Original Research Article



Endogenous bone morphogenetic protein 2 plays a role in vascular smooth muscle cell calcification induced by interleukin 6 in vitro

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Abstract

Systemic inflammation is involved in vascular calcification and cardiovascular disease which is the leading cause of mortality in rheumatoid arthritis (RA). A high level of serum interleukin (IL)-6 plays a key role in local and systemic inflammation in RA. However, the underlying mechanisms remain unclear. We established a human umbilical artery smooth muscle cell (HUASMC) culturing method to investigate the possible role of IL-6 on vascular calcification. HUASMCs were obtained from umbilical arteries of healthy neonates. To detect calcification effects, HUASMCs were treated with (experimental group) or without (control group) recombinant human (rh) IL-6. The calcium deposition stain and calcium concentrations were measured, as well as the mRNA and protein levels of the regulating factor of osteogenic differentiation-bone morphogenetic protein (BMP) 2 and those calcifying related molecules including bonespecific alkaline phosphatase (BAP), osteoprotegerin (OPG), and osteopontin (OPN). Our study showed that rhlL-6 induced calcification of HUASMCs in a time- and dose-dependent manner, and upregulated expressions of BMP2, BAP, OPG, and OPN of HUASMCs. We then used the anti-BMP2 siRNA to knockdown the expression of endogenous BMP2 to confirm its role. HUASMCs were transfected with negative siRNA (control group) or the valid anti-BMP2 siRNA (experimental group) before they were treated with rhIL-6. Cells transfected with negative siRNA without IL-6 stimulating served as the blank group. The results showed that anti-BMP2 siRNA markedly decreased expressions of BMP2, BAP, OPG, and OPN, and also partly reduced the calcification of HUASMCs induced by rhIL-6. Collectively, according to our study, rhlL-6 could induce the extracellular calcification and osteogenic differentiation of human artery smooth muscle cells through upregulating endogenous BMP2 in vitro. This may be one of the underlying mechanisms of the overwhelming vascular calcification in RA.

Keywords

bone morphogenetic protein 2, interleukin 6, rheumatoid arthritis, RNA interfering, vascular calcification

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Rheumatoid arthritis (RA) is a disabling autoimmune disease characterized by chronic erosive arthritis and extra-arthritic manifestations. Recent studies revealed that the mean lifespan of RA patients was a few years less than that of the coetaneous group from the general population despite the improved outcome of arthritis and overall life quality. The leading cause of mortality in RA patients is cardiovascular disease, especially in ¹Department of Rheumatology and Clinical Immunology, the Affiliated Hospital of Qingdao University, Qingdao, China ²Department of Cardiac surgery, the Affiliated Hospital of Qingdao University, Qingdao, China ³Department of Nephrology, Renji Hospital, Shanghai Jiaotong University Medical College, Shanghai, China

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Creative Commons Non Commercial CC-BY-NC: This article is distributed under the terms of the Creative Commons Attribution-NonCommercial 3.0 License (http://www.creativecommons.org/licenses/by-nc/3.0/) which permits non-commercial use, reproduction and distribution of the work without further permission provided the original work is attributed as specified on the SAGE and Open Access pages (https://us.sagepub.com/en-us/nam/open-access-at-sage). voung adults.^{1–3} Studies in diabetes mellitus, chronic kidney disease (CKD), other inflammatory arthritis like ankylosing spondylitis, and aging have suggested that the systemic inflammation is associated with vascular calcification and congenital heart failure.^{4–7} RA is the prototypic inflammatory arthritis with outstanding systemic inflammation. Much research indicated that RA patients are more likely to suffer widely and severe medial artery calcification (MAC). Chronic inflammatory markers are associated with high cardiovascular mortality, high prevalence, and greater extent of peripheral and coronary arterial calcification.7-10 Among these markers, interleukin (IL)-6 is the most prominent one. Tumor necrosis factor (TNF)- α could induce cardiac myocyte apoptosis and remodeling, and was related to congestive heart failure,¹¹ while results of clinical studies in RA patients remained controversial in terms of vascular calcification.¹²⁻¹⁴

