

NIH Public Access

Author Manuscript

Biomacromolecules. Author manuscript; available in PMC 2011 July 12.

Published in final edited form as:

Biomacromolecules. 2010 July 12; 11(7): 1856–1862. doi:10.1021/bm100374n.

Synergistically enhanced osteogenic differentiation of human mesenchymal stem cells by culture on nanostructured surfaces with induction media

Mi-Hyeon Youa,†, **Moon Kyu Kwak**b,†, **Deok-Ho Kim**^c , **Keesung Kim**b, **Andre Levchenko**^c , **Dae-Yong Kim**a,*, and **Kahp-Yang Suh**b,d,*

^aDepartment of Veterinary Pathology, Seoul National University, Seoul 151-742, Korea

bSchool of Mechanical and Aerospace Engineering, Seoul National University, Seoul 151-742, Korea

^cDepartment of Biomedical Engineering, Johns Hopkins University, Baltimore MD 21218, USA

^dWorld Class University (WCU) Program on Multiscale Mechanical Design, Seoul National University, Seoul 151-742, Korea

Abstract

We have examined the effects of surface nanotopography on *in vitro* osteogenesis of human mesenchymal stem cells (hMSCs). UV-assisted capillary force lithography was employed to fabricate a scalable (4 cm \times 5 cm), well-defined nanostructured substrate of a UV curable polyurethane polymer with dots (150-, 400-, 600-nm diameter) and lines (150-, 400-, 600-nm width). The influence of osteogenic differentiation of hMSCs was characterized at day 8 by alkaline phosphatase (ALP) assay, RT-PCR, and real time PCR analysis. We found that hMSCs cultured on the nanostructured surfaces in osteogenic induction media showed significantly higher ALP activity compared to unpatterned PUA surface (control group). In particular, the hMSCs on the 400-nm dot pattern showed the highest level of ALP activity. Further investigation with realtime quantitative RT-PCR analysis demonstrated significantly higher expression of core binding factor 1 (Cbfa1), osteopontin (OP), and osteocalcin (OC) levels in hMSCs cultured on the 400-nm dot pattern in osteogenic induction media. These findings suggest that surface nanotopography can enhance osteogenic differentiation synergistically with biochemical induction substance.

Introduction

Cell fate is dictated in part by adhesive, mechanical interactions between cells and surrounding extracellular matrix (ECM) substrates as well as the milieu of soluble, diffusible factors. ¹⁻² In particular, surface nanotopography has been shown to exert influence over adhesion, proliferation, and gene expression in many cell types. Recent advances in microand nanofabrication techniques have great potential to exercise a high degree of control over the physical properties of biomaterials such as elastic modulus, $3-4$ roughness, $5-6$ size, $7-8$ and topography $9-12$ at cell/tissue-implant interface, allowing for the study of mechanical interactions between cells and their local environment on the sub-cellular scale. $13-18$ In the past few years, for example, exploration of synthetic nanometer-scale features to mimic natural matrices has revealed a significant influence of surface nanotopography on cellular behaviors including changes in gene expression, cell proliferation, migration, adhesion, and

^{*}To whom correspondence should be addressed. sky4u@snu.ac.kr or daeyong@snu.ac.kr.

[†]These authors contributed equally to this work.

Engineered nanometer-scale scaffolds hold great promise for stem cell differentiation and transplantation. $6, 12, 26-27$ Arguably, to facilitate this research, more facile and efficient fabrication methods need to be developed to direct differentiation of stem cells into a particular lineage for clinical application. Recent studies have shown that mammalian cells including hMSCs are capable of responding to the substratum nano-topography. 22, 28-33 In particular, cues arisen from nanotopographically-defined surfaces apparently directed differentiation into neuronal lineage 30 or fibroblast of human bone marrow stromal cells. ³⁴-36 In addition, hMSCs cultured on the nanograting of 350-nm width lines showed significant neuronal and muscular gene expression. 30

