

# Menaquinone-4 modulates the expression levels of calcification-associated factors to inhibit calcification of rat aortic vascular smooth muscle cells in a dose-dependent manner

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**Abstract.** Vascular calcification (VC) caused by chronic kidney disease (CKD)-mineral and bone disorder is a common complication of CKD. Recent studies have demonstrated that menaquinone-4 (MK-4) is negatively associated with VC in patients with CKD. Furthermore, we have previously shown that runt-related transcription factor 2 (Runx2) is important in the phenotypic transformation process of rat vascular smooth muscle cells (VSMCs), which is the key step for the development of VC. The present study investigated the influence of MK-4 on the phenotypic transformation process of rat VSMCs in order to illustrate its role in the process of VC. Calcification assays were performed to assess the calcified degree of rat VSMCs. Additionally, the genes and proteins related to phenotypic transformation were measured by reverse transcription-polymerase chain reaction and western blotting methods. It was revealed that calcium deposition in the cells was evidently increased with an addition of  $\beta$ -glycerophosphate ( $\beta$ -GP) and could be completely prevented by co-incubation with MK-4 in a dose-dependent manner. Furthermore, the expression of Runx2 in the  $\beta$ -GP-induced VSMCs was inhibited by MK-4. It was also revealed that the expression of SMAD1 and bone morphogenetic protein (BMP)-2 were decreased in the  $\beta$ -GP-induced VSMCs treated with MK-4 in a dose-dependent manner; however, the expression of SMAD7 was increased in the  $\beta$ -GP-induced VSMCs treated with MK-4 in a dose-dependent manner. These observations suggest that MK-4 reduces mineralization by regulating the BMP-2 signaling pathway in order to attenuate the expression of Runx2.

## Introduction

Cardiovascular disease (CVD) is the main cause of mortality in patients with chronic kidney disease (CKD), with a 10-20-fold higher risk compared to the age and sex-matched general population (1). One of the potentially life-threatening cardiovascular (CV) consequences of CKD are vascular calcifications (VCs), which contribute to the high prevalence of CV morbidity and mortality in CKD (2-4). Furthermore, risk factors for the VC of CKD patients include abnormal phosphorus and calcium metabolism, bone disorders, phenotype transformation of vascular smooth muscle cells (VSMCs) and deficiency of endogenous calcification inhibitors including Klotho, matrix Gla protein (MGP), pyrophosphate and fetuin-A (5-7).

It has previously been reported that the pathogenesis of VC is similar to bone formation (8-10). One of the inhibitors of VC is the vitamin K-dependent protein (VKDPs), MGP, which regulates ectopic mineralization. Vitamin K functions as a cofactor in the enzymatic reaction that converts specific glutamate residues in MGP into  $\gamma$ -carboxyglutamate ones, which are needed for its ability to inhibit calcification (11-13). Furthermore, vitamin K has the same role in the process of converting growth arrest-specific-6, which can also prevent VC, to their biologically active forms.

Several previous studies have shown that a suboptimal vitamin K status is common in patients with CKD (14-16). Insufficient levels of vitamin K result in VKDPs that are functionally inactive and could therefore represent a modifiable risk factor for VC in CKD patients. Additionally, vitamin K is a fat-soluble compound and consists of two natural forms, vitamins K1 and K2. Both vitamins function as cofactors for the enzyme  $\gamma$ -glutamylcarboxylase. Menaquinone (MK)-4 is a member of the vitamin K2 family, and there is data of the Rotterdam study (17) hypothesizing that vitamin K2 is of greater importance than K1 for vascular health. Additionally, MK-4 deficiency is a predictor of aortic calcification (18). To the best of our knowledge, there are no previous studies that evaluate the effect of MK-4 on the VSMCs in a  $\beta$ -glycerophosphate ( $\beta$ -GP)-induced VC model and the target genes in the regulation of VSMCs differentiation. Therefore, the primary aim of the present study was to investigate how MK-4 inhibited the VC and to characterize the underlying mechanisms.

