A novel nano-sized bioactive glass stimulates osteogenesis via the MAPK pathway

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The aims of this study were to compare the osteogenic effects of a novel nano-sized bioactive glass (BG) and a traditional micron-sized BG, and to verify whether mitogen-activated protein kinases (MAPKs) are involved and play a part in BG's osteoblast gene activation. It was found that the effect of a nano-sized BG on MAPK phosphorylation is better than the traditional 45S5 BG. We prepared extractions of the novel nano-sized 58S BG and traditional 45S5 BG and compared their effect on osteoblast-like cells’ (MG-63 cell) proliferation, osteogenic gene and protein expressions, matrix mineralization and MAPK activation. We further investigated the signal transducing effect of the MAPK pathway on the BG's osteogenic gene activation. Our results showed that nano-58S extraction enhanced the MG-63 cells’ proliferation and osteogenic gene expressions of alkaline phosphatase (ALP), collagen type I (Col I), bone sialoprotein (BSP), Runt-related transcription factor 2 (Runx2) and osteocalcin (OCN). The results of ELISA showed more Col I and OCN protein production in the BG groups than in the control group. Greater mineralized nodule formation was found in the nano-58S BG group using alizarin red S staining. We also found that nano-58S and 45S5 BG activated MAPKs, specifically the ERK and p38 pathways, using western blotting. After blocking the ERK or p38 pathway, real-time PCR showed the osteogenic gene activation induced by the BG extractions was inhibited. Blocking the ERK pathway induced a more obvious inhibitory effect on the genes’ normal expression and activation. The significance of this study is that we found that the ERK and p38 MAPK pathways are involved and play an important part in BG's osteogenic gene activation. The effect of the nano-sized BG on MAPK phosphorylation, osteogenic gene activation, and osteoblast differentiation and mineralization is better than the traditional 45S5 BG.

1. Introduction

The demands on osteoinductive biomaterials have been rapidly growing in recent years due to bone defects, which are common, caused by trauma, post-cancer surgery, periodontal disease and other skeletal diseases. Bioactive glass (BG) is a kind of highly biocompatible, osteoinductive and osteoconductive calcium silicate-based biomaterial. Since it was invented in the 1970s, BG has been used in the clinical treatment of periodontitis, maxillary cystic bone defects and implantation for almost 30 years and proven to have strong direct bonds to bone tissue and excellent osteogenic effects. Recently, several investigators have found that nano-sized sol–gel BG possesses higher specific surface area, better bioactivity and induces more osteoblast adhesion compared to traditional micron-sized BG particles.

Studies have proven that BG regulates the expression of several genes related to cell proliferation, differentiation, and bone matrix formation such as alkaline phosphatase (ALP),
replacement of primary osteoblast cells is widely used in studies on osteogenesis.21,22 

Besides, due to their involvement in the regulation of cells’ proliferation, differentiation, and apoptosis, the function and performance of MAPKs are affected by cell viability. The expression of MAPKs often changes because of different passages’ cell viability and individual differences. Therefore, it is difficult to investigate the real and stable performance of MAPKs in the different passages of primary osteoblasts. On the other hand, because of their infinite passage capacity, MG-63 cells show similar and stable viability and function performance like osteoblasts. The stable and repeatable osteogenesis-related MAPK activation can be observed in this cell line.

The aim of the present study is to verify two hypotheses: (1) MAPKs are involved and play a role in BG’s regulation of osteoblasts’ gene expression; and (2) the effects of nano-sized 58S BG on osteogenic gene up-regulation and MAPK phosphorylation are higher than the traditional 45S5 BG.

2. Materials and methods

2.1 Characteristics of BG

Nano-58S BG (58% SiO2, 33% CaO and 9% P2O5 (w/w)) and traditional melt 45S5 BG (45% SiO2, 24.5% CaO, 6% P2O5 and 24.5% Na2O (w/w)) were prepared by the National Engineering Research Center for Human Tissue Restoration & Reconstruction, South China University of Technology (Guangzhou, China). The preparation methods for the novel nano-58S BG were described previously.22 Both of the bioactive glasses were mainly composed of amorphous silicate phases.24,25 The surface morphology of the particulates was observed by field emission scanning electron microscopy (FE-SEM) (S4800, JEOL, Tokyo, Japan).