The hMSCs have been isolated and established from several sources including bone marrow, adipose tissue, and umbilical cord blood. Of these, the bone marrow derived hMSCs, with the surface markers of CD105⁺, CD166⁺, CD29⁺, CD14⁻, CD34⁻, CD45⁻, have self limited renewal and differentiation capability into diverse cells of mesodermal origin, e.g., bone, cartilage, muscle, and connective tissues. $37-39$ It has been revealed that the differentiation of hMSCs into osteogenesis is influenced by Runx2 (Runt-related transcription factor 2) gene. ⁴⁰ The Runx₂ gene regulates bone development by osteogenesis using G protein-coupled signaling pathway, promoting up-regulation of bone specific extracellular matrix, such as alkaline phosphatase (ALP), osteopontin (OP), osteocalcin (OC), and bone sialoprotein (BSP). Runx2 expression and its activity are influenced by external signal, cell to cell interaction, and growth regulatory factor. $22, 41-42$

In the present study, we investigated the osteogenic differentiation of bone marrow-derived hMSCs by culturing these cells on a range of diverse nanostructured surfaces. To achieve this goal, well-defined, large-area $(3 \times 3 \text{ cm}^2)$ nanopatterns (dots and lines) were fabricated on glass coverslips using UV-assisted capillary force lithography (CFL), which allowed for a simple and scalable approach to creating a structured biomaterial interface for cell adhesion studies. $15, 20, 43$ This technique was developed by combining the crucial element of soft lithography – using an elastomeric or soft mold – and the use of capillarity for polymer molding. 44 This method allows one to exploit topographic definition of the substratum while eliminating the need to use an extremely high pressure that is typically needed in nanoimprint lithography. CFL has proven to be effective in fabricating various complex nanopatterns using a mold material with differences in permeability, mechanical modulus, and surface tensions.¹⁵

With UV-assisted CFL technique, various scalable $(>\frac{3\times3}{\text{cm}^2})$ nanopatterns were fabricated such as nanoscale dots (150-, 400-, 600-nm diameter) and lines (150-, 400-, 600-nm width). For the patterning material, a UV curable polyurethane polymer functionalized with acrylate groups (polyurethane acrylate, PUA) was used for its biocompatibility, fast curing time (< 20 s), and easy release from the mold. ⁴⁵⁻⁴⁶ Using these nanopatterns, the effects of surface nanotopography on *in vitro* osteogenesis of human mesenchymal stem cells (hMSCs) were examined. We found that osteogenic differentiation of hMSCs was much enhanced on the nanostructured surfaces of 150-nm width lines, 150-nm diameter dots, and 400-nm diameter dots in osteogenic induction media (OM), as verified by alkaline phosphatase (ALP) assay, RT-PCR analysis. Further investigation with real-time quantitative RT-PCR analysis showed that the significantly higher Cbfa1 level of hMSCs was expressed from culture on the nanostructured surface of 400-nm diameter dots $(p<0.05)$ compared with the unpatterned PUA surface (control group). These results suggest that surface nanotopography, in the

presence of suitable biochemical signals, can exert significant influence on regulating stem cell differentiation.

Experimental section

Fabrication of nanostructured substrates with UV-assisted CFL

A small amount (∼0.1 – 0.5 mL) of a UV curable PUA prepolymer was drop-dispensed on silicon master having positive patterns (features sticking out) and a supporting poly(ethylene terephthalate) (PET) film was carefully placed on top of the surface to make conformal contact. The PET film used in this study was surface modified with urethane groups to increase adhesion to the acrylate-containing monomer (Minuta Tech., Korea). The silicon masters had been prepared by photolithography or electron-beam lithography. To cure, the resin was exposed to UV (wavelength: 250 ∼ 400 nm) for 17 s at an intensity of 100 mW/ cm², and the cured 1st replica was peeled off from the master using a sharp tweezer. The 1st replica was additionally exposed to UV overnight to remove any uncured active groups on the surface. The 2nd replica was prepared by using a capillary molding process on glass coverslips with the over-cured 1st PUA as a mold, resulting in the same pattern as the silicon master. After curing, the 1st replica was removed from the surface using a sharp tweezers. The fabricated PUA nanopatterns were sterilized by rinsing with IPA (isopropyl alcohol) and D.I. water, and coated with 0.1 % gelatin for 1 hr at room temperature prior to cell culture. A schematic diagram of the fabrication procedure is illustrated in Figure 1 along with the self-replication characteristic of the PUA material.