MAC is mainly occurred in vascular smooth muscle cells (VSMCs). Previous studies, including ours, have shown that the concentration of serum IL-6 was positively related to some calcification indicators like bone alkaline phosphatase (BAP), osteopontin (OPN), osteoprotegerin (OPG), osteocalcin (OC), and the ultimate aortic calcification score.^{6–9,15} Since BAP, OPN, OPG, and OC are all proteins secreted by osteoblasts, it is reasonable to hypothesize that IL-6 is involved in an osteogenic differentiation and calcification of the VSMCs. Pathological studies found that VSMCs in calcified vessels expressed the core regulating factor of osteogenic differentiation-bone morphogenetic protein (BMP) 2, the key transcription factor of osteogenesis-core binding factor α -1 (Cbfa1), and downstream bone-related proteins.^{16–19} those Therefore, we established a human umbilical artery smooth muscle cell (HUASMC) culturing method to observe the role of IL-6 on vascular calcification in vitro. To confirm the role of endogenous BMP2, small RNA interfering method was used to silence or knockdown the expression of BMP2 of the HUASMCs. Those results may provide a better understanding of the mechanism underlying the overwhelming vessel calcification in RA patients.

Materials and methods

Cell culture and identification

HUASMCs were obtained from artery explants of the umbilical cords of healthy donors. Briefly, arteries of umbilical cords from healthy neonates were separated within 4 h after delivery and thoroughly washed by saline. The sleeve-like adventitia was removed and the remaining tissues were chopped into pieces of 1-2 mm³. The explants were planted on the bottom of T-25 flask in 3-5 pieces/cm² and merged into the Dulbecco's modified Eagle's medium (DMEM) (31600-034, Invitrogen) which was supplemented with 20% fetal bovine serum (Gibco Laboratories), 100 unit/ mL penicillin, and streptomycin. The elongate spindle-shaped cells would migrate out from the edge of explants at approximately one week. After 2-3 weeks, cells migrating from the explants would overgrow the plate (Figure 1a). The confluent cells would appear the classic peak-and-valley pattern at prolonged culture (Figure 1b). All cells were synchronically cultured by serum-free DMEM for 24 h before they were used for any experiment (Figure 1c).

Cells identified by morphological characters were ultimately confirmed to be VSMCs by indirect immunofluorescence (IIF) test of α -SM actin, the marker of SMCs, using monoclonal antibodies against α -SM actin (A2547, Sigma) (Figure 1d). IIF tests of factor VIII and vetamin with monoclonal antibodies against factor VIII (ab41187, abcam) and vimentin (ab8069, abcam) were also applied to expel contaminations from endothelial cells and fibroblasts.

The protocols were approved by the Ethics Board of the Affiliated Hospital of Qingdao University. All participants provided written informed consent.

Calcification staining and measuring

For calcification staining, Alizarin red S method was used. Cells cultured with rhIL-6 50 ug/L (R&D Systems) for six days served as the experimental group and the routinely cultured cells as the control group. Cells were washed with D-Hank's solution, fixed with pure iced ethanol, and were then exposed to 0.5% Alizarin red S (PH 7.0) for 15 min at room temperature (red/orange means a positive staining of extracellular matrix). For calcium concentration measuring, O-cresolphthalein complexone method was used. Cells cultured with rhIL-6 10 ug/L or 50 ug/L served as experimental groups and routinely cultured cells served as controls. The calcium measuring was applied at baseline and after being



Figure 1. Culture and identification of the human umbilical artery smooth muscle cells. (a) The elongated spindle-shaped cells grew out from the edge of the explants after 2–3 weeks' culture ($100\times$). (b) Cells appeared the classic peak-and-valley pattern of morphology in prolonged culture ($100\times$). (c) Cells were synchronically cultured by serum-free DMEM for 24 h ($100\times$). (d) The fiber filaments in the cytoplasm were stained with green fluorescence in the indirect immunofluorescence test of α -SM actin ($400\times$).

treated for 3, 6, 9, and 12 days. Total calcium in the cell layer was extracted by 0.6 mol/L HCl solution for 24 h at 37°C. The remaining components of cells were dissociated by 1 mL of 0.1 N NaOH/0.1% SDS to obtain the total protein. Protein concentrations were determined using a BCA protein assay kit (Pierce, Cat. no.23227) and final concentrations of calcium were adjusted by the protein content of the cell layer.

Quantitative real-time RT-PCR

The control cells were cultured with regular DMEM and the experimental cells were cultured with extra 10 ug/L rhIL-6. Cells were harvested at 12, 24, and 72 h. Total RNA extraction, reverse-transcription, and real-time RT-PCR reactions were carried out according to the manufacturer's instructions (TaKaRa, Japan; ABI PRISM 7900, America). All data were analyzed using ABI PRISM SDS 2.0 software (Applied Biosystems). The mRNA expressions of the target genes were represented using the Δ Ct method; B-actin was co-amplified as the housekeeping gene.¹⁵ Every gene level was transformed into the ratio of experimental group to control group at the same time point during statistical analyzing. The oligonucleotide primers for BMP2, BAP, OPN, and OPG were listed in Table 1.

