RESEARCH



Evaluation of the anti-inflammatory, antioxidant and regenerative effects of microbiota-derived postbiotics in human periodontal ligament mesenchymal stromal cells

Hazal Kibar Demirhan^{1,2} · Emine Omer Oglou^{1,2} · Zeynep Busra Aksoy³ · Fadime Kiran¹

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Abstract

Objective This study investigates the regenerative and protective effects of postbiotics (cell-free supernatant) derived from the *Lactiplantibacillus plantarum* EIR/IF-1 strain on human periodontal ligament mesenchymal stromal cells (hPDL-MSCs). **Materials and methods** hPDL-MSCs were isolated from periodontal ligament tissues (PDL) of wisdom teeth using enzymatic digestion and subsequently characterized through immunophenotyping. The effect of postbiotics on the viability of hPDL-MSCs was assessed using the MTT assay and flow cytometry, while their impact on cell migration was evaluated via the scratch assay. Anti-inflammatory effects of postbiotics were investigated on lipopolysaccharide (LPS, derived from *Porphyromonas gingivalis*)-stimulated hPDL-MSCs through Enzyme-Linked Immunosorbent Assay (ELISA). Additionally, the antioxidant effects of postbiotics were analyzed in hydrogen peroxide (H₂O₂)-induced hPDL-MSCs by measuring reactive oxygen species (ROS) levels using flow cytometry. The expression of collagen type I (*COL1A1*) gene was further assessed by quantitative reverse transcription PCR and immunofluorescence staining.

Results Treatment with postbiotics (250 μ g/mL) significantly increased the viability and migration capability of hPDL-MSCs, while enhancing collagen production for PDL repair. Treatment with postbiotics for 24 h resulted in a 54.53 \pm 2.01% reduction in intracellular ROS levels compared to untreated H₂O₂-induced hPDL-MSCs. Furthermore, postbiotics significantly decreased the production of pro-inflammatory cytokines (IL-8, IL-6, and IL-1 β), and increased the anti-inflammatory cytokine IL-10 (2.67-fold) compared to untreated LPS-stimulated hPDL-MSCs.

Conclusion Our findings indicate that postbiotics exhibit biological activity throughout all stages of the healing process, beginning with the modulation of the inflammatory response to LPS stimulation, followed by the promotion of cell migration, proliferation, and collagen synthesis. Given the unmet need for safe and adjuvant therapeutic approaches that promote comprehensive periodontal regeneration in periodontal diseases, this study presents postbiotics as a promising candidate. **Clinical relevance** Postbiotics could be integrated into regenerative therapies as a novel bioactive material to improve the healing and regenerative outcomes in periodontal defects by both controlling inflammation and stimulating tissue repair processes.

Keywords Periodontal disease · Postbiotics · Regeneration · Inflammation · Migration · Oxidative damage

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Introduction

Periodontal diseases, such as gingivitis and periodontitis, are one of the most widespread oral health conditions globally, with a prevalence reaching up to 50% worldwide [1]. Gingivitis represents the initial and mildest form of the disease. However, if neglected and untreated, it can progress, leading to further degenerative changes in the tissues and ultimately resulting in periodontitis [2]. As an inflammatory condition driven by biofilm-induced dysbiosis, periodontitis



Fadime Kiran fkiran@science.ankara.edu.tr

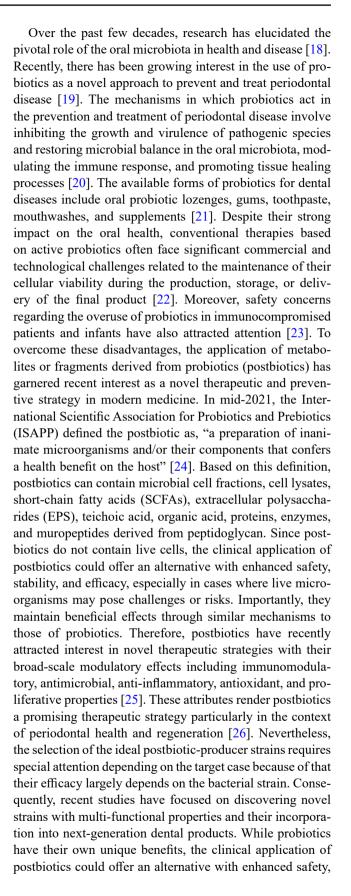
Pharmabiotic Technologies Research Laboratory, Department of Biology, Faculty of Science, Ankara University, Ankara 06100, Turkey

Graduate School of Natural and Applied Sciences, Ankara University, Ankara 06110, Turkey

Stem Cell Institute, Ankara University, Ankara 06520, Turkey

has emerged as a significant public health concern [3] affecting approximately 1.1 billion individuals worldwide, which represents 13.1% of the global population [4]. To date, numerous studies [1, 5, 6] have highlighted the consequences of periodontitis, including the destruction of soft tissues such as gingiva, periodontal ligament (PDL), and periodontal pocket, as well as the resorption of hard tissues (e.g., marginal alveolar bone loss), that may ultimately lead to tooth loss. Although current conventional periodontal therapies (improving oral hygiene, dental biofilm control, mechanical removal of plaque, local or systemic antibiotic therapy) focus on biofilm removal that mitigates the inflammation and aggressive progression of periodontitis, they cannot achieve complete periodontal regeneration which involves the formation of new cementum, alveolar bone, and functional PDL on the periodontitis root surface [7–9]. In light of the challenges associated with treatment of periodontitis, there is an urgent need for alternative therapeutic strategies to achieve complete periodontal regeneration [10].