Human mesenchymal stem cell culture

Cultured hMSCs were purchased from FCB-Pharmicell CO., Ltd., Sungnam, South Korea. Initially, hMSCs were obtained from 20 mL of aspirates from the iliac crest of normal human doners. hMSCs isolation and culture procedures were performed under GMP conditions as previously reported. $47-48$ hMSCs were maintained in growth medium (GM; DMEM, 10% FBS, 0.3 mg/ml glutamine, 100 units/ml penicillin, and 100 μg/ml streptomycin). Fifth passage hMSCs was used for experiments. For osteogenic differentiation, hMSCs were cultured on diverse nanopatterns (150-, 400-, 600-nm width lines and 150-, 400-, 600-nm diameter dots) in osteogenic differentiation medium (OM, 50 μM ascorbic acid-2-phosphate, 10mM β-glycerophosphate, and 100 nM dexamethasone) for 8 days at a cell density of 3000 cells/cm². As a negative control, hMSCs were cultured in GM for 8 days at the same density using the same patterns. To ensure reproducibility and biological significance of the data, each experiment was run three times under the identical conditions with subsequent analysis of osteogenic differentiation of hMSCs by using ALP activity assay as well as real time RT-PCR.

ALP activity assay

After 8 days culture, hMSCs $(3000 \text{ cells/cm}^2)$ were harvested, and total lysate was prepared using a RIPA lysis buffer supplemented with a protease inhibitor cocktail (Calbiochem, San Diego). Protein concentrations of the lysate were determined by a Bio-Rad Protein Assay Kit (Bio-Rad, Hercules, CA), and ALP activity was analyzed by incubating the lysates with p-nitrophenylphosphate (4.5 mg/ml) at 37 °C for 2 hours prior to the measurement of OD 410 by a spectrophotometer (Bio-Rad, Japan). By performing the assay using known concentration of ALP in parallel with the samples, the concentrations of ALP for the samples were calculated and normalized to metabolic activity.

Cell staining

ALP was stained using Sigma kit #85 according to the manufacturer's instructions. In brief, samples were fixed in acetone/citrate, rinsed in water, and stained with Fast Blue RR/ naphthol.

RT-PCR analysis

Total RNA was purified from the cells using TRIzol reagent, according to the manufacturer's protocol (Invitrogen). 1 μg of total RNA was used in cDNA synthesis with random hexamers as primers (Promega). PCR amplification was subsequently performed with the following primer sets: for Cbfa1, sense: 5'-AGAGCTGAACAGGAACAACGT-3' and antisense: 3′- CACCAGCAAGAAGAAGCCTTTG-5′; for glyceraldehyde-3-phosphate dehydrogenase (GAPDH), sense: 5′-ATTACAGTCCATGCCATCA-3′ and antisense: 5′- TCCACCACCCTGTTGCTGTA-3′; for alkaline phosphatase(ALP), sense: 5′- TTAGTGCCAGAGAAAGAG-3′ and antisense: 5′-CTTGGCTTTTCCTTCATGGTG-3′; for peroxisome proliferator activator receptor γ 2(PPARγ2), sense: 5′- GCTGTTATGGGTGAAACTCTG-3′ and antisense: 5′- ATAAGGTGGAGATGCAGGTTC-3′; for Osteopontin (OP), sense: 5′- CACATCGGAATGCTCATTGC-3′ and antisense: 5′-ATCACCTGTGCCATACCAGT-3; for Osteocalcin (OC) sense: 5′-ATGAGAGCCCTCACACTCCTC-3′ and antisense: 5′- GCCGTAGAAGCGCCGATAGGC-3; for Sox 9, sense: 5′- ACGTCATCTCCAACATCGAGACC-3′ and antisense: 5′- CTGTAGTGTGGGAGGTTGAAGGG-3 All PCR was carried out with an initial denaturation at 94 °C for 2 minutes followed by 35 cycles at 94 °C for 20 seconds, annealing temperatures of 57.4 °C (Cbfa1), 41.5 °C (ALP), 53.8 °C (GAPDH), 59.3 °C (PPAR γ 2), 56.0 °C (OP), 55.0 °C (OC), 55.0 °C (Sox 9) for 30 seconds, and a final extension at 72 °C for 5 minutes. All samples were run in triplicates in each experiment.

