

Exposure of periodontal ligament progenitor cells to lipopolysaccharide from *Escherichia coli* changes osteoblast differentiation pattern

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ABSTRACT

Periodontal ligament mesenchymal stem cells (PDLMSCs) are an important alternative source of adult stem cells and may be applied for periodontal tissue regeneration, neuroregenerative medicine, and heart valve tissue engineering. However, little is known about the impact of bacterial toxins on the biological properties of PDLMSCs, including self-renewal, differentiation, and synthesis of extracellular matrix. Objective: This study investigated whether proliferation, expression of pro-inflammatory cytokines, and osteogenic differentiation of CD105-enriched PDL progenitor cell populations (PDL-CD105⁺ cells) would be affected by exposure to bacterial lipopolysaccharide from *Escherichia coli* (EcLPS). Material and Methods: Toll-like receptor 4 (TLR4) expression was assessed in PDL-CD105⁺ cells by the immunostaining technique and confirmed using Western blotting assay. Afterwards, these cells were exposed to EcLPS, and the following assays were carried out: (i) cell viability using MTS; (ii) expression of the interleukin-1 beta (IL-1 β), interleukin-6 (IL-6), interleukin-8 (IL-8), and tumor necrosis factor alpha (TNF- α) genes; (iii) osteoblast differentiation assessed by mineralization *in vitro*, and by mRNA levels of run-related transcription factor-2 (RUNX2), alkaline phosphatase (ALP) and osteocalcin (OCN) determined by quantitative PCR. Results: PDL-CD105⁺ cells were identified as positive for TLR4. EcLPS did not affect cell viability, but induced a significant increase of transcripts for IL-6 and IL-8. Under osteogenic condition, PDL-CD105⁺ cells exposed to EcLPS presented an increase of mineralized matrix deposition and higher RUNX2 and ALP mRNA levels when compared to the control group. Conclusions: These results provide evidence that CD105-enriched PDL progenitor cells are able to adapt to continuous *Escherichia coli* endotoxin challenge, leading to an upregulation of osteogenic activities.

Keywords: Cell differentiation. Bacterial antigens. Periodontal ligament. Stem cells. Osteogenesis.

INTRODUCTION

The great differentiation capacity of mesenchymal stem cells (MSCs), the possibility of grafting, and its expansion in culture led to an increased interest in the clinical use of these cells through directed administration on specific sites in many pathological situations. In this sense, regenerative therapy based on stem cells has gained attention since the

isolation of mesenchymal stem cells from various tissues.

The search for similar cells to mesenchymal stem cells in specific tissues has led to the discovery of a variety of stem cells in almost all organs and tissues of the body in recent decades²⁰. Dental tissue is a potent reservoir of postnatal stem/progenitor cells isolated from dental pulp⁴, periodontal ligament from permanent²¹ and deciduous teeth²², apical papilla²⁴, or dental follicles¹⁷. These postnatal

populations cells have been characterized by the expression of specific surface markers to mesenchymal stem cells (CD105, CD166, CD73 e STRO1)^{21,22}, and have stem cells-like properties, including the capacity for self-renewal and multilineage differentiation potential⁶. Periodontal ligament mesenchymal stem cells (PDLSCs) are an important alternative source of adult stem cells and may be applied in periodontal tissue regeneration^{21,25}, neuroregenerative medicine³, and heart valve tissue engineering¹⁴. Although PDLSCs can be routinely isolated and cultivated from periodontal ligament, many issues have to be approached before these cells can be used for therapies, such as their growth and differentiation in the presence of bacterial toxins, which may be present in many clinical situations.

Lipopolysaccharide (LPS) is an endotoxin present in Gram-negative bacteria such as *Escherichia coli*. This endotoxin, when in contact with cells of the host immune system, as well as gingival fibroblasts, induces an immune-inflammatory response characterized by the release of pro-inflammatory cytokines, which are mediated by Toll-like receptors (TLR2 e TLR4)⁹. The feasibility of using PDLSCs in regenerative procedures by virtue of their capacity to differentiate into multiple cell lines depends on whether the growth and osteogenic differentiation of PDLSCs are affected by bacterial stimuli. Some studies suggest that periodontal ligament cells may alter their phenotype in response to inflammation promoted by bacterial lipopolysaccharide^{23,27}. However, little is known about the interactions between bacteria and MSCs. Recently, it was observed¹⁶ that an *in vitro* change in osteogenic differentiation pattern occurs when mesenchymal stem cells isolated from the bone marrow of healthy individuals were exposed to the *Escherichia coli* LPS.