For years, mesenchymal stromal cells (MSCs) have been investigated as a promising therapeutic strategy, not only for tissue regeneration [11], but also for immunotherapy [12] and chronic diseases [13]. Recently, dental MSCs, isolated from various sources within the oral cavity, have emerged as a promising approach for inducing functional dental tissue regeneration [10]. Among these, human periodontal ligament mesenchymal stromal cells (hPDL-MSCs) are particularly noteworthy for their role in maintaining periodontal tissue homeostasis and promoting regeneration, particularly in the context of periodontitis. Their high therapeutic potential is likely attributed to their immunomodulatory properties including the suppression of peripheral blood mononuclear cell and CD4+T cell proliferation, as well as the promotion of macrophage polarization towards an anti-inflammatory M2 phenotype. However, the inflammatory environment can modulate each of these mechanisms [14, 15]. Given their promising therapeutic potential, the number of clinical trials exploring the use of hPDL-MSCs, either in combination with scaffolds, membranes, or signaling molecules, or as standalone therapies, has been steadily increasing in the field of regenerative periodontology [16]. Despite the development of various therapeutic approaches utilizing hPDL-MSCs, their application in contemporary clinical practice for repairing the affected periodontium remains a significant challenge. This is primarily due to factors such as the limited availability of cells and the lack of multifunctional efficacy in treatment strategies that address different stages of periodontitis. To overcome these challenges, natural compounds have been proposed to enhance the survival, biological function, and therapeutic potential of hPDL-MSCs [17].





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stability, and efficacy, especially in cases where live microorganisms may pose challenges or risks.

This study aims to evaluate the regenerative, antioxidant, and anti-inflammatory effects of postbiotics derived from the Lactiplantibacillus plantarum EIR/IF-1 strain, previously isolated from the fecal microbiota of a breastfed infant, on hPDL-MSCs. This strain was selected for its demonstrated antimicrobial, anti-biofilm and anti-quorum sensing properties, particularly against pathogens implicated in dental diseases (Filifactor alocis, Fusobacterium nucleatum, Porphyromonas gingivalis, Prevotella denticola, Streptococcus mutans, and Streptococcus sanguinis) [27–29]. The observed effects of postbiotics on periodontitis-associated pathogens highlight their potential for integration into therapeutic strategies. To further strengthen our hypothesis, this study includes analyses investigating their role in tissue regeneration, as well as their capacity to mitigate oxidative damage and inflammation. Our findings not only advance our understanding of the effects of postbiotics on periodontal tissues but also introduce innovative approaches with the potential to revolutionize dental therapies and reshape the future of periodontal treatments.

Materials and methods

Bacterial strain and culture conditions

The Lactiplantibacillus plantarum EIR/IF-1 strain (NCBI GenBank accession number: OP810909.1), previously isolated and identified, was used in this study as a postbiotic producer. This strain was obtained from the Culture Collection of the Pharmabiotic Technologies Research Laboratory, Department of Biology, Ankara University (Ankara, Turkey), and preserved at -80 °C in de Man, Rogosa, and Sharpe (MRS; Merck, Germany) broth supplemented with glycerol (50:50%, v/v). Prior to use, the strain was cultured in MRS broth at 37 °C under static aerobic conditions for 24 h.

Preparation of postbiotics (cell-free supernatant)

The bacterial strain (100 µL) was cultured in 100 mL of MRS broth at 37 °C for 24 h. After incubation, the supernatant of bacterial culture was obtained by centrifugation at 15.000×g for 20 min, and sterile filtration was performed using a membrane filter with a pore size of 0.22 µm (Sartorius, Germany) to completely remove the cells. The filtered supernatant was subsequently lyophilized under freezing conditions of -20 °C, vacuum pressure of 0.120 mB, and a condenser temperature of -58 °C, using a freeze dryer (Buchi, Switzerland) for 48 h. The resulting lyophilized

powder was dissolved in distilled water (1 g/mL), sterilized through membrane filtration as previously described, and stored at -20 °C for subsequent assays [27]. For the purpose of standardization, identical conditions were maintained across all batches.

Isolation of human periodontal ligament mesenchymal stromal cells

Human periodontal ligament mesenchymal stromal cells (hPDL-MSCs) were isolated from periodontal ligament (PDL) tissues attached to one-third of the roots of wisdom teeth from four volunteers (two males and two females), aged 20-40 years, who were undergoing orthodontic treatment (Fig. 1a). All experimental protocols were approved by the Ethics Committee of Karamanoglu Mehmetbey University, Karaman, Turkey (Protocol code: 01-2023/20). Briefly, four healthy human teeth were collected in a container containing Dulbecco's phosphate buffered saline (DPBS) supplemented with 1% antibiotic antimycotic (Gibco, Thermo Fisher Scientific, USA) under aseptic conditions and immediately transferred to the laboratory. The extracted teeth were then washed three to five times using 1× phosphatebuffered saline (PBS, Sartorius, Germany) solution in a sterile petri dish (Fig. 1b). The middle-third of the roots were scraped gently using a Bard-Parker's blade number 15 into a sterile petri-dish and segmented into pieces of 1–3 mm size (Fig. 1c). The collected PDL tissues were digested in a solution of 3 mg/mL collagenase Type I (Gibco, UK) and 4 mg/ mL dispase (Gibco, UK) for 1 h at 37 °C. The mixture was then transferred through a 70 µm pore size cell strainer (BD Falcon Labware, Franklin Lakes, NJ, USA) to obtain single-cell suspension. The resulting cells were subsequently pelleted by centrifugation at 1.800 rpm and cultured in a 25 cm² filtered flask (Sarstedt, Germany) containing alpha modified of Eagle's medium (α-MEM Sigma-Aldrich, USA) supplemented with 15% fetal bovine serum (FBS; Gibco, UK), 2mM L-glutamine (Gibco, UK), and 100 U/mL penicillin/100 mg/mL streptomycin (Gibco, UK) for 7 days in a humidified atmosphere of 5% carbon dioxide (CO₂) at 37 °C, reaching 70-80% confluency under standard culture conditions. The medium was refreshed every 3-4 days. Once the cells reached 80% confluency, they were rinsed with PBS, harvested with 0.25% trypsin-EDTA (Sartorius, Germany), and centrifuged at 200×g for 5 min. The cells were stained with trypan blue (Sartorius, Germany) and counted using a TC20 Automated Cell Counter (Bio-Rad, USA) for further assays [30]. Experiments were performed using either passage (P) 4 or 5, and the cells were regularly tested for bacteria, fungi, and mycoplasma contamination using commercial PCR-based kit protocols (Thermo Scientific, USA).