BAP, bone specific alkaline protein; BMP2, bone morphogenetic protein 2; OPG, osteoprote-gerin; OPN, osteopontin.

Fluorescent quantitative method

This method was used to detect the BAP protein level. The cell treating process was the same as

Genes	Upstream primers	Downstream primers
BMP2	5'-AACACTGTGCGCAGCTTCC-3'	5'-CCTAAAGCATCTTGCATCTGTTCTC-3'
BAP	5'-GGACCATTCCCACGTCTTCAC-3'	5'-CCTTGTAGCCAGGCCCATTG-3'
OPN	5'- ATGGAAAGCGAGGAGTTGAATG-3'	5'-TGCTTGTGGCTGTGGGTTT-3'
OPG	5'-AGCTGCAGTACGTCAAGCAGGA-3'	5'-TTTGCAAACTGTATTTCGCTCTGG-3'

 Table 1. Primer sequences of the target genes.

described above. Total proteins were obtained by 1% triton X-100 NaCl solution. Total protein content was detected by BCA method (see "Calcification staining and measuring" section). BAP protein concentration was assayed by fluorescent quantitation kit (Sigma, Cat. no. APF) and adjusted by total protein content.

Western blot analysis

This method was used to detect the protein levels of BMP2, OPN, and OPG. The cell treating process and total protein extraction were the same as before. Protein sample (25 ug) was loaded to 10% polyacrylamide gels, electro-blotted, and incubated with monoantibodies of OPN (2 mg/L, room temperature, 2 h) (Chemicon, Cat. no. 14331), OPG(0.4 mg/L, room temperature, 2h) (Chemicon, Cat. no. AB2125P), BMP2 (2 mg/L, 4°C, overnight) (Abcam, Cat. no. ab6285) according to the standard procedure, followed by horseradish persecondary oxidase-conjugated antibodies (PIEREC, Cat. no. 31430; Chemicon, Cat. no. AP132P). The ECL detection system (PIERCE, Cat. no. 34081) was used to detect immunereactive proteins. The films were scanned and analyzed by density analysis system (GS-8000, Bio-RAD). Equal protein loading for western blots was adjusted by immune-blotting for GAPDH. Every protein level was transformed into the ratio of experimental group to control group at the same time point during statistical analysis.

Anti-BMP2 siRNA interfering

Target gene sequence was obtained through CorNucleotide in the NCBI official website (http://www.ncbi.nlm.nih.gov) and the BMP2 mRNA serial number was NM_001202. Four candidate siRNAs and one negative control siRNA labeled with FAM fluorescein were designed and prepared by Shanghai GenePharma. The transfect agent was Lipofectamine2000 (Invitrogen, Cat. no. 11668-019) 8 uL/well; medium was non-serum OptiMEM (GIBCO, Cat. no. 31985). Optimization of the transfect system and the siRNA validities screening were followed the standard procedure. The negative FAM labeled siRNA of 60 nM achieved the highest efficiency according to visual fluorescence brightness. The following experiments were based on this optimized concentration. One of the four candidate siRNAs showed the lowest expression level of BMP2 mRNA (the mean level was 0.19) and the highest inhibiting efficiency (the mean value was 81%). This anti-BMP2 siRNA was chosen as the valid siRNA: 5'-GUGCUAUCUCGAUGCUGUATT-3' and 5'-UACAGCAUCGAGAUAGCACTG-3'. The BMP2 mRNA being interfered was: GTGCTA TCTCGATGCTGTA.

To test the effect of siRNA on mRNA level, cells were divided into two groups: the control group (transfected with negative siRNA) and experimental group (transfected with the valid anti-BMP2 siRNA). After transfection, the cells of both groups were changed to regular medium culture and both were treated with rhIL-6 10 ng/mL. The cellular mRNA level of BMP2, BAP, OPG, and OPN were tested at 12 h and 48 h.

To observe the effect of siRNA on calcification, 50 ng/mL rhIL-6 was used. Cells were divided into three groups: the control group and the experimental group (the same as described before), and the blank group (transfected with negative siRNA but not treated with rhIL-6). The calcium concentrations of cells were measured at baseline, two and four days after treatment.