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**Key words:** menaquinone-4, vascular calcification, runt-related transcription factor 2, bone morphogenetic protein-2, SMAD1, SMAD7

## Materials and methods

**Cell culture of VSMCs.** Rat VSMCs were isolated from the thoracic aorta tunica media of adult male Sprague Dawley rats (Experimental Animal Center of Hebei Medical University, Hebei, China) adopting the method of explant culture as previously described (19). Briefly, the rats were anesthetized by 400 mg/kg chloral hydrate and sacrificed via cervical dislocation. The thoracic aorta was then removed in gnotobasis. The thoracic aorta was then cut into 1-2 mm<sup>2</sup> pieces following the washing off of any residual blood. The tissue pieces were then cultured in dishes with Dulbecco's modified Eagle's medium (DMEM; Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 15% fetal bovine serum (FBS; Gibco; Thermo Fisher Scientific, Inc.), 4.5 g glucose, 100 µg/ml streptomycin and 100 U/ml penicillin in a 5% CO<sub>2</sub> incubator at 37°C. Cells that migrated from tissue pieces were collected when they reached confluence. Furthermore, the cells were cultured in DMEM replenished with 15% FBS, and the medium was changed twice per week. VSMCs were identified by positive staining of  $\alpha$ -smooth muscle actin (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) and used for all experiments between generations 3-4.

The current study conformed to the Guide for Care and Use of Laboratory Animals as adopted and promulgated by the United National Institutes of Health. All experimental protocols were approved by the Review Committee for the Use of Animal Subjects of Hebei Medical University (Shijiazhuang, China).

**Calcification assays.** In order to induce calcification, the VSMCs were incubated with a calcifying medium, which consisted of growth medium supplemented with 10 mM  $\beta$ -GP (Sigma-Aldrich; Merck KGaA). In some experiments, the cells were exposed to calcifying medium accreted with 10, 25 and 50 µM of MK-4 (Sigma-Aldrich; Merck KGaA). After incubation for 10 days, the cells were washed twice with phosphate-buffered saline and fixed with 95% ethanol. Next, the cells were incubated with 0.2% Alizarin red [(pH 8.3); Beijing Solarbio Science & Technology Co., Ltd., Beijing, China]. Subsequent to washing, an LH50A inverted phase contrast microscope (Olympus Corporation, Tokyo, Japan) was used for visualizing cells and capturing images to record the incidence of induced calcification. NIS-Element F3.0 software was used to capture the images (Olympus Corporation). Subsequently, calcium deposited in the extracellular matrix was treated with 0.6 M HCl for 24 h at 37°C, and the content of calcium in the supernatant was measured with the o-cresolphthalein complexone method using a calcium assay kit (BioSino Biotechnology & Science, Inc., Beijing, China). This was then normalized relative to the protein concentration of the same culture.

**Reverse transcription-polymerase chain reaction (RT-PCR).** Total RNA was extracted from VSMCs by reference methods using TRIzol reagent (Invitrogen; Thermo Fisher Scientific, Inc.) in accordance with the manufacturer's instructions, and quantified by an UV absorbance of 260-280 nm. In total, 2 µg RNA was used for synthesizing cDNA with the RT-for-PCR kit (Clontech Laboratories, Inc., Mountainview, CA, USA) with oligo (dT) primers as recommended in the protocol provided.

cDNA was used as regular RT-PCR template. Furthermore, the internal control was the glyceraldehyde-3-phosphate dehydrogenase (GAPDH) gene. All of the primers used are listed in Table I. PCR was performed using the PCR Master Mix kit (Promega Corporation, Madison, WI, USA) according to the manufacturer's protocol and purified prior to sequencing. The PCR conditions were as follows: Initial incubation for 2 min at 95°C; 35 cycles of 30 sec at 95°C for denaturation; 30 sec at 55°C for annealing; 45 sec at 72°C for extension; and a final extension at 72°C for 5 min. Cycle sequencing was performed with the Dye Terminator Cycle Sequencing Ready Reaction kit (Applied Biosystems; Thermo Fisher Scientific, Inc.). For regular RT-PCR, the PCR products were detected by a 1.5% agarose gel with electrophoresis and visualized by staining with ethidium bromide. Furthermore, the reaction of each sample was repeated once for quality control. The optical density of the band was measured using a Gel Documentation System (CST Biological Reagents Co., Ltd., Shanghai, China) and the final data is expressed as the mRNA level relative to that of GAPDH.