Thus, this study investigated the effects of lipopolysaccharides from *Escherichia coli* (EcLPS) on cell viability, expression of pro-inflammatory cytokines, and osteogenic differentiation potential of CD105-enriched PDL progenitor cell populations.

MATERIAL AND METHODS

Cell culture

This study was approved by the Institutional Review Board (#022/2011). Five populations of mesenchymal progenitor cells (STRO1⁺ CD105⁺ CD166⁺ CD34⁻ CD45⁻ cells) from periodontal dental ligament (PDL) of permanent teeth were obtained and characterized in a previous study²². Briefly, teeth extractions of completely erupted third molars of five subjects (aged 19 to 22 years) were performed at the Dental Clinic of the School of Dentistry, State University of Campinas, Piracicaba, São Paulo, Brazil. Teeth were placed in biopsy media composed

of Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum, 250 mg/ml gentamicin sulfate, 5 mg/ml amphotericin B, and 1% penicillin/streptomycin (Gibco BRL, Life Technologies, Carlsbad, CA, USA), and transferred to the laboratory facilities. The PDL tissues from permanent teeth were gently scraped from the mid-portion of the root surface and enzymatically digested in a solution of 3 mg/ml collagenase type II (Gibco BRL) and 4 mg/ml dispase (Gibco BRL) for 1 hour at 37°C. Heterogeneous single-cell suspensions were obtained by passing the cells through a 70-mm cell strainer (Falcon, BD Labware, Franklin Lakes, NJ, USA). Samples were expanded in 25 cm² culture flasks (Falcon) at 37°C, 5% CO₂ in DMEM supplemented with 10% fetal bovine serum, 1% L glutamine and 1% penicillin/streptomycin (Gibco BRL) (standard media). CD105⁺ enriched cell subsets from PDL (PDL-CD105⁺ cells) were isolated by magnetic cell sorting, and characterized by flow cytometry, and immunostaining. Flow cytometric analysis demonstrated a higher proportion of CD105⁺ cells co-expressing CD166 surface antigen, whereas immunostaining demonstrated expression of STRO-1. These cell populations showed multipotential capabilities to differentiate *in vitro* into an osteoblast phenotype and adipocyte-like cells²². All five PDL-CD105⁺ cell populations were cultured in Dulbecco's modified Eagle's minimal essential medium (DMEM) supplemented with 10% fetal bovine serum, 1% L glutamine and 2% penicillin/streptomycin (standard media) (Gibco BRL), frozen with Recovery™ Cell Culture Freezing Medium (Gibco BRL) and kept in liquid nitrogen for subsequent experiments.

Preparation of LPS solution from *Escherichia coli*

Bacterial LPS were suspended in 1 mL of sterile and endotoxin-free water as a stock solution (concentration 1 mg/mL) of *Escherichia coli* (Sigma-Aldrich, St Louis, MO, USA), and stored at -20°C. At the time of each experiment, the concentrations of 100 ng/mL, 1 µg/mL e 10 µg/mL¹⁰ were obtained by diluting each LPS in DMEM from the stock solution.

Immunostaining for TLR4

To evaluate the expression pattern of Toll-like receptor 4 (TLR4), PDL-CD105⁺ cell populations were seeded at 4×10⁴ cells/well on glass coverslips (13 mm, Knittel® GmbH, Braunschweig, Germany), placed in 24-well plates (Falcon), and cultured for 24 hours in standard media. Afterwards, cells were fixed in 4% paraformaldehyde for 10 minutes, blocked with 3% bovine serum albumin (BSA, Sigma-Aldrich) for 30 minutes, and incubated with a mouse antihuman TLR4 antibody (1:50; Abcam, Cambridge, MA, UK) for 1 h. Subsequently, cells

were incubated with a secondary antibody Goat anti-mouse IgG Alexa fluor 488 (1:1000) (Invitrogen, Carlsbad, CA, USA) for 1 hour. In order to identify nuclei, the samples were counterstained with TO-PRO[®]-3 iodide 642/661 (1:2000) (Invitrogen) for 15 minutes. As a negative control, cells were incubated only with the secondary antibody. Samples were then analyzed by confocal laser scanning microscopy (Leica TCS SP5A OBS, Mannheim, Germany).

