: lncR-2271 was overexpressed human BMSCs; Group B: lncR-2271 was knocked down in human BMSCs by transfection with lncRNA-2271 siRNA; Group C: human BMSCs were transfected with scrambled plasmids - control; Data are presented as mean \pm SD; ** $P < 0.01$ compared to group C.

calcification by day 21 (Figure 4A). The optical density of alizarin red staining also demonstrated that calcification in group A was significantly higher compared to groups B and C (A: $215 \pm 0.05\%$ vs. B: $44 \pm 0.02\%$, $P < 0.01$; A vs. C: $100 \pm 0.04\%$, $P < 0.01$). The calcification level in BMSCs in group B was also significantly lower compared to group C ($P < 0.01$) (Figure 4B). The results show that overexpression of lncR-2271 promotes calcification, while decrease of lncR-2271 expression inhibits calcification. Calcium deposition is a late indicator of osteogenic differentiation in human BMSCs and a marker of late osteogenic differentiation. Therefore, our data suggested that lncR-2271 regulates the osteogenic differentiation of human BMSCs.

4 Discussion

At present, bone defects, cartilage defects and degenerative diseases caused by trauma and tumor, osteonecrosis and the collapse of bone remain challenging

in clinical treatment. In this study, we found that lncR-2271 overexpression promoted ALP activity, calcification, and expressions of the osteogenic transcription factors RUNX2 and OSX; whereas lncR-2271 down-regulation hampered osteogenesis. Our data demonstrated that lncR-2271 plays an important role in osteogenic development in human BMSCs and is of great importance to bone regeneration and bone formation. This study provides new potential therapeutic target for the treatment of bone defects, cartilage defects and osteonecrosis diseases using human BMSCs.

Despite the current understanding of the role of lncRNAs in a number of physiologic and pathologic conditions, little is known about lncRNAs in bone health. However, with the basic completion of human genome sequencing, more and more studies have confirmed the important role of lncRNAs in promoting osteogenic differentiation of BMSCs [18]. The process of osteogenic differentiation of BMSCs is regulated by