Real-time quantitative RT-PCR (sybr method)

Total RNA was extracted by using Trizol (Invitrogen). RNA was reverse-transcribed into cDNA. Two microliters of each RT reaction was amplified in a 20 μL PCR assay volume containing 10 μM each primer and power SYBR Green PCR Master Mix (applied Biosystems). Samples were incubated in the ExiCycler for an initial denaturation at 94 °C for 10 min, followed by 40 PCR cycles. Each cycle consisted of the following times and temperatures: 94°C for 40 s, 60°C for 30 s, and 72°C for 30 s. All gene expressions were normalized according to glyceraldehydes 3-phosphate dehydrogenase using the comparative cycle thresholod (ddCt) methods. The Cbfa1, sense: 5′-CACTGGCGCTGCAACAAGA-3′ and antisense: 5′- CATTCCGGAGCTCAGCAGAATAA-3′ (product size 127 bp); for OP, sense: 5′-ATTCTGGGAGGGCTTGGTTG-3′ and antisense: 5′-

TGTGGTCCCGACGATGCT-3′; for OC, sense: 5′-CTCCAGGCACCCTTCTTTCC-3′ and antisense: 5′- ATTCCTCTTCTGGAGTTTATTTGGG-3′ for glyceraldehyde-3-phosphate dehydrogenase (GAPDH), sense: 5′-GCACCGTCAAGGCTGAGAAC-3′ and antisense: 3′- ATGGTGGTGAAGACGCCAGT-5′ (product size 142 bp).

Data analysis

All data were presented as mean \pm SD. The results were analyzed using one-way analysis of variance (ANOVA) test to assess the statistically significant difference of overall differences in ALP activity level and real time PCR results. When one-way ANOVA test revealed $p <$ 0.05, the data were further analyzed by Dunnett's t-test to assess the statistical difference between treatment and control groups. Differences were considered statistically significant, only when p values were less than 0.05 ($p < 0.05$).

Results and discussion

Nanotopography and substrate characterization

ECM is composed of various protein fibers interwoven with glycosaminoglycan chains. *In vivo*, cells sense and respond to topographical features of the local microenvironment on multiple length scales such as bone roughness, molecular complexes controlling cell to cell interaction, and collagen banding. Aiming at mimicking the cellular microenvironment, *in vitro* studies have been centered on modulating the roughness by blasting. ⁴⁹⁻⁵¹ The previous findings, however, have often been conflicting, hard to reproduce, and performed on the micro- rather than the more relevant nano-meter scale. To overcome these limitations, we developed a well-defined, scalable cell adhesion substratum to direct differentiation of hMSCs into osteoblasts.

