

Preliminary findings on the possible role of B-lymphocyte stimulator (BLyS) on diabetes-related periodontitis

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Abstract: The possible role of B-cell growth and differentiation-related cytokines on the pathogenesis of diabetes-related periodontitis has not been addressed so far. The aim of this study was to evaluate the effects of diabetes mellitus (DM) on the gene expression of proliferation-inducing ligand (APRIL) and B-lymphocyte stimulator (BLyS), two major cytokines associated to survival, differentiation and maturation of B cells in biopsies from gingival tissue with periodontitis. Gingival biopsies were obtained from subjects with periodontitis (n = 17), with periodontitis and DM (n = 19) as well as from periodontally and systemically healthy controls (n = 10). Gene expressions for APRIL, BLyS, RANKL, OPG, TRAP and DC-STAMP were evaluated using qPCR. The expressions APRIL, BLyS, RANKL, OPG, TRAP and DC-STAMP were all higher in both periodontitis groups when compared to the control group (p < 0.05). Furthermore, the expressions of BLyS, TRAP and RANKL were significantly higher in the subjects with periodontitis and DM when compared to those with periodontitis alone (p < 0.05). The mRNA levels of BLyS correlated positively with RANKL in the subjects with periodontitis and DM (p < 0.05). BLyS is overexpressed in periodontitis tissues of subjects with type 2 DM, suggesting a possible role of this cytokine on the pathogenesis DM-related periodontitis.

Keywords: Periodontitis; Diabetes Mellitus; Inflammation; Cytokines.

Introduction

Periodontitis has long been linked to diabetes mellitus (DM).^{1,2} Nowadays, DM is considered a risk factor for periodontitis and has to be included as a descriptor in grading periodontitis based on the new clinical classification of periodontal diseases.³ The pathogenesis of DM is multifactorial, involving several metabolic and hemodynamic disorders like insulin resistance, dyslipidemia, hypertension, hyperglycemia and immune-inflammatory dysfunctions. The immune-inflammatory dysfunctions associated to DM include increase in reactive oxygen species and a shift to an overall pro-inflammatory profile.^{4,5}

In recent years, the role of DM in modulating mediators involved in the pathogenesis of periodontal diseases has been studied by our research group. In brief, higher levels of pro-inflammatory T-helper (Th)1 - and



Th17-cytokines have been reported in periodontitis from type 2 diabetic subjects when compared to non-diabetics.^{6,7,8,9} Furthermore, subjects with periodontitis and type 2 DM exhibited elevated ratio of receptor activator of nuclear factor kappa-B ligand (RANKL)/osteoprotegerin (OPG), a key indicator of bone resorption,¹⁰ as well as increased expression of sclerostin and Dickkopf-1, important inhibitors of the Wnt/ β -catenin pathway and suppressors of bone formation.¹¹ Moreover, increased expression of both peroxiredoxin II and superoxide dismutase II has been observed in sites with periodontitis in diabetic subjects, probably as a consequence of protective and adaptive mechanisms against increased levels of reactive oxidative species, as seen in a DM context.¹²

Although most studies have focused on the role of T cell-mediated immunity in the pathogenesis of periodontal diseases, pre-clinical and clinical studies have indicated a critical contribution of B cell to periodontal breakdown. B cell-deficient animals were protected from bacterial infection-induced alveolar bone loss.^{13,14} Furthermore, the number of multiple subsets of B and plasma cells exceeds the number of T cells in established periodontitis in human.^{15,16} A possible role of B cells on osteoclastogenesis and alveolar bone loss during periodontitis has therefore been proposed. A previous study demonstrated that approximately 90% of B cells express RANKL in periodontitis lesions, whereas the percentage of RANKL-positive B cells in healthy gingiva is very low.¹⁷ Furthermore, there was a significant increase in B-cell RANKL expression in mice infected with *Porphyromonas gingivalis*.¹⁴ Moreover, memory B cells supported osteoclast differentiation *in vitro* in a RANKL-dependent manner, and B cells in the gingiva of animals submitted to experimental periodontitis produced RANKL and sustained osteoclastogenesis beyond T and other lymphocytes.^{18,19} Finally, B lineage cells were an important source of secreted osteoclastogenic factor of activated T cells (SOFAT) that induces bone resorption in a RANKL-independent manner in inflammatory states.²⁰