Statistical analysis

All tests were repeated three times at cell level. The results of all tests were expressed as mean \pm SD. One-way ANOVA was used for data analysis through SPSS 13.0 for windows, and *P* value <0.05 was considered statistically significant.



Figure 2. IL-6 induced calcification of the HUASMCs. (a) The control group was treated without rhIL-6 for six days and showed a negative result of calcium stain by alizarin red S staining. (b) The experimental group was treated with rhIL-6 (50 ng/mL) for the same time period and the extracellular matrix was positively stained with orange. The two arrows indicated the sand-like and the nodular calcium stain in the extracellular matrix, respectively. Both panels were magnified 200 times. (c) Cells were treated with different concentrations of rhIL-6 for different durations as indicated. Calcium concentrations were elevated in a time- and dose-dependent manner. *P<0.05 vs. control of the same time point or vs. the sample of the earlier time point within the same group. **P<0.05 vs. control and sample of the other group at the same time point.

Results

RhIL-6 induced calcification of HUASMCs

The control cells were negatively stained by Alizarin red S on the extracellular matrix (Figure 2a), while the extracellular matrix of the experimental cells was positively stained in two pattern, sand-like stain and nodular stain (Figure 2b). Calcification measure showed that the calcium concentration of the 50 ng/L group at days 6, 9, or 12 was $1.81 \pm 0.03 \text{ mmol/g}$, $2.08 \pm 0.10 \text{ mmol/g}$, and $3.22 \pm 0.18 \text{ mmol/g}$, respectively, and were all elevated compared to the 10 ng/L group at the same time point (P < 0.05). Calcium concentrations of the 10 ng/L group at days 9 and 12 were $0.76 \pm$ 0.02 mmol/g and $1.54 \pm 0.11 \text{ mmol/L}$, respectively, and were both higher than the control group at the same time point (P < 0.05). The calcium concentrations of all three groups showed a time-dependent elevation in this study (Figure 2c).

RhIL-6 induced expression of BAP, OPN, OPG, and BMP2 in HUASMCs

The experimental group (cells treated with IL-6 10 ng/mL) upregulated gene and protein expression of BMP2, BAP, OPN, and OPG sequentially. Compared with the control group, BAP mRNA levels of the experimental group increased at 12 and 24 h and regressed to baseline at 72 h (2.51 \pm 0.11 at 12 h, 1.94 ± 0.03 at 24 h, and 0.90 ± 0.03 at 72 h); BAP protein level only increased at 12 h $(3.96 \pm 0.54 \text{ at } 12 \text{ h}, 1.45 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 24 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.27 \pm 0.46 \text{ at } 1.28 \text{ h}, \text{ and } 1.28$ 0.66 at 72 h). OPN mRNA levels of the experimental group increased at 24 and 72 h (1.03 \pm 0.09 at 12 h, 1.90 ± 0.10 at 24 h, and 3.14 ± 0.32 at 72 h), while the protein level of OPN only increased at 72 h (0.98 \pm 0.32 at 12 h, 1.37 \pm 0.47 at 24 h, and 2.57 \pm 0.43 at 72 h). OPG mRNA level of the experimental group increased at 24 and 72 h (0.99 ± 0.13 at 12 h, 1.77 ± 0.14 at 24 h, and 4.06 ± 0.24 at 72 h); and the protein level increased at 72 h only $(1.25 \pm 0.42 \text{ at } 12 \text{ h}, 1.46 \pm 0.42 \text{ at } 24 \text{ h}, \text{ and } 3.46 \pm$ 0.34 at 72 h), just in the same way of OPN. The mRNA level of BMP2 in experimental group elevated at 12 and 24 h (3.04 \pm 0.07 at 12 h, 1.89 \pm 0.14 at 24 h, and 0.98 ± 0.14 at 72 h). The protein level of BMP2 in experimental group only elevated at 12 h (8.14 ± 0.41 at 12 h, 1.78 ± 0.76 at 24 h, and 1.14 ± 0.10 at 72 h). Part of the data is shown in Figure 3.