For cell culture substrates, various nanoscale dot and line patterns of PUA material were fabricated on optically transparent glass coverslips using UV-assisted CFL and selfreplication (Figure 1). Subsequently, the hMSCs were cultured on the nanopatterns, whose feature sizes were at least one or two order of magnitude smaller than that of the cell body. Figure 2 shows representative SEM images of such nanostructured substrates, demonstrating that the patterns exhibited high structural fidelity with well-defined vertical edge profiles. In order to generate uniform patterns, care should be taken during the 1st and 2nd replication steps so as to preserve the original geometry of the silicon master. This is because air cannot easily permeate out of the cured PUA replica and might be trapped in the spaces between nanostructures, resulting in a reduced height or a non-uniform height distribution. To alleviate this problem, a roller was mildly rounded several times on the surface, which helped expel trapped air prior to UV exposure. It is noted that the PUA material has been extensively used as a cell culturing substrate for many cell types with good biocompatibility. ¹⁵, 23, 52-54 In the current experiment, no considerable level of cell death was observed during a long culture of < 1 week, as evidenced by tests with a cell toxicity maker with control group (hMSCs cultured in normal 25 flask) (data not shown).

In the experiments, two types of nanopatterns, varying in size from 150-nm to 600-nm was used: dots (150-, 400-, 600-nm in diameter) and lines (150-, 400-, 600-nm in width). For consistency, the pattern height was fixed at a constant height of 500-nm, as verified by atomic force microscopy (AFM) measurements (not shown). As shown in Figure 2, the patterns were prepared over a large area (4 cm by 5 cm) with good physical integrity, allowing for large-area investigation of hMSCs differentiation into osteoblasts. It is noted in this regard that for cell biological applications, it would be beneficial to have large area patterns, not smaller than 1×1 cm², for various cell density and conventional biochemical assays (e.g. western blotting, RT-PCR, etc).

Expression of ALP Activity in hMSCs

To investigate the role of surface nanotopography, hMSCs were cultured for 8 days at the initial cell density of 3000 cells/cm² on the nanopatterns as well as on the flat PUA surface (control) in GM and OM, respectively. The initial plating density of hMSCs was fixed at 3000 cells/cm² because the lineage-specific differentiation of hMSCs is significantly affected by initial cell density. 55 As surface roughness is a defining feature of the bone tissue, we hypothesized that osteogenic differentiation of hMSCs might be enhanced on nano-topographically defined surfaces. To determine whether surface nanotopography could enhance hMSCs differentiation into osteoblasts, we characterized the ALP activity level of hMSCs cultured on various nanostructured substrates. ALP is widely used as a marker of the osteogenic phenotypes, because cell membrane of osteoblasts is composed of glycoprotein, an isoform of ALP. 56 The ALP activity of hMSCs was measured using cell samples

collected on day 8. As shown in Figure 3a-b, the ALP activity level of hMSCs was increased for the patterned substrates both in OM and GM. In particular, the hMSCs that were cultured on 150-nm line, 150-nm dot, and 400-nm dot patterns in OM showed significantly higher level of ALP activity compared to the other nanopatterns and the control group ($p < 0.01$). Among the patterns tested, the hMSCs on 400-nm dot patterns showed the highest level of ALP activity, with respect to which all the data were normalized both in OM and GM. The osteogenic differentiation is well represented by the characteristic spindle shapes of cells, as shown in Figure 3c. Also, a fluorescent image in Figure 3d demonstrates that the cells express a higher intensity of ALP when cultured on a nanopatterned substrate in OM (400 nm dot patterns used here).

As mentioned above, the hMSCs in OM showed higher up-regulation of osteogenic marker (Figure 3a). Interestingly, the hMSCs in GM also showed increasing ALP activity for osteogenesis as compared to control group (Figure 3b). The increased level, however, appears to be less significant than that for OM and the variations among different patterns were observed to be small. These findings suggest that mechanical topographic cues induce the osteogenic differentiation of hMSCs, the level of which can further be enhanced when combined with a suitable chemical definition of the cell medium. In addition, a specific topography (e.g, 400-nm dot patterns) could more synergistically affect osteogenic differentiation in the presence of OM. In fact, in clinical uses of dental implants, rough surfaces in bone-implant contact have already shown a better osseous fixation than smooth surface. $57-58$

