

Activation of JNKs is essential for BMP9-induced osteogenic differentiation of mesenchymal stem cells

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Although BMP9 is highly capable of promoting osteogenic differentiation of mesenchymal stem cell (MSCs), the molecular mechanism involved remains to be fully elucidated. Here, we explore the possible involvement and detail role of JNKs (c-Jun N-terminal kinases) in BMP9-induced osteogenic differentiation of MSCs. It was found that BMP9 stimulated the activation of JNKs in MSCs. BMP9-induced osteogenic differentiation of MSCs was dramatically inhibited by JNKs inhibitor SP600125. Moreover, BMP9-activated Smads signaling was decreased by SP600125 treatment in MSCs. The effects of inhibitor are reproduced with adenoviruses expressing siRNA targeted JNKs. Taken together, our results revealed that JNKs was activated in BMP9-induced osteogenic differentiation of MSCs. What is most noteworthy, however, is that inhibition of JNKs activity resulted in reduction of BMP9-induced osteogenic differentiation of MSCs, implying that activation of JNKs is essential for BMP9 osteoinductive activity. [BMB Reports 2013; 46(8): 422-427]

INTRODUCTION

Mesenchymal stem cells (MSCs) are multipotent cells present not only in the bone marrow, but in a number of other tissues (1). MSCs can self-renew and differentiate into osteoblasts, chondrocytes, myoblasts or adipocytes (1). Bone morphogenetic proteins (BMPs) belong to the transforming growth factors β (TGF β), and have been proved to play pivotal roles in the processes such as embryogenesis, hematopoiesis, neurogenesis and osteogenesis (2, 3). Several forms of recombinant BMPs, in particular BMP2 and BMP7, have been extensively used to augment bony healing in the clinical setting (4, 5). Though much attention has

been directed to BMP2 and BMP7 in bone regeneration, it is unclear whether these are actually the most osteogenic BMPs

Recent studies have demonstrated that BMP9 is a more potent inducer in inducing osteogenic phenotype of MSCs *in vitro* and *in vivo* (6-9). It was found that BMP9 regulates a distinct set of downstream targets such as Runx2 and CTGF to promote osteogenic differentiation of MSCs (7). We have conducted a comprehensive analysis of seven functional type I receptors and four type II receptors in BMP9-induced osteogenic differentiation of MSCs. The results showed that ALK 1 and ALK 2 are functional type I receptors necessary for BMP9 osteogenic signaling (8). Moreover, we found that TGF β type II receptors BMPRII and ActRII are the type II receptors in BMP9-induced osteogenic differentiation (9). Nevertheless, the signaling mechanism underlying BMP9-induced osteogenic differentiation of MSCs is still largely unknown.

The classic BMPs signaling pathway operates by activation of the transcription factors Smads (2, 10), and there are growing evidences that it can also act through a Smads-independent mitogen activated protein kinases (MAPKs) signaling pathway (2, 11-14). Our recent study has demonstrated that p38 and ERK1/2, two subfamilies of MAPKs, play important roles in regulating BMP9-induced osteogenic differentiation of MSCs (15). Therefore, we spontaneously asked whether JNKs, an important member of MAPKs, is also relevant to BMP9-induced osteogenic differentiation. Herein, we presented evidence that BMP9 can promote activation of JNKs in MSCs. Moreover, inhibition of JNKs led to a decrease in BMP9-induced osteogenic differentiation, suggesting that BMP9 can induce osteogenic differentiation of MSCs through activation of JNKs.

RESULTS

BMP9 induced phosphorylation/activation of JNKs in MSCs

To determine if JNKs can be activated by BMP9 in MSCs, C3H10T1/2 cells were infected with Ad-BMP9 or Ad-GFP with infection efficiency at 40% (Fig. 1A). As illustrated in Fig. 1B, BMP9 significantly increased the levels of phosphorylated JNKs, without altering the total amounts of JNKs proteins. However, SP600125, which is a selective inhibitor for JNKs activation, was able to suppress BMP9-induced phosphorylation of JNKs. Similar results were also observed in C2C12 (Fig. 1C)