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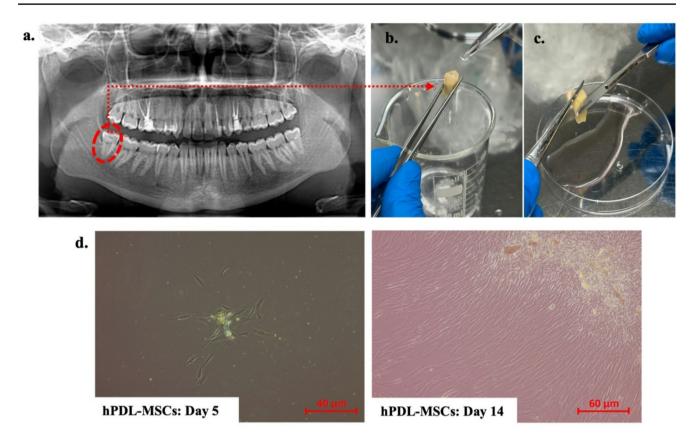


Fig. 1 Isolation of human periodontal ligament mesenchymal stromal cells (hPDL-MSCs). (a) Panoramic radiograph of the patient, (b) Washing the extracted teeth with 1× phosphate-buffered saline, (c)

Extracting the periodontal ligament tissue, (d) Micrographs of cells observed under inverted microscope on day 5 and day 14 after seeding

Flow cytometric analysis of hPDL-MSCs phenotype

To verify the identity of the isolated cells, expression of surface markers was analyzed using flow cytometry. Cells were cultured in 25 cm² polystyrene cell culture flasks in aMEM, washed twice carefully in PBS, detached from the culture flasks using 0.25% trypsin-EDTA, and subsequently centrifuged at 200xg for 5 min at 4 °C. Harvested cells were fixed with 4% paraformaldehyde for 30 min at 4 °C, and then permeabilized with 0.1% Triton X-100 for 10 min at room temperature, and blocked with a solution of 0.5% bovine serum albumin (BSA) and 0.02% Tween-20 for 20 min at room temperature. Next, the cells were incubated with primary monoclonal antibodies raised against CD31, CD34, CD45, CD73, CD90, and CD105 according to the standard quantities recommended by the manufacturer (Thermo Fisher Scientific, USA), at 37 °C for 1 h in the dark. After incubation, the cells were washed twice with PBS, and treated with fluorescent secondary antibody for 40 min at 4 °C in the dark [30]. The cells were then analyzed by flow cytometry (NovoCyte, ACEA Biosciences, USA) using FlowJo software (version 10.5.3, Tree Star, San

Carlos, USA). Unstained cells served as negative control were used to establish gating parameters.

Cell viability assays

An MTT assay (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) was conducted to evaluate the cytotoxic effects of postbiotics on hPDL-MSCs [31]. Briefly, 1×10^4 cells per well were seeded into 96-well cell culture plates (Sarstedt, Germany) and allowed to adhere for 24 h under standard culture conditions (37 °C with 5% CO₂ and 95% relative humidity). Subsequently, the culture medium was replaced with fresh α-MEM containing different concentrations of postbiotics, ranging from 2.5 to 1000 µg/ mL. Untreated cells were used as negative controls. Following 24 h of incubation, 10 µL of MTT solution (5 mg/mL, Sigma-Aldrich, USA) was added to each well, and the plates were incubated in darkness for 4 h. The formazan crystals formed by the viable cells were then solubilized in 100 µL of dimethyl sulfoxide (DMSO, Serva, USA). Finally, the optical density of each well was measured at 570 nm using a microplate reader (Epoch, USA).



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The Annexin V-FITC Apoptosis Detection Kit (Bio-Legend, USA) was employed to quantitatively assess cell viability via flow cytometry, following the manufacturer's instructions. A total of 1×10^5 cells per well were seeded into 6-well cell culture plates (Sarstedt, Germany) and treated with postbiotics for 24 h, as previously described. Untreated cells were used as negative controls. After incubation, the cells were stained with FITC-conjugated Annexin V and propidium iodide (PI) and subsequently analyzed using a flow cytometer. Data analysis was performed using FlowJo software.

Migration assay

The impact of postbiotics on the motility of hPDL-MSCs was assessed by using a scratch assay, as described by Liang et al. [32]. Briefly, cells were seeded in 6-well cell culture plates at a density of 1×10^5 cells per well and incubated under standard conditions until a confluent monolayer was formed. A sterile pipette tip was used to create a scratch across the center of each well, and cellular debris was removed by gently washing the cells twice with PBS. The cells were subsequently incubated in a fresh medium supplemented with postbiotics at the highest non-cyotoxic doses, as determined by the cell viability assay, while untreated cells served as controls. Cell migration into the wound area was monitored every 3 h using an inverted microscope (Olympus, Japan). The distance between the two edges of the wound area was measured using ImageJ software (NIH, USA), and the percentage of wound closure was calculated using the equation: $(At_0 - At/At_0) \times 100$, where At_0 is represents the scratch area at 0 h and At represents the corresponding scratch area at different time intervals.

Determination of intracellular reactive oxygen species (ROS) levels

Intracellular ROS levels were quantified by flow cytometry analysis. Hydrogen peroxide ($\rm H_2O_2$) was employed to induce ROS production in hPDL-MSCs. Initially, the effects of varying $\rm H_2O_2$ concentrations on hPDL-MSCs viability were determined using the MTT assay. The $\rm H_2O_2$ concentration that reduced cell viability by 50% ($\rm IC_{50}$) after 24 h of exposure and led to significant ROS accumulation was selected for subsequent treatments. Intracellular ROS levels in $\rm H_2O_2$ -challenged hPDL-MSCs, treated with or without postbiotics for 24 h, were evaluated using the Cellular ROS Assay Kit (Abcam, UK) according to the manufacturer's instructions. Briefly, cells were seeded in 6-well cell culture plates at a density of 1×10^5 cells per well and treated with postbiotics for 24 h under standard conditions. Untreated cells were used as negative controls. After treatment, cells

were trypsinized, centrifuged, and resuspended in 100 μL of 25 μM 2,7-dichlorodihydrofluorescein diacetate (DCFH-DA) solution, a cell-permeable ROS-sensitive fluorescent probe. The suspension was incubated at 37 °C for 30 min in darkness. ROS accumulation was subsequently measured at an excitation wavelength of 485 nm and an emission wavelength of 528 nm using a fluorescence multi-detection reader integrated with flow cytometry. Data analysis was performed using FlowJo software.