Expressions of Cbfa1, ALP, OC, OP, PPARγ2, and Sox 9

As reported above, we observed that the ALP activity level of hMSCs was increased on all the nanopatterns tested as compared to the control group. To explore the differentiation outcomes further, we conducted semi-quantitative RT-PCR on day 8 to detect molecular markers of adipocyte, chondrogenic, and osteoblast lineage. For this purpose, hMSCs were cultured in OM and GM for 8 days at the same density of 3000 cells/ cm^2 on the same surfaces as described in the previous section.

As shown in Figure 4, adipocyte marker (peroxisome proliferator activator receptor γ 2, PPARγ2), osteoblast markers (ALP, Cbfa1: early osteogenic marker, OC, OP: late osteogenic marker), and chondrogenic marker (Sox9) displayed different expressions. In OM, ALP level was expressed for various nanopatterns except for 600-nm lines and dots. ALP level in GM was also expressed for various nanopatterns except for 600-nm lines and dots. The reason for the absence of ALP activity in the case 600-nm patterns is not clear at this stage, which appears to be associated with different cell attachment and penetration into the spaces of nanopatterns. ⁵⁹

As for other osteogenic markers, Cbfa1 was expressed for various nanopatterns both in GM and OM. In OM, the OC level was more intensively expressed for various nanopatterns than that in GM. The OP level also showed higher expression with nanopatterns and especially the highest intensity for 400-nm dots in OM. In contrast, the intensities of OC and OP were relatively lower for the nanopatterns in GM. The fact that the Cbfa1 is similarly expressed both in GM and OM indicates that early differentiation occurs in both media. Moreover, the cell density was properly controlled in the experiment, which can further facilitate osteogenic differentiation even in GM sample. However, late stage differentiation, as evidenced from staining results from OP and OC, was only observed in OM, supporting the existence of synergistic role of nanotopography and osteogenic induction media.

Figure 4 also demonstrates differentiation into other lineage using molecular makers of $PPAR\gamma2$ and Sox 9, in which the markers were not detected for all the nanopatterns in OM

and GM (except for expressing PPARγ2 marker in GM control). On the basis of these results, osteogenesis appears to predominate over adipogenesis and chondrogenesis for 8 days in the presence of the induction media substance. It is noted that the cell proliferation was low, so that the cell density was almost maintained at ~3000 cells/cm² for 2∼3 weeks, in agreement with the previous study. ³¹

Real time PCR

We also conducted real time quantitative RT-PCR using hMSCs on the 400-nm dot pattern and control on day 8. As shown in Figure 5a-c, the expression levels of Cbfa1, OC, and OP on the 400-nm dot pattern (GM and OM) increased significantly in comparison to the control group ($p < 0.05$), further confirming strong influence of the surface nanotopography on osteogenic differentiation. Here, ALP and Cbfa1 are expected to increase early in osteogenesis and decrease with mineralization. Also, OC and OP are the main components of mineralized ECM for osteoblasts. As can be seen from the figures, ALP and Cbfa1 were expressed both in GM and OM, with significantly lower levels of OC and OP in GM samples. This suggests that early osteogenic differentiation can be initiated regardless of the choice of media with the help of a proper cell density; the late stage osteogenesis (mineralization), however, is hardly observed in GM-control, supporting the synergistic role of nanotopography and differentiation induction media. As with other staining and PCR results, the 400-nm dot pattern showed the higher mineralization with some local distribution of the osteogenic differentiation.