Western Blot analysis for TLR4

Total cellular protein was extracted from cultured PDL-CD105⁺ cell populations using the RIPA buffer (Thermo Scientific, Rockford, IL, USA) supplemented with protease inhibitor cocktail (Sigma, P2714), following the manufacturer's specifications. Protein concentration was measured by the Bradford method (Bio-Rad Laboratories, Benicia, CA, USA) with bovine serum albumin (BSA) as a standard, and was measured spectrophotometrically at 595 nm. Equal amounts of protein per sample were separated by sodium dodecyl sulphate polyacrylamide gel electrophoresis (SDS-PAGE) 10% and transferred to a nitrocellulose membrane (Amersham[™] Hybond ECL, Amersham BioSciences[™], GE Healthcare, Little Chalfont, Buckinghamshire, UK). The membranes were blocked with 3% BSA in TBS for 1h at room temperature, incubated overnight with a 1:1000 dilution of primary antibody mouse anti-human TLR4 (Abcam), and then incubated with peroxidase-conjugated secondary antibodies (Anti-mouse 1:2500) (Santa Cruz Biotechnology, Dallas, TX, USA) for 2 h. The membrane was developed using an ECL reagent (SuperSignal West Femto Substrate, Thermo Scientific) and the signals were detected using radiographic films (Kodak, Rochester, NY, USA).

Cell viability assay

The MTS assay was used to determine cell viability of PDL-CD105⁺ cells. Single-cell suspensions (1.5×10^4 cells/well) were seeded on 96-well plates

with standard media for 24 h for cell attachment and spreading. Subsequently, media was changed to standard medium containing 0, 100 ng/mL, 1 µg/mL or 10 µg/mL concentrations of EclPS, and cells were cultured for 10 days. At time points of 1, 3, 7, and 10 days, MTS reagent (Promega Co., Madison, WI, USA) was added, and cells were incubated for 2 h at 37°C in a humidified CO₂ incubator. The absorbance of the formed formazan dye was measured photometrically at 490 nm.

Cytokines gene expression

Cells were seeded on a 60-mm dish (2×10^5 cells/dish) with standard media for 24 h. Subsequently, media was changed to standard media supplemented with 0, 100 ng/mL, 1 µg/mL or 10 µg/mL of EclPS. After 24 hours, total RNA was obtained using TRIzol[®] reagent (Invitrogen) to determine the expression of transcripts corresponding to interleukin-1 beta (IL-1β), interleukin 6 (IL-6), interleukin 8 (IL-8), and tumor necrosis factor alpha (TNF-α).

Osteogenic induction

PDL-CD105⁺ cells were seeded 2×10^5 in 60-mm tissue culture dishes in standard media for gene expression analysis and in 24-well plates to assess the mineral nodule formation *in vitro*. After 24 hours, standard medium was changed to an osteogenic-inducing medium (DMEM 10% FBS, 50 µg/ml ascorbic acid, 10 mM β-glycerol-phosphate, 10^{-8} M dexamethasone). To evaluate the effect of LPS on osteogenic differentiation, the differentiation medium was also supplemented with 1 µg/mL *E. coli* LPS. Total RNA was obtained using a reagent (TRIzol[®], Invitrogen) at days 3, 7, and 14 for the determination of the expression of transcripts corresponding to ALP, OCN, and RUNX2. In parallel, *in vitro* mineral nodule formation was assessed at day 21 using the alizarin Red assay (Sigma-Aldrich). For that, after 21 days of differentiation, cells were fixed with 70% ethanol and stained with alizarin Red (Sigma-Aldrich) for calcium detection. For