Measurement of immunological markers

The effect of postbiotics on cytokine production in hPDL-MSCs was assessed following lipopolysaccharide (LPS)induced inflammation. LPS derived from P. gingivalis (MQ300, Sigma-Aldrich, USA) was used as the inflammatory agent. Initially, hPDL-MSCs cells were cultured in α-MEM medium with or without different concentrations of P. gingivalis LPS (0 µg/mL, 1 µg/mL, 5 µg/mL, and 10 µg/mL) for 24 h [33]. The lowest non-cytotoxic dose of LPS was determined via MTT assay as described previously. hPDL-MSCs were then seeded at a density of 1×10^4 cells per well in 96-well cell culture plates and treated with the non-cytotoxic concentration of LPS, either alone or in combination with postbiotics. Following a 24 h incubation period, cytokine levels, including interleukin (IL)-10, IL-8, IL-6, and IL-1β, were quantified in the cell culture supernatants using Enzyme-Linked Immunosorbent Assay (ELISA) kits (MabTech, Switzerland), in accordance with the manufacturer's protocols.

Analysis of collagen deposition

The expression of collagen type I (COL1A1) gene was assessed using quantitative reverse transcription PCR (qRT-PCR) and immunofluorescence staining. For qRT-PCR, total RNA was extracted from hPDL-MSCs (1×10^6 cells per well) treated with or without postbiotics for 24 h, using PureZOLTM reagent (Bio-Rad, Hercules, CA, USA) according to the manufacturer's protocol. Following RNA extraction, complementary DNA (cDNA) was synthesized using the iScriptTM cDNA Synthesis Kit (Bio-Rad, USA), qRT-PCR was performed on a Bio-Rad CFX96 Touch Real-Time PCR system using specific primers designed for the genes of COL1A1 (F:5'-CCCTCCACTCCTTCCCAAAT-3'; R:5-CGAAGGGGAGATGTTGCAAG-3') and β -actin (F:5'- G CTCTTTTCCAGCCTTCCTT-3'; R:5'-CAGCAATGCC AGGGTACATG-3') as an internal control, using Primer3 software and the NCBI-Primer BLAST program. The PCR reaction mixture included 5 µL of 1x SYBR Green Master Mix (Bio-Rad, USA), 0.5 µL of forward and reverse primers (0.5 mM each), and 1 µL of cDNA. Amplification was



carried out under the following conditions: initial denaturation at 95 °C for 3 min, followed by 40 cycles at 95 °C for 10 s, annealing at 60 °C for 30 s, and DNA strand synthesis at 72 °C for 30 s. The relative quantification of target gene expression was expressed as threshold cycle numbers (Ct) and normalized to internal controls. The fold change in gene expression was quantified using the $2^{-\Delta\Delta Ct}$ method, where ΔCT represents the difference between the mean CT values of triplicate samples.

Immunofluorescence detection of COL1A1 was performed using fluorescence microscopy. Briefly, hPDL-MSCs $(1 \times 10^6 \text{ cells/mL})$ grown on glass coverslips with or without postbiotics for 24 h were fixed in 4% paraformaldehyde (PFA, Sigma-Aldrich, USA) in PBS for 20 min. The fixed cells were subsequently treated with a primary antibody against COL1A1 (collagen type I). This was followed by the application of a secondary antibody conjugated with FITC (Fluorescein isothiocyanate). Nuclear DNA was stained using 4',6-diamidino-2-phenylindole (DAPI, Thermo Fisher Scientific, USA), which emits blue fluorescence for cellular nuclei visualization. Slides were then mounted in ProLong Gold Antifade Reagent (Invitrogen, USA) and examined using a fluorescence microscope (Olympus BX60) equipped with a Color View III cooled CCD camera. The excitation wavelength was set to 488 nm, and the emission wavelength was monitored at 520 nm for capturing fluorescence signals.

Data analysis

All assays were conducted across three independent experiments, with each measurement performed in triplicate. Data analysis was carried out using GraphPad Prism software version 8.0.2 (San Diego, CA, USA). The student's t-test was employed to compare the means between two groups, while analysis of variance (ANOVA) was utilized to compare the means across three or more groups. All results were presented as the mean \pm standard deviation and p-values less than 0.05 were considered to indicate a statistically significant difference.

Results

Characteristics of hPDL-MSCs

hPDL-MSCs were isolated from wisdom teeth using enzymatic digestion and cultured in the α -MEM. Initially, the cells appeared small and thin by day 3, exhibiting a spindle-shaped morphology characteristic of fibroblast-like cells by day 5. By day 14, the fibroblast-like cells reached approximately 80% confluence (Fig. 1d). The hPDL-MSCs

were further characterized through cell surface marker analysis. Flow cytometry results revealed that hPDL-MSCs expressed mesenchymal stem cell-associated markers, including CD73 (95.90%), CD90 (95.80%), and CD105 (94.81%), while lacking expression of hematopoietic markers such as CD34 (2.72%) and CD45 (2.19%), as well as the endothelial marker CD31 (0.65%) (Supp. Figure 1).

