

COLLAGENATED HETEROLOGOUS CORTICO-CANCELLEUS BONE MIX STIMULATED DENTAL PULP DERIVED STEM CELLS

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Collagenated heretologous cortico-cancellous bone mix (CHCCBM) is largely employed in maxillary and dental surgery for regeneration procedures, and is similar to human bone from chemical and physical point of view and promotes osteogenesis. In order to get more inside how this biomaterial induces osteoblast gene expression to promote bone formation, the mRNA levels of bone related genes were compared in human osteoblasts and dental pulp stem cells, using real time RT-PCR. The obtained results demonstrated that CHCCBM enhance stem cells differentiation and deposition of matrix by the activation of osteoblast related genes SP7, FOSL1 and SPP1.

Collagenated heretologous cortico-cancellous bone mix (CHCCBM) is a bovine bone derived employed in maxillary and oral surgery. Some reports have demonstrated its clinical efficacy to restore alveolar ridge in pre-prosthetic surgery (1-3).

To investigate the molecular mechanism by which CHCCBM promotes osteoblast differentiation and proliferation, the quantitative expression of the mRNA of specific bone related genes, was examined in derived dental pulp stem cells treated CHCCBM. Dental pulp stem cells (DPSCs) represents an ideal source of stem cells because approachable niches containing a high number of stem cells compared to equal volumes of bone marrow (4-6). In this study the expression levels of specific genes were examined by means of real time RT-PCR in DPSCs after treatment with CHCCBM.

Gene expression in DPSCs was then compared with the gene expression in treated Human

Osteoblasts (HOb) to evaluate the potential effect of this material in osteoblasts differentiation.

MATERIALS AND METHODS

Stem cells isolation from dental pulp, flow cytometric analyses and primary human osteoblasts cell culture were performed as previously described (4-6).

Cell treatment

DPSCs and HOb were seeded with CHCCBM (Gen-Os, Tecnooss srl, Giaveno, Torino, Italy) at the concentration of 10 mg/ml. Another set of wells containing untreated cells were used as control. The cells were maintained in a humidified atmosphere of 5% CO₂ at 37°C. Cells were harvested at two time points, 15 and 30 days, for RNA extraction. RNA processing, RT-PCR and statistical analyses were conducted as previously described (4-6).

Key words: bone, dental pulp, stem cells, gene expression, osteoblast differentiation

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RESULTS

Cell cultures were phenotypically characterized by flow cytometric analyses as previously reported (4-6). To study if CHCCBM stimulates osteoblasts differentiation and proliferation in DPSCs, several osteoblast genes and mesenchymal stem cells marker, were analyzed by quantitative real-time PCR after 14 days of treatment with CHCCBM. Treated DPSCs resulted in up-regulation of SP7, FOSL1 and SPP1 genes and down-regulation of ENG, RUNX2, COL3A1, COL1A1 and ALPL during the first 14 days of treatment.

Analyzing the results obtained in HOb, we observed that after 14 days of treatment SP7, ENG, FOSL1, COL1A1, COL3A1 and SPP1 were up-regulated. The expression of RUNX2 and ALPL was down-regulated. "Two tailed ANOVA" comparison between DPSCs and HOb after the treatment showed that all genes were significantly differentially expressed after 14 days of treatment. (Table I).

DISCUSSION

CHCCBM (Gen-Os, Tecnos srl, Giaveno, Torino, Italy) is widely used in several bone regeneration procedures in oral surgery due to its characteristics of resorption rate and capability

of promotes osteogenesis. It is of paramount importance since several situation determine bone loss. Among them are socket preservation after tooth extraction, peri-implants bone maintenance (7-16), ridge reconstruction in syndromic patients (17-23) and oral rehabilitation after oncological treatments (24-35)

In order to get more inside how CHCCBM acts on DPSCs, changes in expression of bone related marker genes (RUNX2, SPP1, SP7, COL1A1, COL3A1, ALPL and FOSL1) and mesenchymal stem cells marker (ENG) were investigated by real-time RT-PCR. DPSCs were isolated by enzymatic digestion and phenotypically characterized by flow cytometric analyses. Dental pulp derived cell were homogenously CD105⁺, CD90⁺, CD34⁻, CD45⁻, CD14⁻, which is a typical mesenchymal stem cells surface antigen profile.

The osteoinductive properties of CHCCBM were demonstrated by the up-regulation of SP7 during all the entire treatment both in treated DPSCs than in treated HOb. SP7 is a transcriptional factor involved in bone formation and osteoblast differentiation downstream of RUNX2 pathway. CHCCBM also modulate the expression of collagenic extracellular matrix genes, collagen type 1 α 1 (COL1A1) and collagen type 3 α 1 (COL3A1).

After 14 days, in treated DPSCs was observed the

Table I. Differentially expressed genes between DPSCs and HOb after 14 days of treatment.

Genes	DPSCs	HOb	Differentially expressed genes
	Log10 RQ	Log10 RQ	p<0,005
SP7	1,22	0,52	0,001
ENG	-0,42	0,03	0,009
FOSL1	0,25	0,26	0,001
RUNX2	-0,42	-0,03	0,002
COL3A1	-1,46	0,02	0,001
COL1A1	-0,71	0,13	0,001
ALPL	-0,54	-0,23	0,001
SPP1	0,93	1,64	0,001

down-regulation of COL1A1 and COL3A1. Instead, in HOOb, the two collagens type were up-regulated. Osteopontine (SPP1) was up-regulated during all the treatment both in DPSCs than in HOOb. The up-regulation of this gene suggests the differentiation effect of treatment.

ENG, also named CD105, a surface marker used to define a bone marrow stromal cell population capable of multilineage differentiation was down-regulated in DPSCs and up-regulated in HOOb during the entire treatment. Another gene involved in osteoblast differentiation and modulated by CHCCBM was FOSL1. Both in DPSCs than in HOOb this gene was up-regulated. FOSL1 is a component of the dimeric transcription factor AP-1 involved in the transcription of bone related genes. This study demonstrates that CHCCBM is capable of supporting osteoblast differentiation and extracellular matrix deposition and mineralization in mesenchymal stem cells by the activation of osteoblast related genes SP7, SPP1 and FOSL1.

Within the limits of this *in vitro* study related to the period of observation (2 weeks), it possible to infer that CHCCBM is a bone substitute that positively active gene related to bone formation.

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