Anti-BMP2 siRNA partly reversed the calcification of HUASMCs induced by rhlL-6

The mRNA levels of BMP2 and BAP in the experimental cells at 12 h were 0.02042 ± 0.0069 and 0.1350 ± 0.0093 , respectively. Both were significantly lower than those of the control cells (*P* <0.05) and both were recovered at 48 h, while the mRNA levels of OPG and OPN in the experimental cells were at the same level as the control cells at 12 h and significantly decreased at 48 h to 0.2896 \pm 0.0176 and 0.2533 \pm 0.0211, respectively (*P* <0.05). These data suggest that with siRNA interfering, IL-6 failed to induce BMP2 expression at 12 h and the effect on BMP2 downstream genes could last at least 48 h. The statistical results are shown in Figure 4a.

The calcium concentrations in the cellular layer had no statistical significance among all three groups at baseline. When incubated with rhIL-6 50 ng/mL, the calcium concentrations in the experimental cells increased to 1.95 ± 0.2472 mmol/g protein at day 2 and 2.17 ± 0.2368 mmol/g protein at day 4, which were both lower than those in the control cells (2.95 ± 0.2241 and 2.89 ± 0.3162 , respectively), but higher than those in the blank group cells (1.11 ± 0.2573 and 0.92 ± 0.1073 , respectively) at the same time points (P < 0.05). The statistical results were showed in Figure 4b. These results indicated that BMP2 RNA interfering could partly reverse the calcification in cell matrix induced by IL-6.

Discussion

Extensive exposure to inflammation of organs all over the body is prominent phenomenon in RA. As a pleiotropic cytokine, IL-6 has been demonstrated to be closely associated with the pathogenesis of RA. Circulating IL-6 is involved in a group of conditions caused by the systemic inflammation of RA, including small-vessel vasculitis, anemia, weight loss, general osteoporosis, extensive artery calcification, and accelerated CVD.20-22 These issues are common in RA patients and are also alarming problems in diabetes, CKD, aging, and chronic stress.^{23,24} We thus assumed that IL-6 may play a role in systemic inflammation associated vascular calcification. The relationship between IL-6 and vascular calcification has never been proved directly before, in vitro or in vivo. Since vascular smooth muscle cells are the core and origin of vascular calcification tested by several studies in other fields,^{25–27} we established a system of HUASMCs culturing (Figure 1) to observe the role of IL-6 on vascular calcification in vitro.

The concentration of circulating IL-6 is lower than 10 pg/mL in normal human and is usually above 50 pg/mL in patients with active RA.²⁸ In vitro studies like this would prefer using a high concentration of IL-6 as the stimulator to accomplish the effects in a relatively short time.^{29,30} Data from our preliminary experiment showed that HUASMCs had normal proliferation curves when co-cultured with IL-6 under 100 ng/mL. We therefore chose up to 50 ng/mL of IL-6 as the treating condition in this study. Our research showed that IL-6 50 ng/mL could induce calcium deposition in extracellular matrix of HUASMCs detected by positive Alizarin red S staining. The calcium concentration increased in both time- and dosedependent manners. These results indicate precisely



Figure 3. IL-6 induces the expressions of BAP, OPN, OPG, and BMP2 in the HUASMCs. Cells treated with or without recombinant human IL-6 10 ng/mL served as the experimental (Exp.) group or the control (Con.) group. (a) The mRNA level of every target gene was expressed by the ratio of the experimental group to the control group at the same time point. (b) The proteins detected by western blot were given on the left. The levels of these proteins are shown on the right, which were expressed by the ratio of the experimental group at the same time point. *P < 0.05 vs. control.

that IL-6 can induce calcification in the extracellular matrix of HUASMCs in vitro.

Calcification of the extracellular matrix is an active process carried out by osteoblasts. With the abduction of BMPs, especially BMP2, the precursor cells will differentiate into osteoblasts and express the calcification associated proteins like BAP, OC, OPN, collagen I, and OPG. These proteins will lead into mineral deposition in the extracellular matrix and eventually result in calcification of the bone matrix. Is there any possibility that the cultured HUASMCs trans-differentiated into osteoblasts under stimulating of IL-6? Next, we detected BAP, OPG, and OPN, which are proteins



Figure 4. Anti-BMP2 siRNA partly reversed the IL-6 induced calcification of HUASMCs. Cells transfected with negative siRNA and treated with rhIL-6 served as the control group. Cells transfected with the valid anti-BMP2 siRNA and treated with rhIL-6 served as the experimental group. Cells transfected with negative siRNA but not treated with rhIL-6 served as the blank group. (a) After the BMP2 RNA interfering, rhIL-6 10 ng/mL failed to increase the mRNA levels of BMP2 and BAP at 12 h, and levels of OPG and OPN at 48 h in the experimental group. (b) When observing the calcification effect, 50 ng/mL rhIL-6 was used. The calcium concentration of the experimental group was lower than that of the control group, but higher than that of the blank group at the same time point. *P < 0.05 vs. control. **P < 0.05 vs. control and the bland group.