Conclusions

In this study, we have presented that nanotopographic surface definition could synergistically induce a significant up-regulation of osteogenic markers such as ALP, Cbfa1, OC, and OP with osteogenic induction media (OM), demonstrating directed differentiation into osteogenic lineage of hMSCs. We have used UV-assisted capillary lithographic technique with a polymeric biomaterial of PUA for simple and scalable $(4 \text{ cm} \times 5 \text{ cm})$ fabrication of surface nanotopography. Based on alkaline phosphatase (ALP) assay, RT-PCR, and real-time quantitative RT-PCR analysis, we found that the hMSCs responded to the underlying surface nanotopography by up-regulation of mRNA osteogenic level, and the 400-nm dot pattern appeared more effective in facilitating osteogenic differentiation synergistically with osteogenesis induction media (OM). In normal culture media (GM), the hMSCs also showed increased level of ALP, OP, and OC activity for osteogenesis but the increased level was less significant than that for OM and the variations among different patterns were observed to be small.

In conjunction with our findings, a previous study by Dalby et al. 22 reported that the differentiation of hMSCs could be enhanced when cultured on dot patterns. They found that when hMSCs were cultured on dot patterns and displaced randomly by up to 50-nm on both axes from their initial position, the cells showed more osteogenic differentiation. This observation, together with our findings, suggests that the hMSCs can sense the underlying surface nanotopography and differentiate into different cell lineage. Furthermore the dot patterns might be more effective in up-regulating osteogenic differentiation presumably due to its biomimetic character with respect to the actual bone surface, stimulating Runx 2 gene of hMSCs. A further study would be required to elucidate the biomechanical or physical role of nanopatterns on the differentiation of hMSCs. Our study further demonstrated that tunable nanotopographic features may have synergistic effect with the chemical medium composition on directed stem cell differentiation into osteogenic lineage.

Acknowledgments

This research was supported by the WCU (World Class University) program (R31-2008-000-10083-0), the Institute of Advanced Machinery and Design, and the Korea Research Foundation Grant funded by the Korean Government (MOEHRD) (Grant KRF-J03003). This work was also supported in part by National Institute of Health (R21- EB008562).

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Figure 1.

A schematic diagram for the fabrication of PUA nanopatterns by self-replication and capillary force lithography (CFL). As shown, the PUA patterns (2nd replica) have the same geometry with the silicon master after double replications.

Figure 2.

Planar SEM images of the fabricated nanopatterns by UV-assisted CFL. Scale bars represent 5 μm and 1 μm in the panel and inset images, respectively. (a) 150-nm diameter and 500-nm pitch dots. (b) 400-nm diameter and 800-nm pitch dots. (c) 600-nm diameter and 1200-nm pitch dots. (d) 150-nm width and 200-nm pitch lines. (e) 400-nm width and 800-nm pitch lines. (f) 600-nm width and 1200-nm pitch lines.

Figure 3.

ALP expression level of the hMSCs plated at 3000 cells/cm², cultured for 8 days in (a) OM and (b) GM. The data were normalized with respect to the highest level of ALP activity for 400-nm dot pattern both in OM and GM. As shown, the ALP activity was increased for the patterned substrates both in OM and GM compared to the control group. Especially, the hMSCs cultured on 150-nm line, 150-nm dot, and 400-nm dot patterns in OM showed significantly high ALP activity level ($p < 0.01$). *: $p < 0.01$. (c) Bright-field images of hMSCs cultured on 400-nm dot pattern in OM, plated at 3000 cells/cm², cultured for 8 days. Bar = 200 μm. Inset shows a magnified image of osteogenic hMSCs exhibiting a spindle shape. Bar = 20 μm. (d) Fluorescent image of hMSCs cultured on 400-nm dot pattern in OM stained for ALP (osteogenic marker). Bar = 100μ m.

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Figure 4.

RT-PCR analysis: (a) ALP, (b) Cbfa1, (c) OC, (d) OP, (e) PPAR γ 2, (f) Sox9, and (g) glyceraldehyde-3-phosphate dehydrogenase (GAPDH). All samples were collected after 8 days in culture.

Figure 5.

Quantification of osteogenic-specific marker expression by real-time PCR at day 8 (* and * p < 0.05): (a) Cbfa1, (b) OP, (c) OC.