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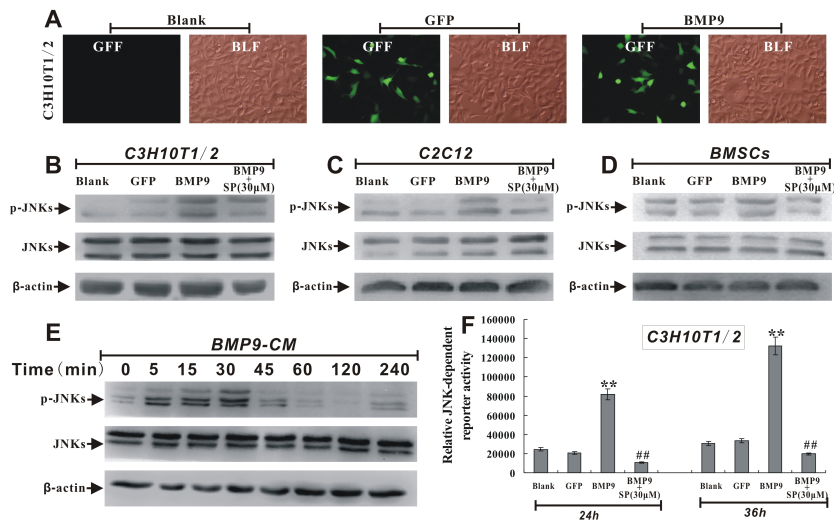


Fig. 1. BMP9 stimulated phosphorylation of JNKs in MSCs. (A) Infection efficiency of Ad-BMP9 in C3H10T1/2 cells. Magnification, 100. GFF: Green Fluorescence Field; BLF: Bright Light Field. (B) C3H10T1/2 cells were infected with Ad-BMP9 or Ad-GFP, at 24 hours post infection, total amount and phosphorylated forms of JNKs were analyzed by western blotting. SP: abbreviation of SP600125. (C) C2C12 cells were infected with Ad-BMP9 or Ad-GFP, at 24 hours post infection, total amount and phosphorylated forms of JNKs was analyzed by western blotting. (D) BMSCs were infected with Ad-BMP9 or Ad-GFP, at 24 hours post infection, total amount and phosphorylation forms of JNKs was analyzed by western blotting. (E) C3H10T1/2 cells were treated with BMP9-CM, total amount and phosphorylated forms of JNKs were analyzed by western blotting at indicated time points. (F) C3H10T1/2 cells were infected with Ad-BMP9, JNKs-dependent reporter activity was quantitatively assessed at 24 and 36 hours post BMP9-stimulation. Data were means \pm SD of three experiments. ** $P < 0.01$ VS GFP; ## $P < 0.01$ VS BMP9.

and BMSCs (Fig. 1D). Moreover, we tested the effect of BMP9-conditioned medium (BMP9-CM) on activation of JNKs in C3H10T1/2 cells. As illustrated in Fig. 1E, an increased level of JNKs phosphorylation was first increased at 5min, and peaked at 30 min post BMP9-CM treatment. These findings indicated that BMP9 was capable of effectively promoting phosphorylation of JNKs in MSCs.

JNKs can phosphorylate and then activate transcription factor AP-1 (16). Therefore, using a JNKs-dependent AP-1 promoter reporter pBG2-AP1-Luc, which contains JNKs-responsive elements and reflects the level of JNKs activation, we found that BMP9 was capable of increasing JNKs-regulated AP-1 transcription activity (Fig. 1F). However, SP600125 was able to effectively inhibit BMP9-induced reporter activity. These data implied that BMP9 can activate JNKs, and subsequently induce JNKs-dependent transcription activity. Together, these above results strongly suggested that BMP9 can effectively induce activation of JNKs in MSCs.

BMP9-induced osteogenic differentiation of MSCs was dramatically inhibited by SP600125, a selective inhibitor of JNKs

Next, to further document the detail role of JNKs in BMP9-induced osteogenic differentiation of MSCs, C3H10T1/2 cells were exposed to BMP9 in the presence of varying concentrations of SP600125 (0, 10, 20 and 30 μ M). SP600125 was able to inhibit BMP9-induced ALP activity of C3H10T1/2 cells mostly in a dose-dependent manner (Fig. 2A and B). Similar phenomena were also observed in C2C12 and BMSCs (Fig. 2C).

Moreover, we found that SP600125 treatment resulted in a significant decrease in BMP9-induced matrix mineralization (Fig. 2D), as well as OCN protein expression of MSCs (Fig. 2E). Collectively, these results strongly implied us that inhibition of JNKs can suppress BMP9-induced osteogenic differentiation of MSCs.