**Western blot analysis.** Total protein was collected from the VSMCs, and the concentrations were measured with the bicinchoninic acid protein assay kit (Sigma-Aldrich; Merck KGaA). Equivalent protein (50 µg) from VSMCs were separated by 10% SDS-PAGE and electrotransferred onto a polyvinylidene fluoride membrane (Amersham; GE Healthcare Life Sciences, Chalfont, UK). Non-specific protein binding was blocked with 5% non-fat dry milk in Tris-buffered saline and Tween-20 [TBS-T; (20 mmol/l Tris-HCl (pH 7.6), 150 mmol/l NaCl and 0.02% Tween-20; Invitrogen, Carlsbad, CA, USA)] by incubating the membrane for 1 h at room temperature with agitation. Anti-mouse runt-related transcription factor 2 (Runx2) primary monoclonal antibody (cat. no. ab76956; Abcam, Cambridge, UK) was incubated with the membrane at a 1:500 dilution in TBS-T and at 4°C overnight with agitation. The secondary antibody (horseradish peroxidase conjugated Goat anti-Mouse Immunoglobulin G; cat. no. 074-1806; KPL, Inc., Gaithersburg, MD, USA) was diluted in TBS-T at 1:2,000 dilution and applied to the membrane, and the reaction was incubated at room temperature for 1 h with agitation. Between each of the three proceeding steps (primary antibody, secondary antibody and visualization) the membrane was washed 3 times with TBS-T for 15 min at room temperature. The membrane was then immediately visualized and analyzed using the enhanced chemiluminescence detection system (Santa Cruz Biotechnology, Inc., Dallas, TX, USA) according to the manufacturer's instructions. Furthermore,  $\beta$ -actin was used as the endogenous control and the experiments were repeated three times.

**Statistical analysis.** Results are presented as the mean  $\pm$  standard deviation, and SPSS 17.0 software (SPSS, Inc., Chicago, IL, USA) was used for the analysis of variance and Dunnett's test.  $P < 0.05$  was considered to indicate a statistically significant difference.

## Results

**MK-4 attenuates calcification in VSMCs induced by  $\beta$ -GP.** The present study investigated the effects of MK-4 on the

Table I. Primers used in reverse transcription quantitative polymerase chain reaction.

Gene	Primer	
	Forward	Reverse
Runx2	5'-CCGCACGACAACCGCACCAT-3'	5'-CGCTCCGGCCCCACAAATCTC-3'
BMP-2	5'-ACTCGAAATTCCCCGTGACC-3'	5'-CCACTTCCACCACGAATCCA-3'
SMAD1	5'-GGTGACTGGGAACGGATCG-3'	5'-TGGTCTTCGGTTCGGAAAGG-3'
SMAD7	5'-CCTCGGAAGTCAAGAGGCTG-3'	5'-CCTCGGAAGTCAAGAGGCTG-3'
GAPDH	5'-CAAGGTCATCCATGACAACCTTTG-3'	5'-GTCCACCACCCTGTTGCTGTAG-3'

Runx2, runt-related transcription factor 2; BMP-2, bone morphogenetic protein-2.