Proliferative effects

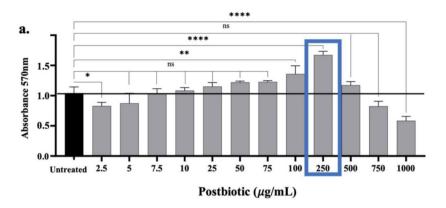
To evaluate the effects of postbiotics on the viability of hPDL-MSCs, an MTT assay was performed. Increasing doses of postbiotics resulted in a plateau effect, with the highest cell viability observed at a concentration of 250 µg/ mL (p < 0.0001). No significant effects were noted at doses ranging from 5 to 75 μg/mL and 500–750 μg/mL. However, a significant increase in cell viability was also observed in 100 μg/mL (p < 0.01). In contrast, treatment with 1000 μg/ mL led to a significant decrease in cell viability (p < 0.0001) (Fig. 2a). Since the MTT assay only provides data on cell viability or cytotoxicity at a single time point, additional experiments were conducted using Annexin V-FITC/PI staining to gain a more comprehensive understanding of the proliferative effects of postbiotics on hPDL-MSCs. The flow cytometry results confirmed the findings from the MTT assay. Compared to untreated cells (86.90% cell viability), a significant increase in the proportion of live hPDL-MSCs (95.64%) was observed following the treatment with 250 μg/ mL of postbiotics (p < 0.05), as indicated by the lower left quadrant of the flow cytometry plots, representing viable cells negative for Annexin-FITC and PI. Furthermore, a significant reduction (p < 0.001) in the proportion of live cells (56.06%) was observed when treated with 1000 µg/mL of postbiotics (Fig. 2b). Based on these findings, the 250 µg/ mL postbiotic concentration which exhibited the most pronounced effects on hPDL-MSCs, was selected for further assays.

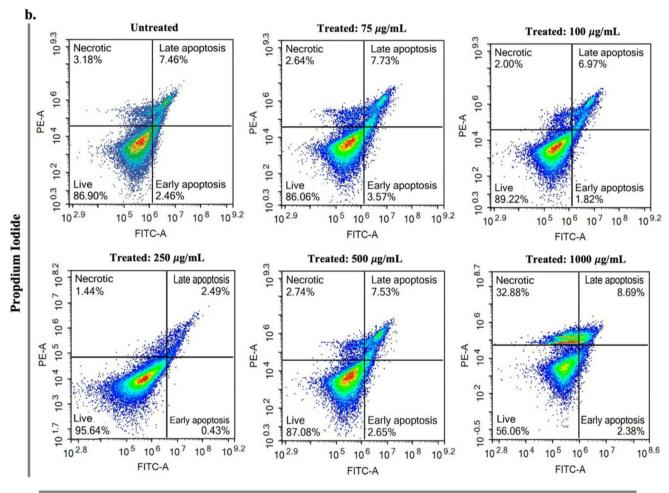
Induction of migration

The effect of postbiotics (250 µg/mL) on the migration ability of hPDL-MSCs was assessed using a scratch assay. At 24 h post-injury, the wound area was reduced to $6.86 \times 10^8 \pm 0.28~\mu m^2$, corresponding to a significant relative wound closure of $97.61 \pm 4.71\%~(p < 0.05)$ compared to the control. In addition, untreated cells achieved $82.09 \pm 3.54\%$ wound closure (Fig. 3a). Microscopic observation further revealed that the scratch wound in the treated wells was significantly filled with cells, which was attributed to accelerated cell proliferation compared to the untreated samples (Fig. 3b), after 24 h of treatment.



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Annexin V

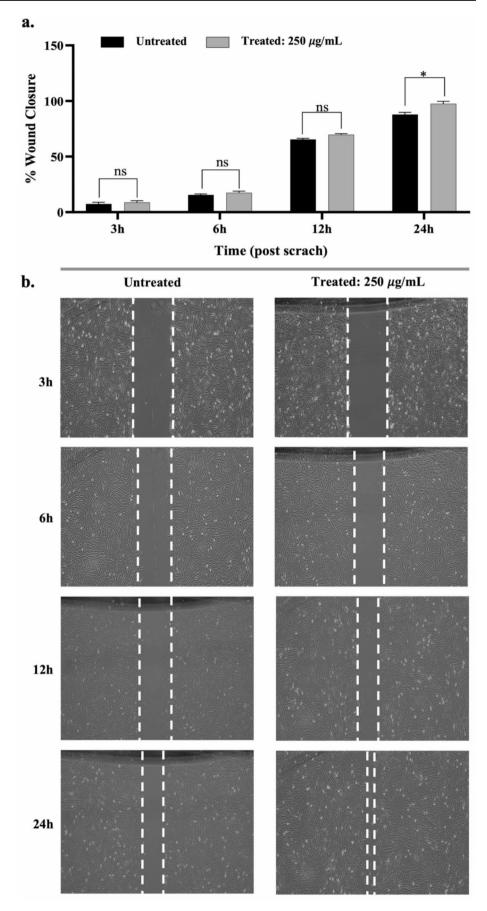
Fig. 2 Cell viability of hPDL-MSCs following postbiotic treatment, assessed by (a) MTT assay and (b) flow cytometry analysis using Annexin V-FITC/PI staining. The flow cytometry quadrants represent necrotic cells (PI-positive only), late apoptotic cells (annexin V and

PI double-positive), early apoptotic cells (Annexin V-positive only), and live cells (non-apoptotic cells). Data are expressed as mean \pm SD (n=3). Statistical significance is denoted as follows: * p<0.05, *** p<0.01, ***** p<0.0001, n.s: non-significant



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Fig. 3 Effects of postbiotics on the migration capability of hPDL-MSCs. (a) Wound closure (%) at different time points post-injury, (b) Representative photomicrographs of hPDL-MSCs during scratch assay at the indicated time points, captured using a 10x objective on an inverted microscope. Data are expressed as mean \pm SD (n=3). Statistical significance is denoted as follows: * p<0.05, n.s. non-significant





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Reduction of ROS accumulation

To induce oxidative stress in hPDL-MSCs, a concentration of 300 µM H₂O₂ was selected, which reduced the viability to $50.69 \pm 1.91\%$ (Fig. 4a and c). Notably, postbiotic treatment significantly (p < 0.0001) improved the viability of H₂O₂induced hPDL-MSCs compared to untreated H₂O₂-induced cells, as observed through both MTT and flow cytometry analyses (Fig. 4b and c). The effects of postbiotics on H₂O₂induced hPDL-MSCs were further assessed by flow cytometry using DCFDA dye to measure intracellular ROS levels. The results showed that treatment with 300 µM H₂O₂ led to a significant increase in intracellular ROS levels (5.23fold, p < 0.0001). However, postbiotic treatment for 24 h resulted in a 54.53 ± 2.01% reduction in intracellular ROS levels compared to untreated H₂O₂-induced hPDL-MSCs (p < 0.0001). Notably, postbiotic treatment alone did not significantly affect the intracellular ROS levels in hPDL-MSCs (Fig. 4d).