It has been evidenced in our previous studies that Id1, Id2, Id3 and Runx2 are targets of BMP9, and are critical to BMP9-induced osteogenic differentiation of MSCs (7). Using semi-quantitative RT-PCR analysis, we found that BMP9-induced gene expression of Id1, Id2, Id3 and Runx2 was strongly decreased by SP600125 (Fig. 2F). We further examined the Runx2 protein expression by western blotting. As shown in Fig. 2G, BMP9-induced Runx2 protein expression was decreased by SP600125 treatment. Next, using a commonly used Runx2-regulated reporter (p6xOSE-Luc), we found that SP600125 was able to inhibit BMP9-induced p6xOSE-Luc activity, which contains Runx2-responsive elements and reflects Runx2 transcription activity (Fig. 2H). These findings revealed that inhibition of JNKs suppressed BMP9-induced activity of Runx2, a key transcription factor associated with osteoblast differentiation.

Blocking of JNKs activity decreased BMP9-induced activation of Smads signaling

We next sought to probe the possible mechanism behind the effects of JNKs on BMP9-induced osteogenic differentiation of MSCs. Our previous study has demonstrated that p38 and ERK1/2 are likely to regulate BMP9-induced osteogenic differ-

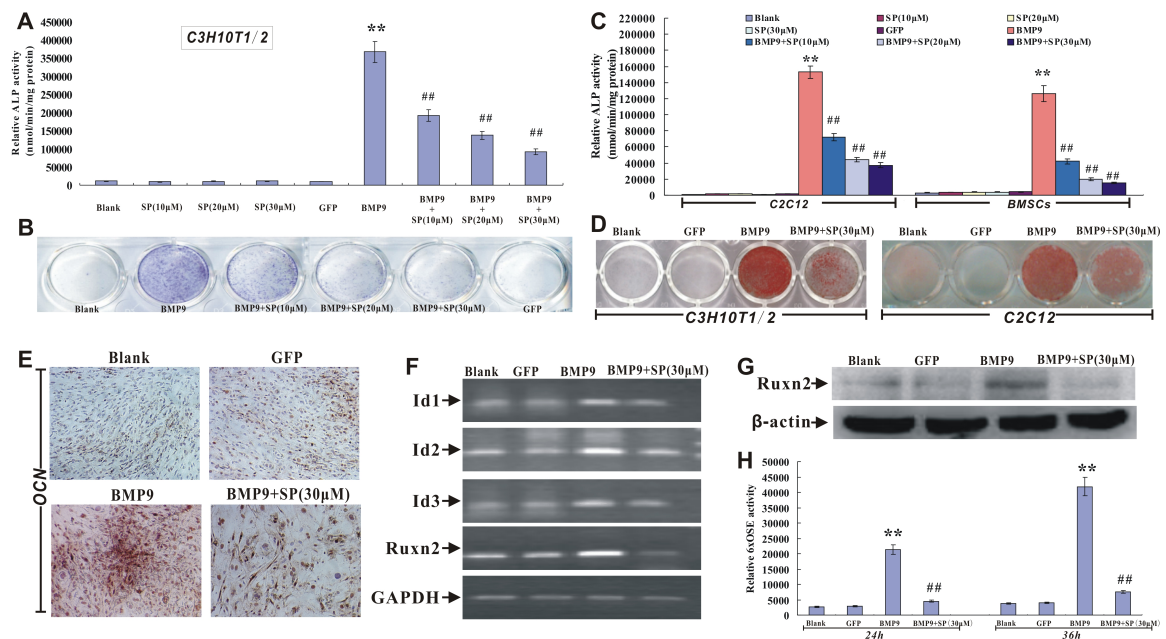


Fig. 2. Inhibition of JNKs activity suppressed BMP9-induced osteogenic differentiation of MSCs. (A) and (B) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (0, 10, 20 and 30 μM). ALP activity was assessed by quantitative assay and staining assay at 7 days post BMP9-stimulation. Data were means ± SD of three experiments. **P < 0.01 VS GFP; ##P < 0.01 VS BMP9. (C) C2C12 and BMSCs cells were infected with Ad-BMP9 in the presence of SP600125 (0, 10, 20 and 30 μM). ALP activity was quantitatively assessed at 7 days post BMP9-stimulation. Data were means ± SD of three experiments. **P < 0.01 VS GFP; ##P < 0.01 VS BMP9. (D) C3H10T1/2 and C2C12 cells were infected with Ad-BMP9 in the presence of SP600125 (30 μM), matrix mineralization was assessed by Alizarin Red S staining at 21 days post BMP9-stimulation. (E) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (30 μM). Osteocalcin (OCN) expression was assessed at 12 days by immunocytochemical staining post infection. Magnification, 100. (F) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (30 μM), the gene expression level of Id1, Id2, Id3 and Runx2 was assessed by RT-PCR at 24 hours post infection by RT-PCR. (G) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (30 μM), the protein expression level of Runx2 was assessed by western blotting at 24 hours post infection. (H) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (30 μM), Runx2-regulated reporter activity was quantitatively assessed at 24 and 36 hours post infection. Data were means ± SD of three experiments. **P < 0.01 VS GFP; ##P < 0.01 VS BMP9.