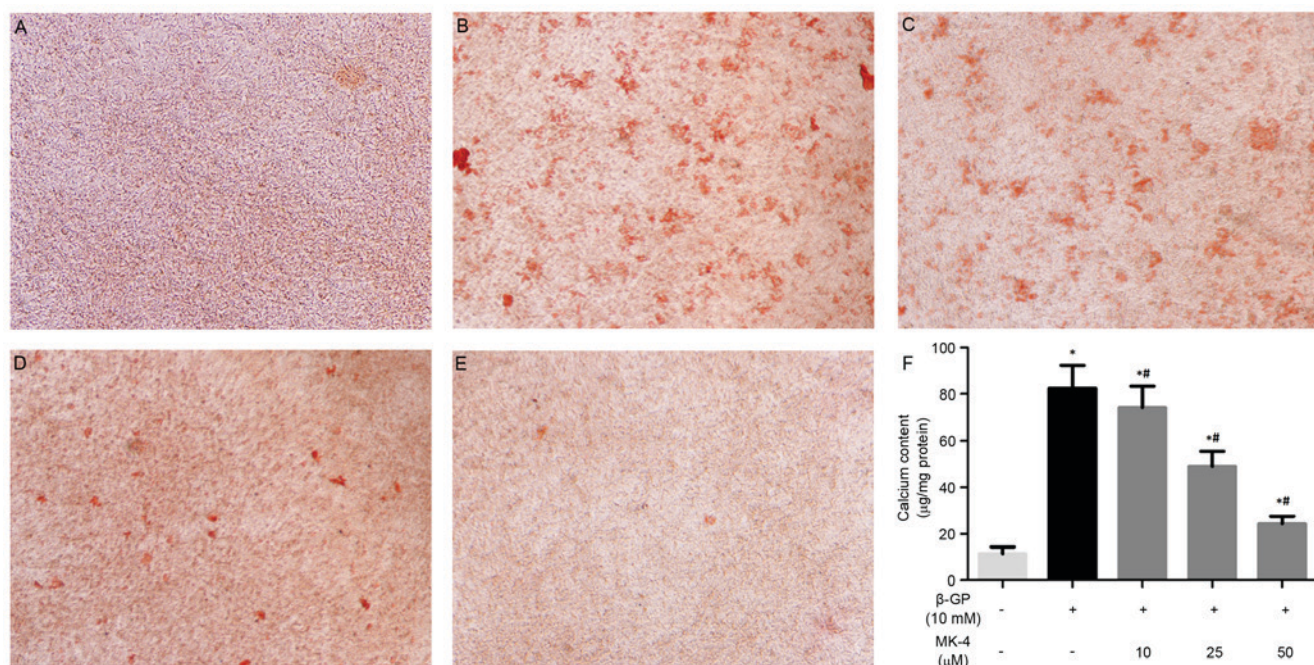


Figure 1. Alizarin red staining was performed on vascular smooth muscle cells incubated under different conditions. (A) control, (B)  $\beta$ -GP, (C)  $\beta$ -GP and 10  $\mu$ M MK-4, (D)  $\beta$ -GP and 25  $\mu$ M MK-4, (E)  $\beta$ -GP and 50  $\mu$ M MK-4 and (F) calcium content. Magnification,  $\times 100$ . \* $P < 0.05$  vs. control; # $P < 0.05$  vs.  $\beta$ -GP.  $\beta$ -GP,  $\beta$ -glycerophosphate; MK-4, menaquinone-4.

calcification of VSMCs induced by  $\beta$ -GP as previously described (20). MK-4 significantly inhibited calcium deposition of VSMCs induced by  $\beta$ -GP at 10 days and in a dose-dependent manner (Fig. 1A-E). In agreement, the quantitative analysis indicated that the calcium content was noticeably reduced in VSMCs maintained in the  $\beta$ -GP + MK-4 medium compared with  $\beta$ -GP medium (Fig. 1F). These results suggest that MK-4 functions in suppressing the calcification of VSMCs induced by  $\beta$ -GP.

#### MK-4 inhibits Runx2 expression of VSMCs induced by $\beta$ -GP.

It has been demonstrated that VSMCs enhanced mineralization under calcifying conditions by phenotype transformation of VSMCs, which is a strictly regulated cellular process that is similar to bone formation (21). Thus, the present study assessed the effects of  $\beta$ -GP on the expression of Runx2 in the presence or absence of MK-4 medium. It was revealed that the expression of Runx2 mRNA and protein in the VSMCs

incubated with the  $\beta$ -GP + MK-4 medium evidently decreased, with the effect remaining enhanced in a dose-dependent manner (Fig. 2).