Gene	Foward Primer	Reverse Primer	Amplicon
ALP	5'-cgggcaccatgaaggaaa-3'	5'-ggccagaccaagatagaggt-3'	184
OCN	5'-agctcaatccggactgt-3'	5'-ggaagaggaaagaagggtgc-3'	150
RUNX2	5'-ccgtccatccactctaccac-3'	5'-atgaaatgcttgggaactgc-3'	139
IL-1β	5'-cttctcgacacatgggataac-3'	5'-tttgggatctactctccagc-3'	283
IL-6	5'-ctagagtacccagaacagattga-3'	5'-tcagcaggctggcattt-3'	154
IL-8	5'-gccaaagagaatatccgaactttaat-3'	5'-ctggctagcagactaggg-3'	172
TNF-α	5'-tccaccatgtactctctac-3'	5'-cctccagatagatgggctcata-3'	155
GAPDH	5'-acatcatccctgcctctac-3'	5'-ccaccttctgatgtcatcatattg-3'	171

ALP, alkaline phosphatase; OCN, osteocalcin; RUNX-2, runt-related transcription factor 2; IL-1β, interleukin-1 beta; IL-6, interleukin 6; IL-8, interleukin 8; TNF-α, tumor necrosis factor alpha; GAPDH, glyceraldehyde-3-phosphate dehydrogenase

Figure 1- Primers sequence used for amplification in real-time PCR

quantitative measurement, alizarin was solved in a 10% cetylpyridinium chloride monohydrate solution (PBS) for 15 min and samples were measured in a plate reader at 562 nm.

Gene expression analysis

Total RNA was DNase treated (Turbo DNase-free, Ambion, Foster City, CA, USA), and single-stranded complementary DNA (cDNA) was synthesized from 1 µg of total RNA using a kit (First Strand cDNA synthesis kit, Roche Diagnostics, Indianapolis, IN, USA) following the manufacturer's recommendations. Primers sequences for ALP, OCN, RUNX2, IL-1β, IL-6, IL-8, TNF-α and glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) are shown in Figure 1. qRT-PCR was performed with the LightCycler 2.0 and software (Roche Diagnostics GmbH, Mannheim, Germany), using the FastStart DNA Master plus SYBR Green Kit (Roche Diagnostics). Water (no template) was used as a negative control for all experiments. Relative quantification of reaction products was accomplished using *GAPDH* as the reference gene.

Statistical analysis

All experiments were performed in triplicate and repeated on at least three separate occasions. Mean and standard deviation were obtained to establish statistical comparisons. Data comparison was performed using one-way analysis of variance test ($\alpha=0.05$), and pairwise multiple-comparison test (Tukey test) was used to identify the difference among the groups. Additionally, non-parametric Kruskal-Wallis test ($\alpha=0.05$) followed by the Student-Newman-Keuls was performed to assess the effect of LPS on cytokines gene expression (BioEstat 5.0, Belém, PA, Brazil).

RESULTS

PDL-CD105⁺ cells express LPS receptor

Immunofluorescence analysis revealed that PDL-CD105⁺ cells exhibited positive staining for TLR4, which is a specific molecule involved in *E. coli* LPS signaling pathways (Figures 2A and 2B). This positivity for TLR4 was confirmed by the Western blotting analysis, showing that all five cell populations assessed expressed this receptor (Figure 2C).

E. coli LPS induces PDL-CD105⁺ response

To verify whether PDL-CD105⁺ cells responded to *E. coli* LPS, a metabolic activity as an indicator for cell viability was first measured using MTS assay. All PDL-CD105⁺ populations from five donors were able to retain the capacity to form adherent colonies and fibroblast spindle-shape after exposure to *E. coli* LPS (data not presented). As shown in Figure 3A, LPS did not affect the proliferation of PDL-CD105⁺ cells during the 10-day period, even after incubation with a relatively high concentration of endotoxin (1 µg and 10 µg/mL). In addition, MTS values showed a time-dependent increase of cell proliferation in all groups, with a higher number of cells after 3 and 7 days in the culture (Figure 3A).