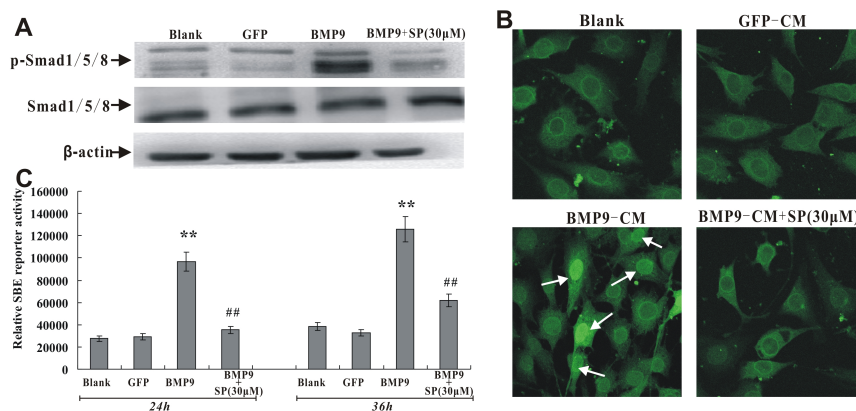


Fig. 3. Blocking of JNKs led to suppression on BMP9-activated Smads signaling in MSCs. (A) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (30 mM), total amount and phosphorylated forms of Smad1/5/8 was analyzed by western blotting. (B) C3 H10T1/2 cells were treated with BMP9-CM in the presence of SP600125 (30 mM), nuclear translocation of Smad1/5/8 were detected by immunocytochemistry staining at 4 hours post BMP9-stimulation, Magnification, 100. (C) C3H10T1/2 cells were infected with Ad-BMP9 in the presence of SP600125 (30 mM), SBE reporter activity was quantitatively assessed at 24 and 36 hours post infection. Data were means ± SD of three experiments. **P < 0.01 VS GFP; ##P < 0.01 VS BMP9.

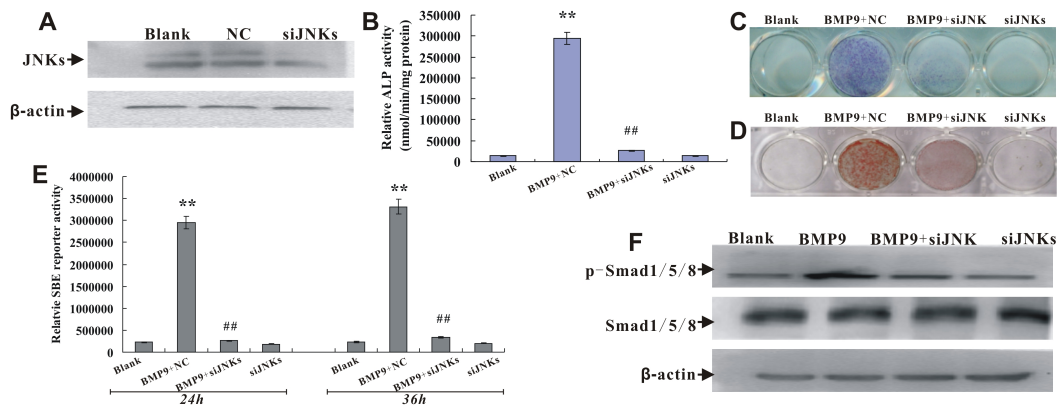


Fig. 4. Silencing of JNKs led to reduction on BMP9-induced osteogenic differentiation of MSCs. (A) Effective knockdown of JNKs expression in C3H10T1/2 by RNAi. NC, negative control. (B) and (C) After achieving effective knockdown of JNKs, ALP activity was assessed at 7 days post BMP9-stimulation by quantitative assay and staining assay. Data were means \pm SD of three experiments. $**P < 0.01$ VS Blank; $^{##}P < 0.01$ VS BMP9+NC. (D) After achieving effective knockdown of JNKs, matrix mineralization was detected by Alizarin Red S staining at 21 days post BMP9-stimulation. Magnification, 100. (E) After achieving effective knockdown of JNKs, SBE reporter activity was quantitatively assessed at indicated time points post BMP9-stimulation. Data were means \pm SD of three experiments. $**P < 0.01$ VS Blank; $^{##}P < 0.01$ VS BMP9+NC. (F) After achieving effective knockdown of JNKs, total amount and phosphorylated forms of Smad1/5/8 was analyzed by western blotting at 24 hours post BMP9-stimulation.