*Effects of MK-4 on bone morphogenetic protein-2 (BMP-2), SMAD1, SMAD7 mRNA expression of VSMCs induced by  $\beta$ -GP.* RT-PCR was performed to detect the mRNA expression of BMP-2, SMAD1, SMAD7 in VSMCs induced by  $\beta$ -GP in the presence or absence of MK-4. The results revealed that MK-4 significantly reduced BMP-2 and SMAD1 mRNA expression in a dose-dependent manner, while MK-4 markedly increased the expression of SMAD7 mRNA in a dose-dependent manner (Fig. 3). In addition, VSMCs were also treated with  $\beta$ -GP in the presence or absence of 1 mg/ml Noggin, which prevents BMP-2 binding to their receptors. As expected, Runx2 expression was upregulated in VSMCs maintained within  $\beta$ -GP medium compared to the control medium, while the mRNA and protein expression of Runx2

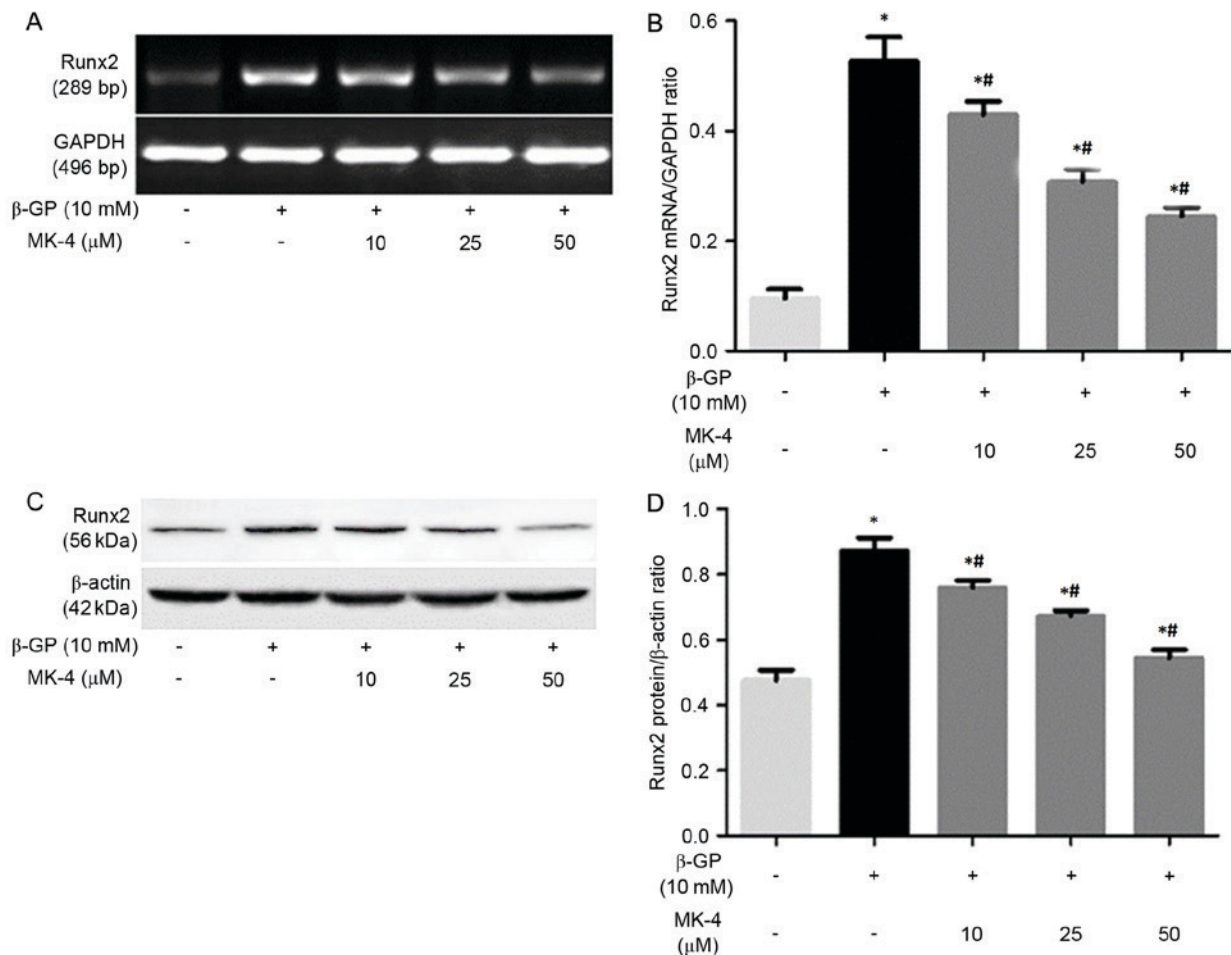


Figure 2. Relative expression levels of Runx2 mRNA and protein. (A) Representative Runx2 mRNA expression levels from vascular smooth muscle cells following treatment with control medium,  $\beta$ -GP or  $\beta$ -GP+MK-4. (B) Quantification of relative expression levels of Runx2. (C) Representative western blots of Runx2 protein in the cells. (D) Quantification of relative protein levels of Runx2. \* $P < 0.05$  vs. control; \*\* $P < 0.05$  vs.  $\beta$ -GP.  $\beta$ -GP,  $\beta$ -glycerophosphate; MK-4, menaquinone-4; Runx2, runt-related transcription factor 2; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

was significantly downregulated ( $P < 0.05$ ; Fig. 4) by noggin. These results indicated that MK-4 partly functions in the modulation of the BMP-2 signaling pathway to inhibit Runx2 expression of VSMCs induced by  $\beta$ -GP.

## Discussion

VC is known to be a high risk factor for mortality in CVD, particularly in patients with CKD (22,23). Although numerous studies have demonstrated that a number of factors are involved in regulating VC, an effective therapy has yet to be found. Accumulated evidence manifested that VSMCs to osteoblast-like cell transdifferentiation (VOT) is important in promoting VC (10,24). Furthermore, the change in a variety of key molecules of progression of VOT may promote the occurrence and development of VC. Pharmacological manipulation of signaling pathways or key molecules to inhibit VOT has been sufficient in the treatment of VC (21). Furthermore, a number of studies have indicated that MK-4 was associated with VC (25-27). The present study investigated the effects of MK-4 on the  $\beta$ -GP-induced calcification processes in rat VSMCs. As predicted, it was identified that adding MK-4 to the medium significantly inhibited calcification induced by  $\beta$ -GP. Furthermore, the results confirmed that

MK-4 is important in the prevention of the  $\beta$ -GP-induced calcification of VSMCs, and may prove useful in the treatment of VC.

However, the mechanisms by which MK-4 inhibited the VC remains unclear. Previous studies have shown that VC is cell-mediated and inactive process, which may result from the osteogenic differentiation of VSMCs and subsequent extracellular matrix mineralization (28-30). Runx2, which is a central transcription factor, is necessary for osteogenic matrix gene expression, bone formation, VSMC calcification and differentiation (22,31,32). Furthermore, the increasing phosphate concentration *in vitro* results in upregulation of Runx2 expression and downregulation of smooth muscle-specific gene expression (10). Furthermore, Runx2 has been shown to be expressed in the calcified vascular lesion of CKD patients (33). Thus, Runx2 has been regarded as the earliest and most specific osteogenic marker of differentiation for the promoting calcification. In the present study, the expression of Runx2 mRNA and protein was evidently upregulated following treatment with  $\beta$ -GP. However, the upregulated expression of Runx2 mRNA and protein were inhibited in cells that were co-cultured in a medium with different concentrations of MK-4, with the effect remaining enhanced in a dose-dependent manner. These results