Production of pro- and anti-inflammatory mediators

The anti-inflammatory potential of postbiotics was further evaluated using LPS-induced hPDL-MSCs. To determine a non-cytotoxic concentration of LPS for hPDL-MSCs, an MTT assay was performed. The results showed that the lowest concentration of LPS (1 µg/mL) did not significantly affect the viability of hPDL-MSCs and was therefore selected for subsequent experiments (Fig. 5a). The effect of postbiotics on the production of IL-8, IL-6, IL-1β and IL-10 involved in the inflammatory and anti-inflammatory response, was assessed by ELISA analysis. The results indicated that LPS, acting as a pro-inflammatory stimulus, significantly increased the production of IL-8 (12.5-fold, p < 0.0001), IL-6 (27.5-fold, p < 0.001), and IL-1 β (6.91fold, p < 0.0001) in untreated hPDL-MSCs. Conversely, treatment with postbiotics markedly attenuated the levels of these pro-inflammatory cytokines in LPS-stimulated hPDL-MSCs. Specifically, postbiotics significantly reduced IL-8 levels by 1.36-fold compared to the elevated secretion of IL-8 in untreated LPS-stimulated hPDL-MSCs (p < 0.0001). Similarly, production of IL-6 and IL-1β were also reduced by 2.33-fold and 1.15-fold, respectively, in postbiotic-treated cells relative to untreated LPS-stimulated hPDL-MSCs. Notably, postbiotics significantly enhanced the production of IL-10, an anti-inflammatory cytokine, in LPS-induced hPDL-MSCs (2.67-fold, p < 0.0001) (Fig. 5b).

Enhancement of collagen deposition

The effects of postbiotics on type I collagen synthesis in hPDL-MSCs were evaluated using qRT-PCR and

immunofluorescence staining. Treatment with 250 μ g/mL of postbiotics for 24 h significantly upregulated the COL1A1 mRNA expression by 1.36-fold (p<0.01) compared to untreated cells (Fig. 6a). Furthermore, Furthermore, immunofluorescence analysis demonstrated increased COL1A1 production in hPDL-MSCs treated with postbiotics, as shown in Fig. 6b. These findings confirm that postbiotic treatment enhances type I collagen production in hPDL-MSCs.

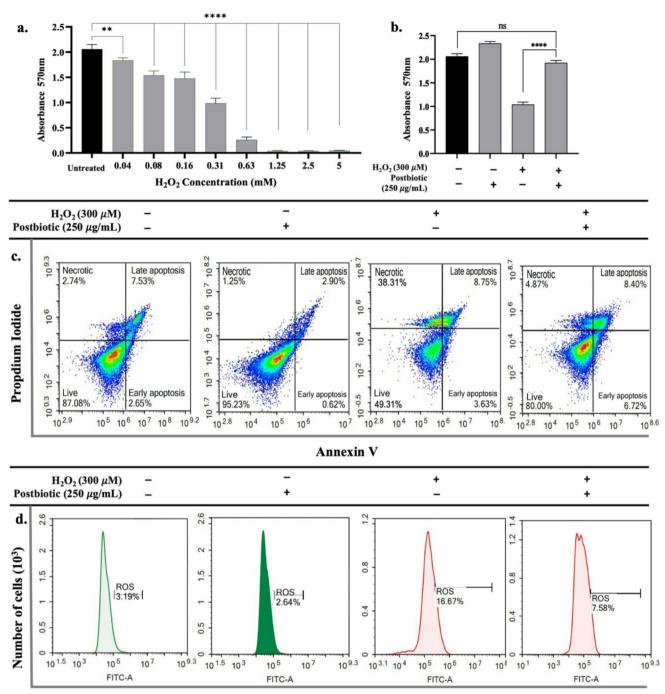
Discussion

Periodontal defects caused by periodontitis or trauma continue to pose significant challenges in the field of dentistry. The structural and biological integrity of the PDL, which plays an essential role in maintaining periodontal health, is critical for the long-term prognosis of periodontal tissues. The primary objective of effective periodontal treatment is to fully restore all components of the periodontium to their original architecture and functionality [34]. However, the limitations of conventional restoration methods in achieving functional periodontal tissue regeneration underscore the urgent need for innovative strategies. In recent years, tissue engineering, a multidisciplinary approach that integrates cells, biomaterials, and bioactive factors, has emerged as a tremendous future in addressing these challenges [17]. Among these approaches, hPDL-MSCs-based therapies hold great potential for achieving functional PDL restoration. These multipotent cells are characterized by minimal criteria, including plastic adherence, and in vitro trilineage differentiation to osteoblasts, adipocytes, and chondrocytes. Furthermore, when cultured, hPDL-MSCs exhibit the expression of specific surface markers such as CD73, CD90 and CD105, while lacking the expression of hematopoietic markers (CD34 and CD45) and the endothelial marker CD31 [14]. Within the scope of this study, hPDL-MSCs were isolated from healthy wisdom teeth and their immunophenotyping analysis confirmed the expression of characteristic surface markers, thereby validating their identity as hPDL-MSCs.