entiation of MSCs through influence on Smads signaling (15). Therefore, we asked whether BMP9-activated Smads signaling was also influenced by JNKs. We found that BMP9-induced phosphorylated Smad1/5/8 was strongly decreased by SP600125 (Fig. 3A). Moreover, using immunochemical staining assay, we showed that SP600125 disrupted translocation of Smad1/5/8 to the nucleus. Lastly, using a BMPs responsive Smad1/5/8 reporter, p12xSBE-Luc (8, 9), we demonstrated that SP600125 was able to neutralized BMP9-induced transcription activity of Smad1/5/8 (Fig. 3C). These results implied us that JNKs are likely to regulate BMP9-induced osteogenic differentiation of MSCs partly through influence on canonical Smads signaling.

Gene silence of JNKs led to a decrease in BMP9-induced osteogenic differentiation of MSCs

Using specific inhibitor for JNKs, we found that inhibition of JNKs activity decrease d BMP9-induced osteogenic differentiation of MSCs. To confirm that the effects of the inhibitor was truly due to JNKs inhibition and not a nonspecific drug effect, we employed adenovirus expressing siRNA targeted JNKs to infect C3H10T1/2 cells. Western blotting was carried out to assess the inhibitory efficiency of the siRNA on expressions of JNKs (Fig. 4A). Then, the effect of JNKs knockdown on BMP9-induced osteogenic differentiation of MSCs was assessed. As shown in Fig. 4B-D, gene silence of JNKs diminished BMP9-induced ALP activity and matrix mineralization of C3H10T1/2 cells. Furthermore, knockdown of JNKs effectively decreased BMP9-activated Smads signaling, leading to reduction of BMP9-induced Smad1/5/8 transcriptional activity (Fig. 4E) and less phosphorylation of Smad1/5/8 (Fig. 4F). Collectively, these above results implied us that JNKs may play a private role in regulating BMP9-induced osteogenic differentiation

and Smads signal activation.

DISCUSSION

BMP9 was originally isolated from fetal mouse liver and is a potent stimulant of hepatocyte proliferation (17). Other roles of BMP9 include inducing the cholinergic phenotype of embryonic basal forebrain cholinergic neurons (18), regulating glucose and lipid metabolism in liver (19), and maintaining homeostasis of iron metabolism (20). Also, BMP9 is a potent inducer for osteogenic differentiation of rat dental follicle stem cells (21). In previous studies, BMP9 has been proved to play a critical role in the processes by which MSCs undergo commitment to the osteoblast lineage (6-9). However, little is known about the downstream signaling pathway(s) involved.

In addition to the Smad1/5/8 transcription factors, MAPKs are also involved in BMPs osteogenic signaling transduction (11-14). Recently, we have reported that p38 and ERK1/2 MAPKs act in opposition to regulate BMP9-induced osteogenic differentiation of MSCs (15). In this current study, we examined the ability of JNKs, another important member of MAPKs, in regulating BMP9-induced osteogenic differentiation of MSCs. The results obtained here indicated that BMP9 was able to activate JNKs. Importantly, blocking of JNKs activity led to reduction on BMP9-induced osteogenic differentiation of MSCs.

JNKs belong to the MAPKs family, and consist of ten isoforms derived from three genes: JNK1, JNK2 and JNK3 (16). JNKs can be activated by several stimuli, such as cytokines, ultraviolet irradiation, and heat shock. The physiological functions of JNKs in osteogenic differentiation and bone formation have been investigated previously (11,13,14). However, the obtained results are controversial, with some studies suggesting a stimulatory role of

JNKs in osteogenic differentiation and bone formation (11,13), and others proposing that JNKs is inhibitory (14). Although these studies about the precise role of JNKs in skeletal development didn't lead to complete unanimity, it is well accepted that JNKs plays a functional role in osteogenesis and bone metabolism.