Next, a qRT-PCR was performed to assess gene expression changes on pro-inflammatory cytokines IL-1β, TNF-α, IL-6 and IL-8 in PDL-CD105⁺ cells. After 24 h of toxin challenge, EclPS at concentrations of 100 ng and 1 µg/mL induced a significant increase of the transcripts for IL-6 and IL-8 compared to the control group (Figure 3B). Despite the fact that PDL-CD105⁺ cells only expressed IL-1β and TNF-α mRNA after LPS challenge, there were no

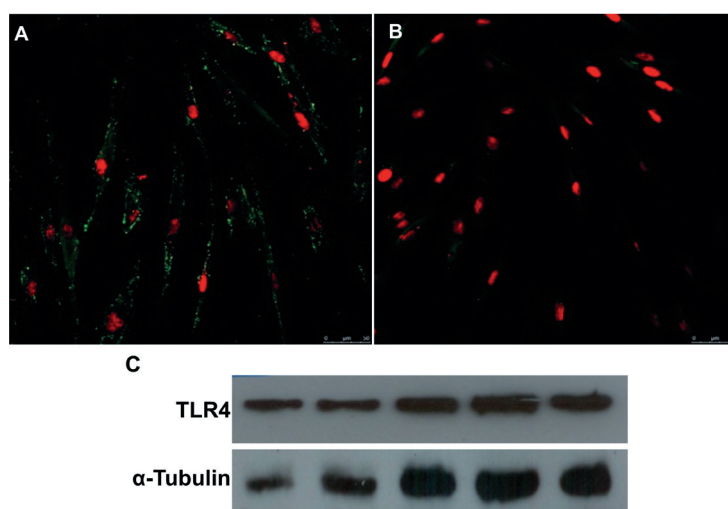


Figure 2- Highly purified CD105⁺ subsets of periodontal ligament (PDL) cells express Toll-like receptor 4 (TLR4). (A) Positive immunofluorescence for the TLR4 with nuclear localization confirmed by TO-PRO[®]-3 staining. (B) Control are represented by PDL-CD105⁺ cells incubated only with secondary antibody. (C) Western blotting analysis confirmed TLR4 in the five PDL-CD105⁺ cells

significant differences between cell exposed *versus* non-exposed, probably due to the greater variability of IL-1 β and TNF- α expression among the five populations (Figure 3B).

LPS challenge enhances osteogenic differentiation of PDL-CD105⁺ cells

Having demonstrated that EcLPS (100 ng and 1 μ g/mL) induced a pro-inflammatory cytokines response in PDL-CD105⁺ cells, next it was investigated whether EcLPS could change the osteogenic properties of these cells. For this evaluation, cells were only exposed to EcLPS at

concentration of 1 μ g/mL, and mineral nodule deposits (Alizarin Red staining) and gene expression of osteogenic markers (RUNX2, ALP, and OCN) were assessed.

After three weeks of osteogenic induction, mineral nodule formation could be observed in osteogenic differentiated cells of the positive control, as demonstrated by the quantification of Alizarin Red crystals (Figure 4A). This mineral deposition was increased when cells were exposed to osteogenic medium and *E. coli* LPS (Figure 4A). No mineralized nodules were observed in cells cultured in the DMEM used as a control (Figure