Recently, the application of hPDL-MSCs in tissue repair and regeneration has consistently attracted the attention of researchers. However, a major limitation in the clinical use of hPDL-MSCs for therapeutic purposes is the insufficient cell populations capable of migrating to the injury site and proliferating effectively. To address these limitations, various cytokines, growth factors, chemicals, and natural compounds have been explored, either individually or in combination, as culture media additives [17]. In this study, we propose a novel approach combining postbiotics with hPDL-MSCs and emphasize the existing gaps in their



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DCF-DA Fluorescence

Fig. 4 Effects of postbiotics on the H_2O_2 -induced ROS levels in hPDL-MSCs. (a) Cell viability of hPDL-MSCs following treatment with various concentrations of H_2O_2 , (b) Viability of hPDL-MSCs treated with selected doses of H_2O_2 with or without postbiotics, as determined by MTT assay and (c) Flow cytometry analysis using Annexin V-FITC/PI

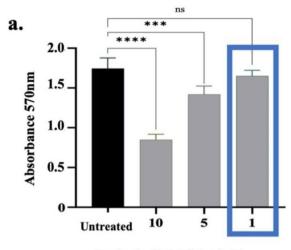
staining, (d) Intracellular ROS levels in hPDL-MSCs, represented as fluorescence intensity (%), following treatment with selected concentrations of H_2O_2 with or without postbiotics, assessed by flow cytometry. Data are expressed as mean \pm SD (n=3). Statistical significance is denoted as follows: ** p<0.01, **** p<0.0001, n.s: non-significant

application for periodontal regeneration. To date, experimental evidence has demonstrated that probiotics contribute to tissue repair by secreting postbiotics [25, 35–37]. Although substantial evidence highlights the beneficial

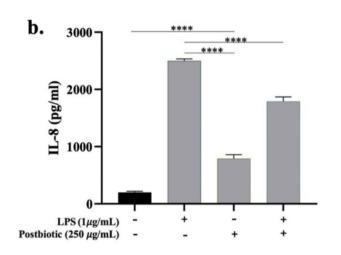
roles of postbiotics, neither their exact role on the dental stem cell proliferation nor the relevance of regulation of the periodontium remains unclear. Therefore, the current study aims to investigate the regenerative and protective effects

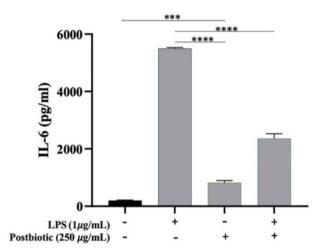


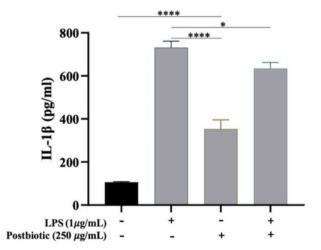
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P. gingivalis LPS (µg/mL)







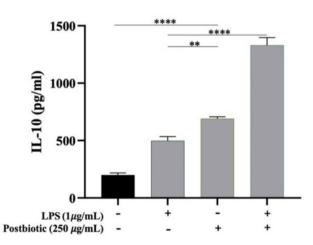
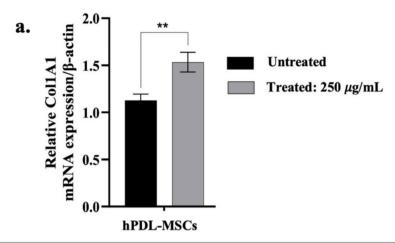


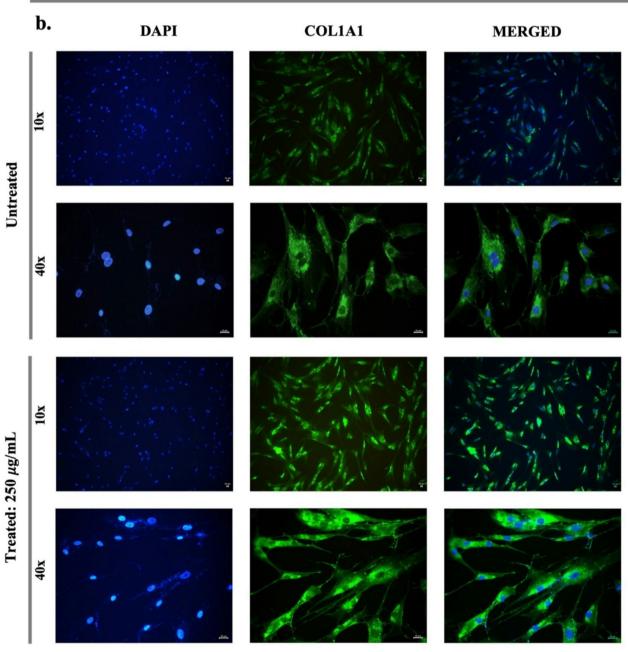
Fig. 5 Immunomodulatory effects of postbiotics. (a) Determination of the lowest non-cytotoxic concentration of LPS using the MTT assay, (b) Levels of the *IL-8*, *IL-6*, *IL-1β* and *IL-10* in LPS-induced

hPDL-MSCs treated with or without postbiotics. Data are expressed as mean \pm SD (n=3). Statistical significance is denoted as follows: *** p<0.001, **** p<0.0001, n.s: non-significant



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◆ Fig. 6 Collagen type 1 (COL1A1) production in hPDL-MSCs treated with postbiotics. (a) Gene expression levels assessed via qRT-PCR analysis, (b) Immunofluorescence detection visualized using fluorescence microscopy (COL1A1, green fluorescence; cellular nuclei: DAPI-stained, blue fluorescence). Data are expressed as mean±SD (n=3). Statistical significance is denoted as follows: ** p < 0.01
</p>

of postbiotics derived from L. plantarum EIR/IF-1 strain on hPDL-MSCs in the context of periodontal diseases. Our findings revealed that postbiotics, at a concentration of 250 µg/ mL, significantly enhanced the proliferation and migration of hPDL-MSCs. Furthermore, postbiotics enhanced cellular proliferation, thereby promoting the production of new collagen fibers by hPDL-MSCs. Consistent with our findings, Wu et al. [38] recently reported that postbiotics from L. reuteri stimulated the proliferation of human-induced pluripotent stem cells (h-iPSC) at lower doses but inhibited their proliferation at higher concentrations. They further demonstrated that sodium butyrate, a well-characterized bacterial metabolite and short-chain fatty acid present in postbiotics, exhibited similar effects on h-iPSC proliferation. Additionally, other recent studies have highlighted that Lactobacillus-derived postbiotics, such as lactate, enhance cell migration and tissue repair by promoting cellular proliferation [39]. Similarly, Neiva et al. [40] observed that B-vitamin complex supplementation accelerated wound healing following periodontal flap surgery. Based on this cumulative evidence, the primary components of postbiotics derived from L. plantarum EIR/IF-1, comprising lactate, stearic acid (C18:0), palmitic acid (C16:0), and B-complex vitamins (B1 thiamine, B5 pantothenic acid, and B12 cobalamin) as determined in our previous study [27], may contribute to their observed mechanism of action. However, further studies are warranted to elucidate the detailed effects of postbiotics on periodontal healing.