In addition to Smads, MAPKs can also be activated by BMPs stimulation (2,11-14), which represents an important mechanism for non-Smads pathway(s) of BMPs signaling. It has been described by various studies that JNKs can be activated by BMPs and lead to various effects depending on the cell context (11,14,22,23) However, the exact role of JNKs on BMPs-induced osteogenic differentiation is diverse and disputable. By now, the studies about JNKs on BMPs-induced osteogenesis were mainly focus on BMP2, and these studies did not reach on consensus. Several studies agree on the notion that JNKs play a positive role in BMP2-induced osteogenic differentiation (11,13). However, other study obtained opposite results, showing that JNKs have a negative role in BMP2-induced osteogenic differentiation (14). All these experiments did not ascertain the exact role of JNKs in BMPs-induced osteogenic differentiation and bone formation because of the controversial results. However, it is convincingly supported that JNKs play essential roles in regulating osteogenic activity of BMPs, with positive or negative effects. In the current study, we found that BMP9 was capable of activating JNKs to induce osteogenic differentiation of MSCs, and supposed that JNKs are positive regulatory components in BMP9 osteoinductive activity.

In conclusion, we have found that BMP9 can activate JNKs in the induction of the osteogenic differentiation of MSCs. Notably, using specific inhibitor and siRNA for JNKs, we found that activation of JNKs are essential for BMP9-induced osteogenic differentiation of MSCs. Furthermore, we found that JNKs are likely to regulate BMP9-induced osteogenic differentiations of MSCs via influence on Smads signaling. This knowledge will provide insights into the molecular mechanisms by which BMP9 induces osteogenic differentiation of MSCs.

MATERIALS AND METHODS

Cell culture and chemicals

C3H10T1/2, C2C12 and HCT116 cells were obtained from ATCC and maintained in complete DMEM supplemented with 10% fetal bovine serum and 100 units/ml streptomycin/penicillin at 37°C in a humidified atmosphere of 5% CO₂.

Isolation of primary bone marrow stromal cells (BMSCs)

A single-step primary BMSCs purification method using adhesion to cell culture plastic was employed as described in the reference (8).

Construction of recombinant adenoviruses

Recombinant adenovirus expressing BMP9 (Ad-BMP9) were generated previously (6,15). Recombinant adenoviruses expressing siRNA targeted JNKs (AdR-si-JNKs) was kindly provided

by Dr. Tong-chuan He of University of Chicago Medical Center.

Preparation of BMP9-conditioned medium

BMP9-conditioned medium (BMP9-CM) were prepared as described (15). Briefly, HCT116 cells were infected with an optimal titer of Ad-BMP9. At 24 hour after infection, the culture medium was changed to serum-free DMEM. Conditioned medium was collected at 48 hrs after infection and used immediately

Determination of ALP activity

ALP activity was assessed by a modified Great Escape SEAP chemiluminescence quantitative assay and a histochemical staining assay as described (15).

Alizarin Red S staining

Matrix mineralization was detected by Alizarin Red S stain, as described previously (15). The staining of matrix mineralization was recorded under bright field microscopy.

Western blotting

Western blotting was performed as previously described (15). Primary antibodies were obtained from Santa Cruz, as follows: anti-phosphor-JNKs, anti-JNKs, anti-phosphor-Smad1/5/8, anti-Smad1/5/8, and anti- β -actin.

Immunocytochemical staining

Immunocytochemical staining was performed as previously described (15). The presence of the expected protein was visualized by DAB staining and examined under a microscope.

RNA isolation and semiquantitative RT-PCR analysis

Total RNA was isolated using TRIZOL Reagents. Total RNA was used to generate cDNA templates by RT reaction with hexamer and Superscript II RT. Semiquantitative RT-PCR was carried out as described previously (15).

Immunofluorescence staining

Immunofluorescence staining was performed as previously described (15). Fluorescence signal was recorded under a fluorescence microscope.

Luciferase reporter assay

Luciferase reporter assay was performed as previously described (15). Cells were seeded in 25 cm² cell culture flasks and transfected with 2 mg per flask of luciferase reporter plasmid using Lipofectamine. At 16 hr after transfection, cells were replated to 24 well plates. Cells were infected with Ad-BMP9 in the presence of SP600125 or co-infected with Ad-BMP9 and AdR-si-JNKs. Cells were lysed and subjected to luciferase assays at indicated time points.

Statistical analysis

Data are expressed as means \pm SD. Statistical analysis was performed using SAS (version 8.1; SAS Institute, Cary, NC), $P <$

0.05 was taken as the level of significance.

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