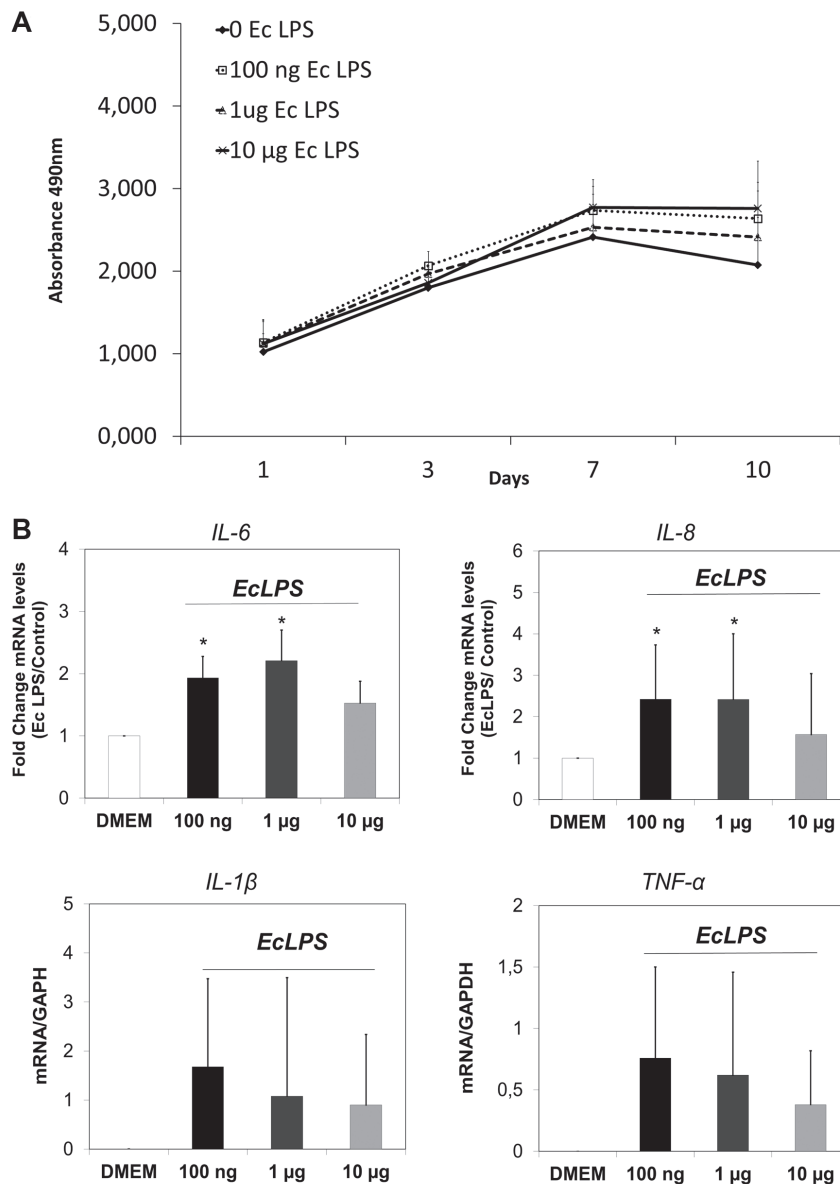


Figure 3- Effect of *E. coli* lipopolysaccharide (LPS) on the viability and pro-inflammatory cytokines expression. (A) PDL-CD105⁺ cells were cultured in the presence of EcLPS at concentrations of 0, 100 ng, 1 μ g and 10 μ g/ml. Metabolic activity as an indicator for cell viability was measured with MTS assay at the following time points: 1, 3, 7 and 10 days. (B) Real-time PCR showed an increase of mRNA levels of IL-6, IL-8, IL-1 β and TNF- α , respectively, in PDL-CD105⁺ cells after 24 hours of EcLPS exposure. Representative data of three independent experiments are shown. Statistical significance intergroup is indicated by * ($p < 0.05$)