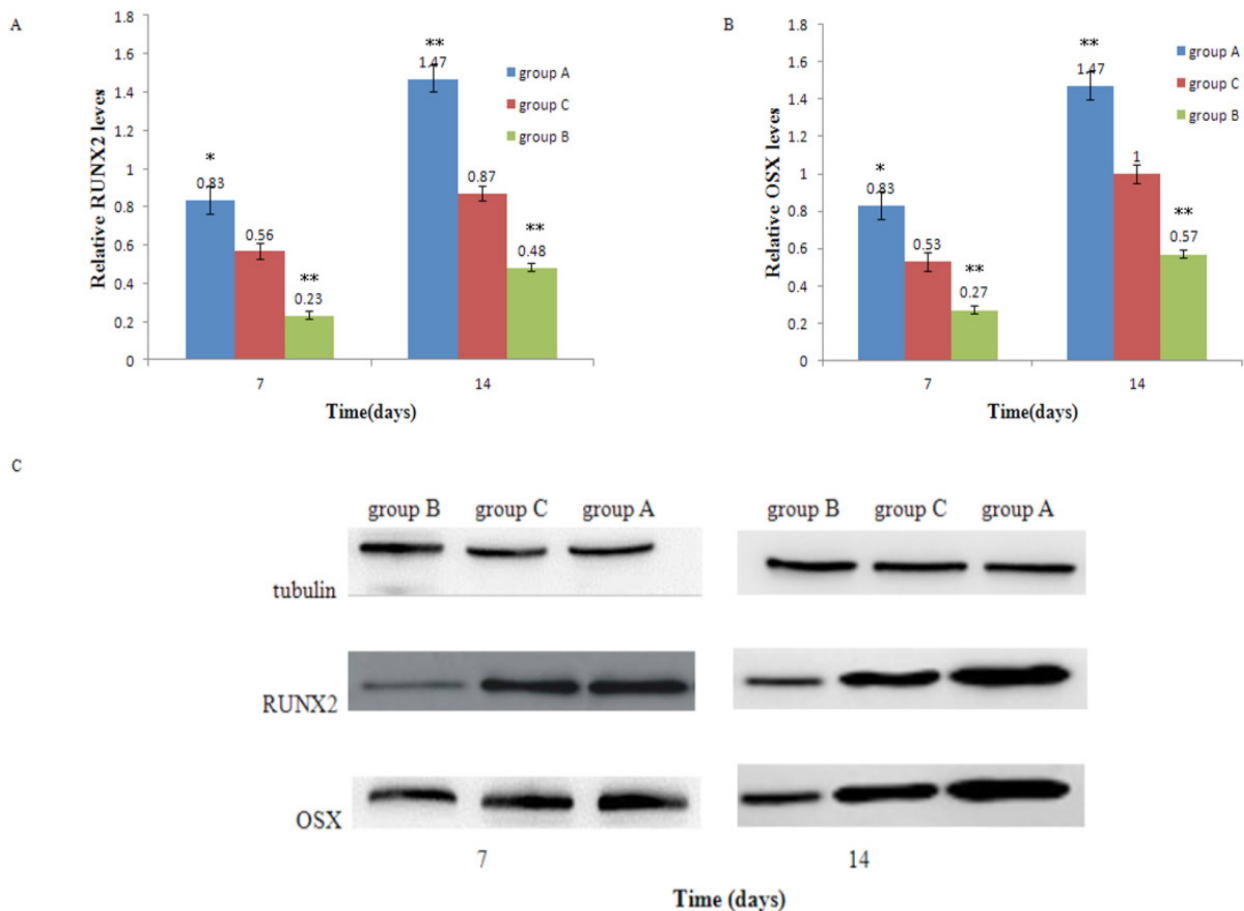


Figure 3. RUNX2 and OSX protein expressions were analyzed in human BMSCs at days 7 and 14 after osteoblastic differentiation. **A)** RUNX2 expression; **B)** OSX expression; **C)** Western blot results for RUNX2 and OSX; group A: lncR-2271 was overexpressed human BMSCs; group B: lncR-2271 was knocked down in human BMSCs by transfection with lncRNA-2271 siRNA; group C: human BMSCs were transfected with scrambled plasmids - control; Data are presented as mean \pm SD; * $P < 0.05$, ** $P < 0.01$ compared to group C.

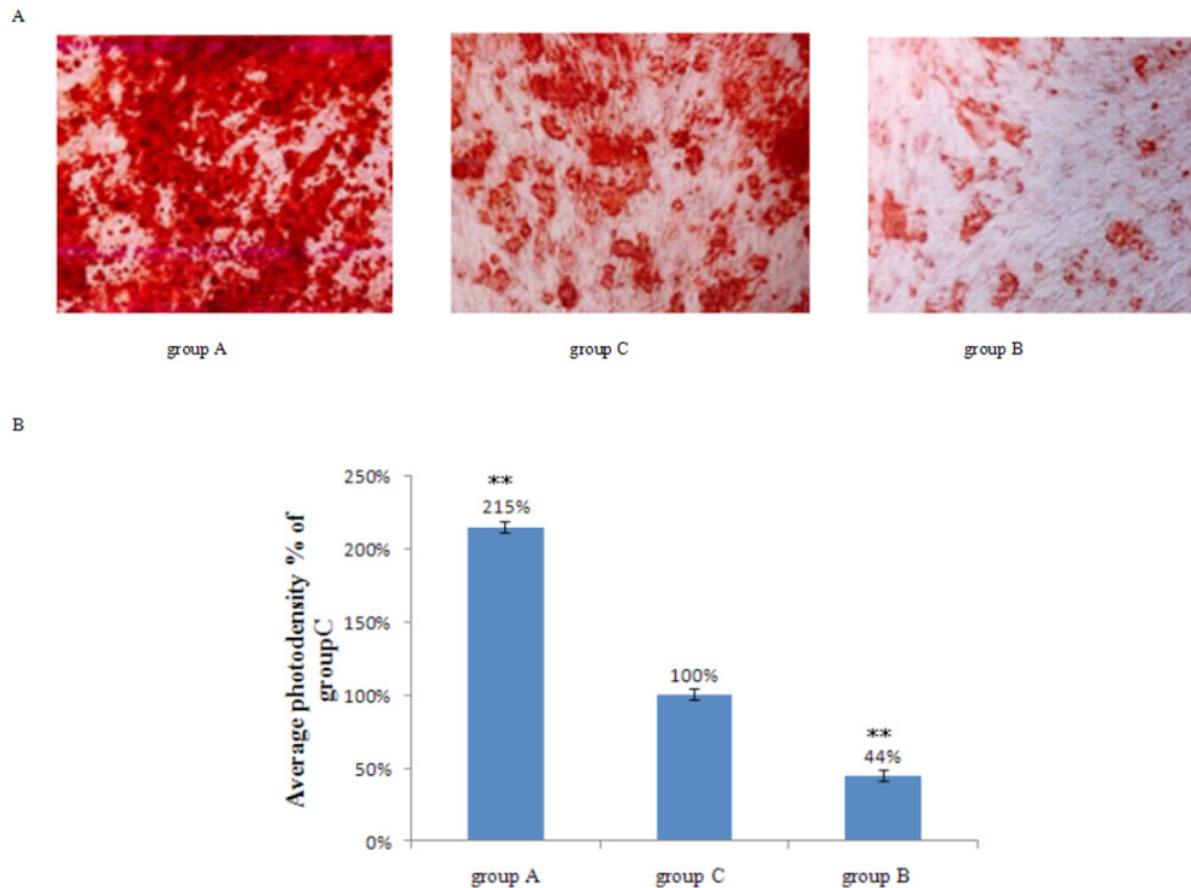


Figure 4. Calcification indicated by Alizarin red staining in human BMSCs after osteogenic differentiation induction for 21 days. A) microscopic images of Alizarin red staining results; **B)** quantification of average staining density of cells. Group A: lncR-2271 was overexpressed human BMSCs; Group B: lncR-2271 was knocked down in human BMSCs by transfection with lncRNA-2271 siRNA; Group C: human BMSCs transfected with scrambled plasmids - control; Data are presented as mean \pm SD; ** $P < 0.01$ compare