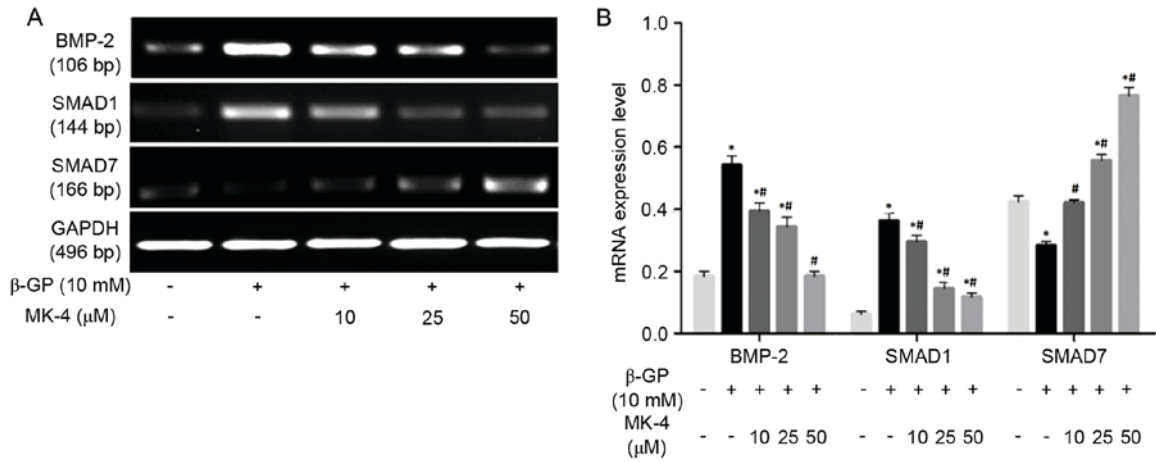


Figure 3. Relative expression levels of BMP-2, SMAD1 and SMAD7 mRNA. (A) Representative mRNA expression following control medium, β-GP or β-GP+MK-4. (B) Quantification of the relative mRNA expression levels of BMP-2, SMAD1 and SMAD7. \*P<0.05 vs. control, #P<0.05 vs. β-GP. β-GP, β-glycerophosphate; MK-4, menaquinone-4; BMP-2, bone morphogenetic protein-2; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