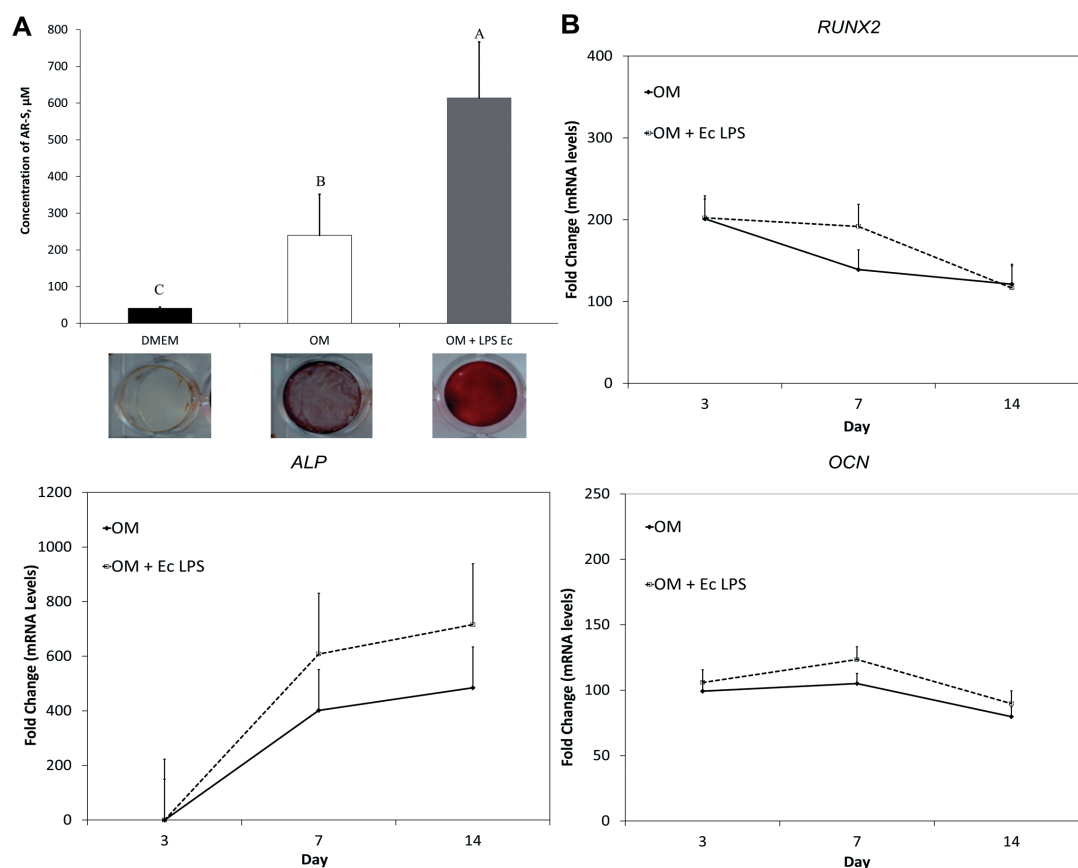


Figure 4- Osteoblast differentiation of PDL-CD105⁺ cells under EcLPS exposure. (A) Quantification of Alizarin Red staining showed that EcLPS at concentration of 1 $\mu\text{g}/\text{ml}$ increased mineral nodule formation at 21 d. (B) PDL-CD105⁺ cells were cultured in osteogenic medium (OM) alone or in OM+EcLPS (1 $\mu\text{g}/\text{ml}$). Data are presented as fold increase relative to non-induced controls for quantitative real-time PCR analysis showing that EcLPS did not alter the expression pattern of mRNAs for RUNX2, ALP, and OCN. All genes tested were normalized to the corresponding housekeeping gene, GAPDH. Representative data of three independent experiments are presented. Bars in (A) represent mean \pm standard deviation (SD) where intergroup statistical differences analysis are indicated by different capital letters ($p < 0.05$)

4A). qRT-PCR analysis further demonstrated an increased expression of a key osteoblast transcription factor, RUNX2, and two osteoblast-specific early/late differentiation transcripts (ALP and OCN, respectively) in osteogenically-induced cells compared to non-induced (Figure 4B). When cells were cultured in an osteogenic medium supplemented with *E. coli* LPS, higher levels of transcripts for RUNX2, ALP, and OCN were found on day 7 after osteogenic induction (Figure 4B), although there is no significant differences between cells induced and non-induced with bacterial toxin. However, taken together, these data suggest that LPS challenge enhances osteogenic differentiation of PDL-CD105⁺ cells.

DISCUSSION

This study investigated the effects of LPS from enterobacteria *Escherichia coli* on CD105-enriched PDL progenitor cell populations (PDL-CD105⁺ cells). The surface marker CD105 was chosen for

sorting progenitor cells from PDL heterogeneous population, since it is the most frequently reported positive surface marker in MSCs²². In addition, the expression of CD105 is one of the minimal criteria to define human MSCs according to the Mesenchymal and Tissue Stem Cell Committee of the International Society for Cellular Therapy². As part of cell characterization, the present study showed that TLR4 was expressed by CD105-enriched PDL progenitor cell populations. This finding is in accordance with previous studies which have also reported that TLR4 are expressed by mesenchymal stem cells such as bone marrow mesenchymal stem cells^{7,13} and periodontal ligament stem cells¹³. The family of the Toll-like receptors is involved in recognition of bacterial cell wall components¹⁵, and it has been described that *E. coli* LPS trigger an inflammatory environment by binding to TLR4^{1,7,13}.