2.2 BG extraction

The BG particulates were sterilized in a dry heat oven at 180 °C for 4 h. The particulates were prepared by incubating nano-58S or 45S5 BG in Dulbecco’s Modified Eagle Medium (DMEM) (Gibco, Gaithersburg, MD, USA) at a concentration of 1 mg mL−1 at 37 °C for 24 h. The mixtures were centrifuged and the supernatant was filtered through a 0.22 μm filter (Millipore, Billerica, MA, US). All BG extractions were supplemented with 10% fetal bovine serum (Hyclone, Logan, UT, USA), 100 mg mL−1 streptomycin, 100 U mL−1 penicillin and 2 mM L-glutamine. The pH of the BG extractions was adjusted to 7.4 for cellular experiments. The concentrations of Si, Ca, and phosphorus (P) in the extracts were detected using inductively coupled plasma optical emission spectroscopy (ICP-OES) analysis (ICAP 6300, Thermo Electron Corporation, Waltham, MA, USA).

2.3 Cell proliferation assay

The MG-63 osteoblast-like cells used in this study were obtained from the American Type Culture Collection (ATCC No. CRL 1427). The cells were cultured with the extractions of nano-58S BG, 45S5 BG and growth medium (control group) for 10 days. Cell proliferation was determined using the methyl-thiazol-tetrazolium (MTT) (Amresco, Solon, OH, USA) method.

2.4 Real-time polymerase chain reaction (PCR)

The total RNA of the MG-63 cells was extracted using Trizol® reagent (Invitrogen, Carlsbad, CA, US) and reversely transcribed into complementary DNA (cDNA) by reverse transcriptase (Promega, Madison, WI, USA) according to the manufacturers’ instructions.

Real-time PCR was performed using a Faststart Universal SYBR Green PCR Master mix (Roche, Indianapolis, IN, US) and Applied Biosystem 7500 thermal cycler (ABI 7500, Applied Biosystems, Foster City, CA, US). Real-time PCR was performed with primers designed by Primer 5.0 according to the cDNA sequences available in the GenBank database for ALP (F: 5’-AGCACCTCCACTTTCATCTGGAA-3’, R: 5’-GAGACCCAATAGGTAGTCCACATG-3’), collagen I (Col I, F: 5’-CGAGACGATCCCACCAATCCC-3’, R: 5’TGTGCAGACCGCAGAT-3’), Runx2 (F: 5’-ACCAGAGAGCAGACAGAAGAT-3’, R: 5’-AGGAAATGCGCCTTAAATCAGT-3’) and OCN (F: 5’-AGGGACGAGGATTAGTG-3’, R: 5’-CCTGAAGAAGGGTTGGT-3’). Glyceraldehyde 3-phosphate dehydrogenase (GAPDH, F: 5’-CAACGGATTTGGTCGTATTGG-3’, R: 5’-GCACAATATCCCTTACCGAGGT-3’) was used for the quantification of relative gene expression. The thermal cycling conditions were as follows: 50 °C for 2 min and 95 °C for 10 min, and 40 cycles of 94 °C for 15 s and 60 °C for 1 min.

2.5 Immunocytochemical assay of Runx2 and OCN

Cells were washed with phosphate buffer solution (PBS) and fixed with 4% paraformaldehyde (Sigma, St. Louis, MO, USA). After being punched with 0.1% Triton X-100 and immersed in 3% H2O2, the cells were blocked with 0.5% goat serum. Afterwards, rabbit anti-human Runx2 and OCN primary antibody (dilution 1:200) (Promega, Madison, WI, USA) was added overnight at 4 °C. Subsequently, the cells were incubated with goat anti-rabbit secondary antibody and were subject to enzymatic immunohistochemical staining using a broad-spectrum immunoperoxidase diaminobenzidine kit (Zhongshan Bioengineering, Beijing, China). Finally, the cells were counterstained with hematoxylin and visualized.