up- or down-regulation of specific genes expression in a specific sequence [19]. But the early stage of osteogenic differentiation determines the direction of BMSCs formation. Therefore, the study of lncRNA on the regulation of early stage of osteogenic differentiation in BMSCs will provide more insights for the research and development of bone tissue engineering and will also provide a new target for the clinical treatment, prognosis and screening of drugs for bone tissue repair. This study first provided information about the important role of lncR-2271 in osteogenesis of human BMSCs. Our findings regarding lncR-2271 are in accordance with previous studies showing the role of lncRNAs in osteogenesis. Xu *et al* showed that lncR-HIF1 α -AS1 promoted osteoblast differentiation via increasing expression of Homeobox D10 [19]. Zhuang *et al.* also demonstrated that lncR-MEG3 enhanced osteogenesis in BMSCs from

multiple myeloma patients through promoter-specific transcriptional activation of bone morphogenic protein 4 (BMP4) [20]. In contrast, a previous study suggested that lncR-NONHSAT009968 was involved in inflammation, which potentially inhibited osteoblast development [21]. Our study is the first to characterize the role of lncR-2271 in promoting BMSC osteogenic differentiation and demonstrated that lncR-2271 controls the osteogenic differentiation of BMSC by regulating the expressions of key regulators RUNX2 and OSX during osteogenic differentiation.

ALP activity, calcification and collagen type I expressions are generally used as indicators of osteogenic differentiation. ALP, as one of the important characteristics of osteoblast phenotype, plays a very important role in calcification *in vitro*. A large amount of calcium deposits occur after ALP is increased. The higher

the ALP activity is, the stronger the osteogenic activity is. Osteocalcin has a role in promoting cell mineralization, and osteocalcin plays an important role in maintaining a certain bone calcification, which is recognized as the most specific marker of osteogenic differentiation. Our study demonstrates that lncR-2271 overexpression increases osteogenic phenotypes, including increased ALP activity and the number, size and color depth of bone calcium nodules, indicating that lncR-2271 can potentially affect multipotent human BMSCs down the osteogenic pathway. However, whether lncR-2271 also has a crucial role in bone formation and repair in physiological homeostasis remains uncertain. The mechanism by which lncRNA modulates osteogenesis is most likely through regulating expression of key differentiation genes. Many molecules play an important role in the process of osteogenic differentiation of BMSCs. In this study, we evaluated two osteogenesis-related transcription factors, RUNX2 and OSX. RUNX2 is known as a critical regulator of osteoblast differentiation, which functions as a transcription factor and is required for osteogenic differentiation. The structure of RUNX2 contains 3 domains, which are the N terminal glutamine / alanine repeat sequence domain (Q / A), the C end consists of proline, serine, threonine, and the PST domain in the middle DNA binding domain [22]. It regulates expression of ossification related genes and therefore plays a pivotal role in the formation, differentiation and maturation of osteoblasts, the formation and absorption of osteoclasts, the differentiation and maturation of chondrocytes, and the production of bone matrix proteins [7, 22]. OSX acts downstream of RUNX2 and is important for bone formation. The final differentiation and maturation of osteoblasts requires the expression of Osterix. When Osterix is downregulated, BMSCs was not able to differentiate into osteoblasts, which is confirmed by no markers expression from endochondral or endochondral osteoblasts [8]. In this study, lncR-2271 overexpression significantly increased protein expression of both RUNX2 and OSX. More importantly, lncR-2271 down-regulation significantly reduced the osteogenic potential of human BMSCs and expression of both RUNX2 and OSX, indicating the indispensable role of lncR-2271 in osteogenesis and bone formation. However, whether lncR-2271 effects are through a direct interaction with RUNX2 and OSX or through other regulating elements requires further study.

The current study has demonstrated that lncR-2271 regulates osteogenesis of BMSC through modulating the expressions of RUNX2 and OSX. This could be possibly mediated through the TGF- β Smad pathway, MAPK/ERK pathway or β -catenin pathway [23] or microRNAs [24, 25]. However, the specific pathway for lncR-2271 to regulate the

expression of RUNX2, the mode and the site of action need to be further verified in our future studies.

5 Conclusions

lncR-2271 promoted osteogenesis of human BMSCs via increasing ALP activity, calcification, and expressions of osteogenic transcription factors RUNX2 and OSX. This study is the first to illustrate the important role of lncR-2271 in bone formation; however, future studies are warranted to investigate the interaction between lncR-2271 and epigenetic regulation of bone-related genes. Further understanding of the epigenetic regulation of osteogenesis will be potentially beneficial for clinically promoting bone repair.

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Conflict of interest: Authors state no conflict of interest.

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