Although regenerative periodontal therapy aims to reconstruct periodontal supporting tissues [41], repairing lesions in this region remains a significant challenge due to the intricate structure of periodontal tissue, the surrounding inflammatory microenvironment, and oxidative damage [42]. In this context, postbiotics with their immunomodulatory and antioxidant properties, offer potential for maintaining periodontal tissue homeostasis. From this point of view, our findings demonstrated that postbiotics from L. plantarum EIR/IF-1 strain increased IL-10 levels, while decreasing IL-8, IL-1β and IL-6 levels in comparison to *P. gingivalis* LPS, thereby mitigating periodontal inflammation. To date, numerous studies have demonstrated the immunomodulatory effects of postbiotics through the regulation of cytokine production [43–45]. Consistent with our results, postbiotics derived from L. plantarum PD18 exhibited immunomodulatory properties in P. gingivalis LPS-induced cells, as evidenced by increased IL-10 levels alongside reduced IL-6 and IL-8 levels [46]. Additionally, recent studies have shown that postbiotics from the *L. curvatus* MG5246 strain downregulated IL-6 expression, a major pro-inflammatory cytokine, in human gingival fibroblast cells and demonstrated immunomodulatory effects in an in vivo mouse model [47].

Considering the close relationship between the various autoimmune or inflammatory oral conditions and cellular degeneration due to increased oxidative stress induced by oxygen-derived free radicals, maintaining homeostasis between oxidative and antioxidant mechanisms is critical for periodontal health [48]. Beyond their role in immune modulation, postbiotics derived from the L. plantarum EIR/ IF-1 strain also act as a natural potent antioxidant. Similar to our results, postbiotics from L. brevis BK3, isolated from Kimchi, a traditional Korean fermented food, have been shown to protect fibroblast cells from oxidative stress induced by H₂O₂ and to reduce intracellular ROS production [49]. Moreover, several studies have also demonstrated that postbiotics as metabolic byproducts of probiotics, such as short-chain fatty acids, and vitamins, may mitigate oxidative damage and reduce the inflammatory burden in periodontal regeneration by modulating signaling pathways, including the Nrf2 pathway [48, 50]. Aligned with these findings, a study by Huang et al. [51] revealed that smallmolecule peptides in postbiotics derived from L. plantarum SCS2 activate the Keap1-Nrf2 signaling pathway, thereby alleviating oxidative damage in H₂O₂-induced rat insulinoma cells. Additionally, postbiotics from L. reuteri SJ-47 contribute to antioxidant defense mechanisms by reducing ROS levels and upregulating antioxidant enzyme expression, ultimately exerting a protective effect on human fibroblasts under oxidative stress [52].

Since periodontal diseases are non-communicable and closely associated with an unbalanced oral microbiota, the application of microbial modulators, including postbiotics, shows promising beneficial effects and warrants further research [53]. Based on studies conducted on patients, lozenges or tablets containing postbiotics have been shown to reduce caries incidence in preschool and schoolchildren compared to standard preventive care. For adults with periodontitis, the adjunctive use of synbiotic and postbiotic products appears to enhance the outcome of conventional scaling and root planing [53]. In another study, a soothing gel containing postbiotics (Lactobacillus Ferment) was found to be a valuable adjunct to non-surgical periodontal treatment, improving periodontal health in patients with Down syndrome and reducing BOP (Bleeding on Probing) after 6 months of treatment [54]. Basir et al. [55] investigated the effect of postbiotic toothpaste (Bifidobacterium animalis subsp. animalis) on salivary levels of Immunoglobulin A (IgA) and pH in children in a triple-blind,



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randomized, placebo-controlled trial. According to the clinical study results, the administration of postbiotics increased oral immunity in children. A recent study also showed that postbiotics co-treatment with dental fillings could improve prognosis for patients with dental caries compared to dental fillings alone [56]. All these results suggest that postbiotics could have a significant role in clinical applications. Therefore, our findings provide a more comprehensive understanding of the therapeutic potential of postbiotics in the context of periodontal regeneration and support their integration into dental product such as toothpaste, mouthwash, scaffolds and gels, as a natural, cost-effective, novel, and safe bio-material. The successful application of postbiotic preparations in the clinical setting can provide crucial support and greater assurance for oral health. However, several questions remain, including the dose-response effects, the most effective mode of administration, and suitable combinations with prebiotics and probiotics [53]. Investigating the molecular mechanisms underlying the immune-modulatory and antioxidant effects of postbiotics is also critical for advancing their clinical application in regenerative medicine. Another knowledge gap is the lack of pre-clinical and clinical studies, which would be essential to assess the sustained effects of postbiotics in periodontal regeneration.

Conclusion

In the present study, we found consistent evidence that postbiotics derived from the *L. plantarum* EIR/IF-1 exhibit significant biological activity throughout all stages of the healing process, ranging from the modulation of initial inflammatory response to LPS stimulation, followed by the promotion of cell migration, proliferation, and collagen synthesis, ultimately improving long-term regenerative potential. Future studies should focus on improving clinical outcomes and enhancing patient quality of life in the fields of oral medicine and dentistry.

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Data availability Data is provided within the manuscript.

Declarations

Ethical approval The research protocol was approved by the Advisory Board of Karamanoglu Mehmetbey University, Karaman, Turkey (Protocol code: 01-2023/20). All examinations were conducted in compliance with the ethical standards set by the Institutional Review Board (IRB) and the modified Helsinki Declaration. Prior to all examinations, written informed consent was obtained from all participants.

Informed consent Written informed consent was obtained from all participants included in the study.

Competing interests The authors declare no competing interests.

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