2.6 Enzyme-linked immunosorbent assay (ELISA) of Col I and OCN

The MG-63 cells were cultured with the extractions of nano-58S BG, 45S5 BG and growth medium (control group) for 4 days and 7 days. The media were collected and stored at −80 °C until use. Protein expression levels of Col I and OCN in the culture supernatant were assayed using a human Col I and OCN enzyme-linked immunosorbent assay kit (R&D Systems, Minneapolis, MN) according to the manufacturer’s instructions. The protein concentration of OCN was calculated from a calibration curve using human Col I and OCN concentrations as the standard.
2.7 Matrix mineralization assay

The MG-63 cells were cultured in the BG extractions with an osteogenic medium (10 mmol L⁻¹ β-glycerophosphate, 10 mmol L⁻¹ dexamethasone, and 50 mg L⁻¹ ascorbic acid) for 28 days. The cells cultured in the growth medium were used as the non-osteogenic control group; those cultured in the growth medium with the osteogenic medium were assessed as the osteogenic control group. For the matrix mineralization assay, the cells were rinsed 3 times with PBS and fixed in 4% paraformaldehyde for 20 minutes. The cells were washed using deionized water 3 times and then stained with 40 mmol L⁻¹ alizarin red S solution (pH 4.2) (Sigma-Aldrich, St. Louis, MO, US) and observed for mineralized nodules.

For matrix mineralization semi-quantification, 1 mL of 100 mM cetylpyridinium chloride (Sigma-Aldrich, St. Louis, MO, US) was used to dissolve the dried alizarin red S stained cellular matrix. The optical density was detected using the absorbance at 562 nm.

2.8 Extraction of proteins and western blotting

Cells were washed with ice-cold PBS and then lysed using RIPA cell lysis buffer (Cell Signaling Technology, Beverly, MA, US). Then the mixtures were centrifuged at 14 000g for 20 min at 4 °C. The concentration of protein in the supernatant was determined using a BCA Protein Assay Kit (Pierce, Rockford, IL, USA). For western blot analysis, protein samples (20 μg) underwent 10% SDS-PAGE (electrophoretic migration at 100 V) and were electrotransferred onto polyvinylidene fluoride (PVDF) membranes. After being incubated in a blocking solution (50 mM Tris/HCl, 200 mM NaCl, 0.2% Tween 20, and 5% nonfat dried milk), the PVDF membranes were incubated in the presence of 1 : 1000 diluted primary monoclonal rabbit anti-human phosphorylated (p-)ERK, anti-human phosphorylated (p-)JNK, anti-human phosphorylated (p-)p38 (Cell Signaling Technology) and β-actin (Santa Cruz, CA, USA) antibodies at 4 °C overnight. After washing with TRIS-buffered saline (0.2% Tween 20), the membranes were incubated with 1 : 10 000 diluted mouse anti-rabbit secondary antibody (Cell Signaling Technology). The immunoreactive bands were visualized using an ECL detection reagent (Thermo Fisher, Waltham, MA, US), exposed to Kodak X-ray films and quantitatively analyzed using ImageJ 1.38(NIH).

3. Statistical analysis

Each experiment was run in triplicate, and each result was averaged using at least three parallel samples and measurements. All data are expressed as mean ± standard deviation (SD). Variance and statistical significance were determined using Student’s t-test using SPSS Statistics 17.0 software. Differences were considered significant at P < 0.05.

4. Results

4.1 Characteristics of BG

FE-SEM showed that the nano-58S BG particulates were regular and spherical with sizes of approximately 20 nm (Fig. 1(a)), while the shape of the 45S5 BG particulates was irregular with sizes ranging between 2 and 15 μm (Fig. 1(b)).

According to the results of the ICP-OES analysis, the highest concentrations of Ca and Si were detected in the nano-58S BG extraction compared to the control and 45S5 BG groups. The concentration of P was lower in the nano-58S BG and 45S5 BG extractions than in the control group (Fig. 1(c)).

4.2 Effect of BG extractions on MG-63 cells proliferation and osteogenic genes expression

The MTT results showed that BG enhanced cell proliferation (Fig. 1(d)). The OD₄₉₀ values of the two BG groups were both significantly higher than that of the control group (P < 0.05).