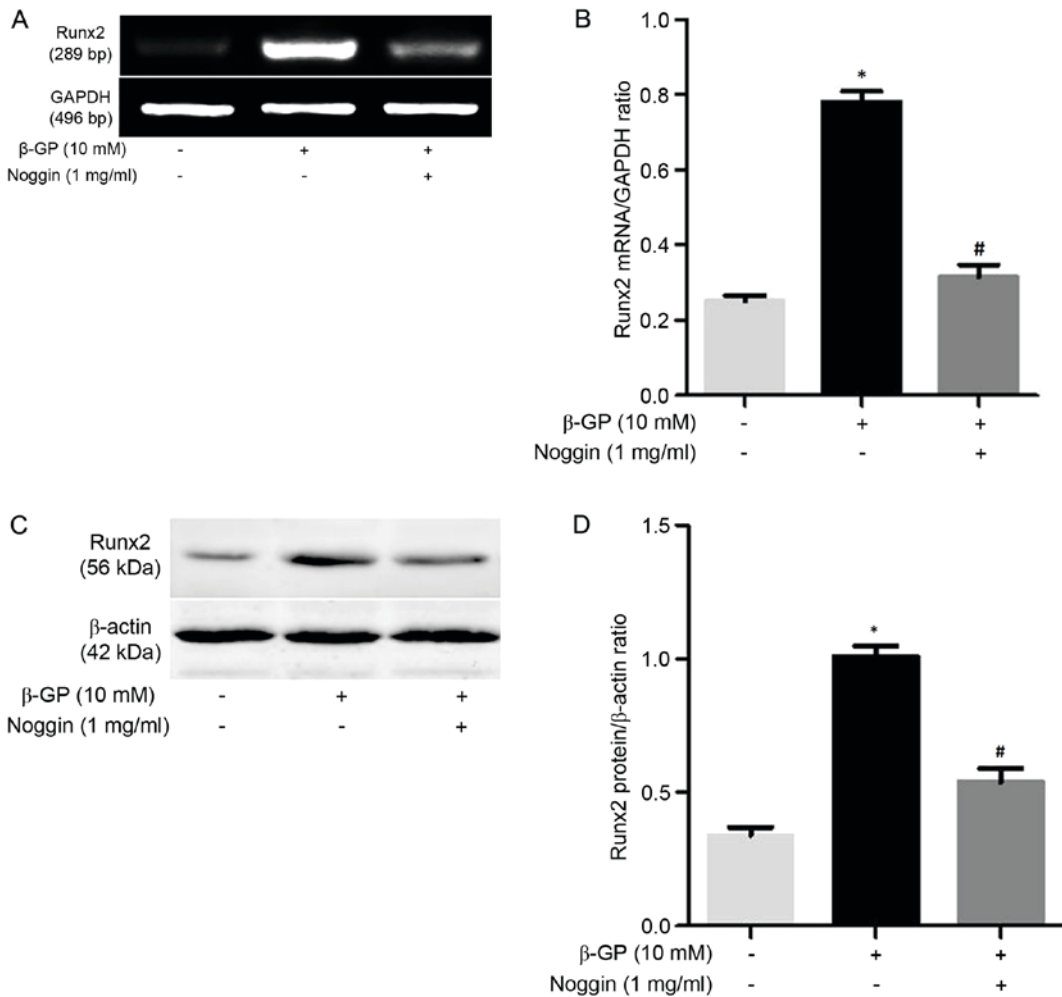


Figure 4. Noggin inhibited β-GP-induced expression of Runx2. (A) Representative mRNA levels of Runx2 in cells following treatment with β-GP and β-GP + noggin. (B) Quantification of the relative mRNA expression levels of Runx2. (C) Representative Western blot analysis of Runx2 in cells following treatment with β-GP and β-GP + noggin. (D) Quantification of the relative protein levels of Runx2. \*P<0.05 vs. control; #P<0.05 vs. β-GP. β-GP, β-glycerophosphate; MK-4, menaquinone-4; Runx2, runt-related transcription factor 2; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

suggested that MK-4 partly reduced VC by reversing the transdifferentiation of VSMCs in calcifying conditions in a dose-dependent manner.

The BMPs are members of the transforming growth factor-β superfamily, and they are important in VC and bone formation (34,35). Additionally, the BMP subfamily

can be further subdivided into several subgroups, including BMP-2/4, BMP-5/6/7/8, BMP-5/6/7 and BMP-9/10 (36-38). A number of studies have shown that BMP signaling is a key regulator of vascular disease (39-41). In order to confirm the effect of BMP signaling in  $\beta$ -GP-induced osteochondrocytic phenotypic change, the effect of Noggin on the Runx2 expression of  $\beta$ -GP-treated VSMCs was investigated. It was shown that Noggin inhibited the  $\beta$ -GP-induced Runx2 expression, suggesting that the BMP pathway is involved in  $\beta$ -GP-induced osteochondrocytic differentiation of rat VSMCs. Next, the present study examined whether the BMP pathway is regulated by MK-4 in VSMCs transdifferentiation induced by  $\beta$ -GP. The results revealed two things. Firstly, MK-4 inhibited the  $\beta$ -GP-induced overexpression of BMP-2 and SMAD1, one of the receptor-regulated SMADs (R-SMADs) that form heteromeric complexes with SMAD4 and then these heteromeric R-SMAD/SMAD4 complexes translocate into the nucleus. Secondly, it also increased the  $\beta$ -GP-regulated down-expression of SMAD7, one of the inhibitory SMADs that antagonize BMP receptor-initiated SMAD signaling by mediating the degradation of receptors and R-SMADs, and suggesting that the inhibitory effect of MK-4 on VSMCs transdifferentiation is mediated, at least in part, through abrogating  $\beta$ -GP-induced activation of the BMP-2 signaling pathway.

In conclusion, MK-4 was capable of validly reducing calcification induced by  $\beta$ -GP in rat VSMCs. Furthermore, MK-4 inhibited the transdifferentiation of VSMCs into osteoblast-like cells by suppressing the expression of Runx2 in a dose-dependent manner. In addition, the downregulated expression levels of the inhibitors BMP-2 and SMAD1 and upregulated expression of the promoter SMAD7 caused by MK-4 in a dose-dependent manner was observed. These observations reveal that MK-4 reduces mineralization by the regulation of the signaling pathway of BMP-2 in order to attenuate the expression of Runx2. Furthermore, the present study may help illuminate the function of MK-4 in the calcification of VSMCs induced by  $\beta$ -GP.

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### Availability of data and materials

All data are fully available without restriction.

### Authors' contributions

LC, JX, and JZ designed the present study and wrote the manuscript. MZ and SZ performed the statistical analysis. VSMC cell culture, calcification assays, reverse

transcription-quantitative polymerase chain reaction and western blotting were performed by SZ and YB.

### Ethics approval and consent to participate

The current study conformed to the Guide for Care and Use of Laboratory Animals as adopted and promulgated by the United National Institutes of Health. All experimental protocols were approved by the Review Committee for the Use of Animal Subjects of Hebei Medical University (Shijiazhuang, China).

### Patient consent for publication

Not applicable.

### Competing interests

The authors declare that they have no competing interests.

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