specifically secreted by osteoblasts. We tested all the three proteins in both mRNA and protein levels. The results showed sequential increasing of them. BAP was expressed in early stage, followed by OPN and OPG. High level of BMP2 was also proved to be induced by IL-6 in both mRNA and protein levels. This mimicked what would happen in osteoblasts.^{31,32}

VSMCs and osteoblasts are both derived from the mesenchymal precursor cells. BMP2 is the most important regulatory factor for osteogenic differentiation and osteoblast proliferation, and is speculated as the key factor promoting calcification of human VSMCs in several circumstances like hyperglycemia and hyperphosphatemia.^{33–36} Several kinds of cells express BMP2, including the bone marrow and mesenchymal stem cells, as well as VSMCs. Based on the results demonstrated above, we thought that BMP2 expressed endogenously by VSMCs initiated osteogenic differentiation of these cells. Pathologic studies revealed that inflammation, uremia, high glucose level, and other conditions could all upregulate the BMP2 expression and ultimately cause calcification.^{31,37,38} However, the BMP2 gene knockout mouse died because of heart defects after gestating for 7–10 days, which makes the gene knockout an impropriate technique in this kind of research.³⁹ Therefore, we established a BMP2 gene posttranscription silence system in VSMCs by small RNA interference technique, to observe the impact of endogenous BMP2 on VSMCs transform and calcification.

Our research showed that the HUASMCs could upregulate the expression of BMP2 in vitro under the stimulating of rhIL-6 and the expression could be effectively silenced by specific anti-BMP2 siRNA. This silence effect on BMP2 and the downstream genes lasted at least 48 h after the siRNA transfect. We believed that this small RNA interference system could serve as a proper method to explore the role of endogenous BMP2. We then tested the role of the transient endogenous BMP2 gene silencing on extracellular matrix calcium deposition. A high dose of rhIL-6 (50 ng/mL) was chosen as the stimulator to get a detectable calcification effect in HUASMCs as quick as possible. According to our data, IL-6 induced calcification could be partly reversed by BMP2 silencing within four days. These results suggested that: (1) the endogenous BMP2 participated in the osteogenic differentiation of VSMCs and the following calcification under IL-6 inducement; and (2) besides the endogenous BMP2, other mechanism might be involved in IL-6 induced calcification of VSMCs.

Vascular calcification is a major life-threatening condition in several diseases, especially in RA. The underling mechanism is not understood clearly. Possible theories include osteogenic differentiation of VSMCs, high serum calcium and phosphorus products deposition, apoptosis vesicle-based calcification, mechanical stress, and oxidative stress.^{33,40,41} Through transient silence of BMP2 of the VSMCs, this study confirmed that endogenous BMP2 and osteogenic differentiation was involved in vascular calcification induced by increased IL-6. Thus, the osteogenic differentiation of VSMCs may be one underlying reason of systemic inflammation related vascular calcification in several situations. Furthermore, serial passage culture of the VSMCs manifested senescent characteristics along with the endogenous BMP2 gene expression.⁴² One study also showed that BMP2 protein was highly expressed at the calcified part of blood vessel along with the prominent phenomenon of apoptosis.⁴³ Another one reported that IL-6 could induce human VSMCs proliferation in vitro.⁴⁴ These studies imply that IL-6 induces VSMCs proliferation, transgenic differentiation, and apoptosis at the same time. This may explain the results of our study. IL-6 might also induce calcification of VSMCs through apoptosis.

However, a few limitations have to be considered. In this study, we used the RNA interfering method and got the transient inhibitory of the BMP2 expression for about 48 h, so the long-term BMP2 inhibitory effect was not observed. Construction of virus vector of anti-BMP2 siRNA will help in long-term silence and a better understanding of the role of endogenous BMP2 on vascular calcification. The HUASMCs obtained from the primary culture method were heterogenic cells which might include smooth muscle cells, mesenchymal stem cells, and perivascular cells at least. The latter two kinds of cells are multi-lineage potential cells. The endogenous BMP2 may induce these cells to differentiate into osteoblasts via autocrine or paracrine effects. But this problem may need a second thought because the primarily cultured cell system is exactly the imitation of the situation in vivo.

Declaration of conflicting interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

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