The mRNA expression of ALP, Col I, Runx2 and OCN are shown in Fig. 1(e)–(h). Compared with the control group, ALP gene expression was significantly up-regulated by nano-58S and 45S5 BG extractions at 2 and 4 days, and the expression in the nano-58S group was significantly higher than that of the 45S5 group at 4 days. The mRNA expression of Col I at 4 days and 7 days, and Runx2 at 2, 4 and 7 days in the nano-58S group was significantly higher than those of the 45S5 group and the control group. 45S5 BG up-regulated Runx2 gene expression significantly at 2 days and 4 days, but with no difference at 7 days. Nano-58S BG induced significantly higher OCN gene expression at 4 days and 7 days than 45S5 BG and the control.

The immunocytochemical assay of Runx2 at day 4 (Fig. 2(a)) and OCN at day 7 (Fig. 2(b)) showed the most intense positive staining of both proteins in the nano-58S group. Moderate and negative staining were found in the 45S5 group and control group, respectively.

ELISA quantified the protein secretion of Col I and OCN, and determined whether the changes in Col I and OCN protein levels were in parallel with gene activation. In agreement with the observed increase in RNA levels, nano-58S BG enhanced Col I and OCN protein secretion compared to 45S5 BG and the control group on day 7 (P < 0.05) (Fig. 2(c) and (d)).

The mineralization assay indicated that nano-58S significantly promoted mineral nodule formation (Fig. 3(a)). At 28 days, no red staining was observed in the non-osteogenic control group and moderate red nodule staining was observed in the osteogenic control group. As for the BG groups, the density of mineralized nodules was significantly higher in the nano-58S group than in the other groups. The 45S5 group showed similar staining to the osteogenic control group.
results of the semi-quantification of Ca deposition were consistent with the mineralization staining (Fig. 3(b)). The OD_{562} of the nano-58S group was significantly higher than the osteogenic control group. No statistical difference between the OD_{562} values was found between the 45S5 group and the osteogenic control groups.

4.3 Effects of BG extractions on MAPK activation

The results of western blotting showed that BG extractions significantly enhanced the phosphorylation of ERK and p38 at 12 h, and 1 and 3 days (Fig. 4(a)). Quantified using ImageJ 1.38 software (Fig. 4(b)), the expression of p-ERK was significantly higher in the nano-58S group than in the 45S5 and control
The expression of p-p38 in the nano-58S group was significantly up-regulated at 12 hours, and 1 and 3 days compared with the control group. The expression of p-p38 in the 45S5 group was similar to the nano-58S group at 12 hours and then showed no significant up-regulation compared to the control group. No expression change of JNK or p-JNK was found throughout the observations.

When the ERK pathway was blocked, the normal expressions of ALP, Runx2, Col I and OCN were inhibited significantly. With the ERK pathway blocked, nano-58S still increased ALP gene expression significantly compared to 45S5 and the control group but it was significantly lower than the normal expression levels of the control group without U0126, also Col I gene expression in the 45S5 group was significantly lower than in the control and nano-58S groups. With U0126, the mRNA levels of Col I, Runx2 and OCN in the nano-58S group and ALP, Runx2 and OCN in the 45S5 group were similar to the ERK blocked control group, which indicated no significant gene activation (Fig. 5(a)). When the p38 pathway was blocked, the gene activation of ALP showed no significant difference between the control, nano-58S and 45S5 groups with or without SB203580. Col I expression was similar in all three groups with SB203580 and was significantly lower than in the control group without SB203580. The gene expressions of Runx2 and OCN in the control, nano-58S and 45S5 groups with SB203580 were similar to the control group without SB203580, indicating no significant gene activation (Fig. 5(b)).

5. Discussion

The MG-63 cell line has been used in several studies due to its ideal osteoblastic characteristics, including the expression of BMP, ALP, Runx2 and OCN. In this study we focussed on ALP, Col I, Runx2 and OCN to observe the osteogenic gene activation effects of BG. ALP is the marker of the mature osteoblastic phenotype and extracellular matrix maturation.
predominant collagen in bone and the structural framework of matrices for inorganic deposition.\textsuperscript{27} Runx2 is an essential transcript factor and master gene of osteoblast differentiation, and regulates the expression of downstream transcript factors and osteogenic markers.\textsuperscript{28} As a late osteoblast differentiation marker, OCN can bond Ca with bone matrix and induce mineralized nodule formation.\textsuperscript{29} The results of our study showed that nano-58S and 45S5 extractions enhanced these gene expressions. The positive Runx2 and OCN protein expressions, the increased protein secretion of Col I and OCN, as well as the matrix mineralization verified the genes’ activation. According to the ELISA results, the increased Col I and OCN protein expression suggested that nano-58S BG and 45S5 BG promoted the early stages of bone matrix formation and mineralization in the MG-63 cells.