To evaluate the influence of *E. coli*-derived LPS on cell proliferation, CD105-enriched PDL progenitor cell populations were initially cultured in the presence of different concentrations of this

bacterial toxin. The findings showed that cells remained viable and proliferative regardless of the LPS concentration. These data corroborate the findings of other studies, which assessed the effect of *Escherichia coli* endotoxins in pre-osteoblastic cells¹¹, bone marrow mesenchymal stem cells¹⁴, adipose-derived stromal cells⁸, and dental follicle stem cells¹⁸.

Since LPS from *E. coli* did not change cell growth, it was determined whether this endotoxin had any effect on cellular inflammatory response. It was found that ECLPS induced a significant expression of IL-6 and IL-8 in PDL-CD105⁺ cells. In addition, this bacterial toxin was able to induce the expression of two pro-inflammatory cytokines, IL-1 β and TNF- α . However, there were no significant differences between PDL-CD105⁺ cells exposed to ECLPS versus non-exposed, probably due to the greater variability of IL-1 β and TNF- α expression among the five cell populations. Similar to our data, previous studies had already observed an inflammatory response of oral cells exposed to LPS from *E. coli*. The expression of IL-6 and IL-8 were induced in periodontal ligament cells²⁶, while in dental follicle stem cells (DFCs) the gene for IL-1 β could also be observed after exposure to ECLPS¹⁸.

Some evidences of the literature supports the notion that inflammatory environment mediated by lipopolysaccharides can inhibit osteoblast differentiation of human periodontal ligament stem cells^{12,13}, mouse bone marrow mesenchymal stem cells⁷, and osteoblastic precursor cell line – MC3T3-E1⁵. It has been suggested that LPS decreases the osteogenic potential of mesenchymal stem cells through TLR4 regulated nuclear factor (NF)- κ B pathway^{7,13}. Further, in osteoblastic precursor cells (MC3T3-E1), LPS seems to induce cell apoptosis and inhibit osteoblast differentiation via c-Jun N-terminal kinases (JNK) pathway⁵.

Our findings demonstrated that the inflammatory response induced by *E. coli* LPS did not affect the osteoblast differentiation of CD105-enriched PDL progenitor cell populations. Unlike the studies cited above^{5,7,12,13}, ECLPS challenge increased mineralized nodules deposition through the osteogenic induction in PDL-CD105⁺ cells. Further, it is possible to suggest that ECLPS exposure induces an early osteoblast differentiation of PDL-CD105⁺ cells as demonstrated by increased expression of RUNX2 and ALP transcripts at day 7 of osteogenic induction. In agreement with our data, the prolonged exposure to ECLPS upregulated ALP activity and calcium deposition in human bone marrow mesenchymal stem cells¹⁶, and could also promote an increase in osteogenic marker genes expression such as ALP, osteopontin and bone morphogenetic protein 2 (BMP-2) in adipose-derived stromal cells⁸.

In the present study, prolonged endotoxin

exposure was designed to mimic a situation in which mesenchymal stem cells interact continuously with bacterial toxins during an inflammatory chronic disease, and then to evaluate whether this bacterial challenge could affect the osteogenic properties of periodontal ligament progenitor cells. Our findings suggest that CD105-enriched periodontal ligament progenitor cell populations acquire an adaptive tolerance under continuous LPS exposure. Previous studies reported endotoxin tolerance in oral mucosa cells associated with chronic periodontitis¹⁹. Also, transcripts for TLR2 and TLR4 were significantly downregulated in the gingival tissue of patients with chronic periodontitis, suggesting that the oral mucosa develops endotoxin tolerance in chronic periodontitis¹⁹. Interestingly, we also observed a downregulation of TLR4 in PDL-CD105⁺ cells under osteogenic induction associated to *E. coli* LPS exposure (data not shown).

However, further studies are required to elucidate the biological mechanism involved in the adaptive response by CD105-enriched periodontal ligament progenitor cells under continuous endotoxin exposure.

CONCLUSIONS

In conclusion, the results provide evidence that CD105-enriched PDL progenitor cells are able to adapt to continuous *Escherichia coli* endotoxin challenge, leading to an upregulation of osteogenic activities.

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