Previous studies have found that higher concentrations of Si and Ca, a suitable CaO/SiO\textsubscript{2} ratio and low concentrations of P can enhance an osteoblasts’ proliferation and

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Fig. 5  (a) Real-time PCR results of osteogenic genes in MG-63 cells cultured in BG extractions and the control medium with and without U0126; (b) real-time PCR results of osteogenic genes in MG-63 cells cultured in BG extractions and the control medium with and without SB203580. The data are relative expression levels shown as the fold difference compared to the control group without U0126 or SB203580 at day 4 and are normalized to the expression level of the housekeeping gene (GAPDH). The data are the mean values ± standard deviation of three individual experiments. Real-time PCR experiments were run twice for each sample (ANOVA, *P < 0.05 versus the control group, #P < 0.05 versus the 45S5 group).
differentiation.\textsuperscript{10,11} In agreement with the previous studies, and compared to 45S5 extractions, the possible reasons leading to the better gene activation of nano-58S BG may be the higher specific surface, better capacity for releasing ions of the particulates, and lower concentration of P in the extraction. However, besides the ion releasing capacity, there are several other features, such as composition, particle size, surface topography, degradation rate, and so on, which determine the bioactivity of bioactive glasses. Although we focused on the ionic production features of BG in this study, we speculated that both the nano-particle and the chemistry of the material enhanced the bioactivity of the nano-58S bioactive glass. To discern the effect of each feature on the BG’s bioactivity, a series of studies are needed in the future.

We hypothesized that osteogenic gene activation by BG is transduced through the MAPK signalling cascades into the cell’s nucleus and the response genes. The results of our study showed that the ERK and p38 pathways were activated, but no change in the JNK pathway was seen throughout the 7 days of observation. These results suggest that the ERK and p38 pathways may be involved in BG-induced osteogenic gene activation.

To find out whether ERK and p38 play a role in signaling transduction and gene expression regulation, we blocked the two pathways. MG-63 cells treated with U0126 showed inhibition of the normal expression of ALP, Col I, Runx2 and OCN. The gene activation of Col I, Runx2 and OCN by BG was also blocked. Blocking the p38 pathway showed no effect on the normal expression of Runx2 and OCN but the genes’ activation by BG was blocked. The up-regulation of ALP by BG showed the same trend with or without SB203580. Inhibition of Col I normal expression was found in all three groups with SB203580. These results indicated that the ERK and p38 pathways play a role in BG-induced gene activation. Based on the results that blocking the ERK pathway inhibited the osteogenic genes’ normal expression and the BG’s gene activation, while blocking the p38 pathway only blocked the BG’s gene activation, it may be that the ERK pathway plays a more important role compared to the p38 pathway.

Several previous studies suggested participation of MAPKs in BG-induced gene activation based on the gene expression change of MAPK members. In agreement with previous studies, the current study focused on the involvement of MAPKs and their phosphorylation and verified their signaling transducing effect for the first time. Previous studies have found that the ERK pathway regulates the proliferation and differentiation of mesenchymal stem cells.\textsuperscript{22} Specifically, ERK pathways can activate the transcription of Runx2 during osteoblast differentiation.\textsuperscript{23,24} As the “master gene” for osteogenesis, Runx2 also regulates the expression of non-collagenous proteins such as OCN and BSP. According to the previous studies and the results of the current study, we propose that BG extraction activates MAPK pathways, specifically the ERK and p38 pathways, to transduce extracellular ionic stimulation into the cell nucleus, and activate the expression of Runx2. Then Runx2 regulates osteoblast differentiation, the expression of OCN and finally mineralization (Fig. 6).

6. Conclusion

The present study verified that the MAPK pathway is involved and plays an important part in BG’s osteogenic gene activation. The effects of the novel nano-sized 58S BG on osteogenic genes up-regulation and MAPK phosphorylation are better than the traditional 45S5 BG.

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