Two major cytokines belonging to the tumor necrosis factor (TNF) superfamily, named proliferation-inducing ligand (APRIL; also identified as TNFSF13 or TALL-2) and B-lymphocyte stimulator

(BLyS; also identified as B cell-activating factor [BAFF], TNFSF13B, or TALL-1), have been proposed as essential factors to the survival, differentiation and maturation of B cells.²¹ Overexpression of BLyS and APRIL has been linked to development of several autoimmune diseases, such as systemic lupus erythematosus, Sjögren syndrome, immune thrombocytopenic purpura, rheumatoid arthritis and even neoplastic lesions.²² Previous studies demonstrated that subjects with periodontitis presented higher serum levels of APRIL and BLyS than periodontally-healthy subjects.^{23,24} In addition, the expression of APRIL and BLyS at mRNA and protein levels was locally higher in periodontitis in human and in experimental periodontitis in mice.²⁵

Despite some evidence on the possible role of APRIL and BLyS on the pathogenesis of periodontitis, the impact of DM on this mechanism has seldom been explored. Therefore, the main aim of this study was to evaluate the effects of DM on BLyS and APRIL mRNA expression in gingival biopsies with periodontitis. The secondary aim was to correlate the expression of APRIL and BLyS with genes related to osteoclast formation and activity, including RANKL and OPG, markers of osteoclastogenesis, tartrate-resistant acid phosphatase (TRAP), an osteoclastic differentiation marker, and dendritic cell-specific transmembrane protein (DC-STAMP), an essential protein for cell fusion during osteoclastogenesis.

Methodology

Systemically healthy subjects without periodontitis (controls), systemically healthy subjects with periodontitis and type 2 diabetic subjects with periodontitis were sequentially selected from the population referred to the Periodontology Clinic of Guarulhos University (Guarulhos, Brazil) between March 2014 and September 2016. Qualified individuals were invited to participate in the study, fully informed of the nature, risks and benefits of the procedures and signed their informed consent. During volunteer screening, medical and dental histories were obtained. This study was approved by the Research Ethics Committee of the Guarulhos University (CAAE: 25526913.8.0000.5506).

The inclusion criteria were individuals aged 30 years or older, presenting a minimum of 15 teeth, excluding third molars. Pregnant or breastfeeding women were excluded, as well as smokers and those with a history of subgingival periodontal therapy in the 6 months preceding the start of the study. Furthermore, individuals frequently using mouth rinses containing antimicrobials in the previous 2 months, systemic conditions other than DM that could influence the pathogenesis of periodontitis and the expression of the studied proteins [e.g. immunological disorders, bone-related diseases e complications (e.g., osteoporosis, ankylosing spondylitis, recent bone fractures, and rheumatoid arthritis)], use of antibiotics in the preceding 6 months, long-term use of anti-inflammatory, immunosuppressive (e.g., glucocorticoids) and antiresorptive agents (e.g., bisphosphonates and denosumab), hormone replacement therapy and orthodontic treatment were also excluded.

Inclusion criteria for subjects with periodontitis were as follow: generalized moderate or severe chronic periodontitis²⁶ including > 30% of sites presenting probing depth (PD) and clinical attachment level (CAL) \geq 4 mm with bleeding on probing (BoP), \geq 6 teeth with \geq 1 sites with PD and CAL \geq 5 mm and BoP, \geq 1 tooth indicated for extraction due to severe periodontitis. According to the new classification patients with generalized stage 3 or 4, grade C.²⁷ Subjects with no periodontitis should have had less than 10% positive marginal bleeding and/or BoP, no sites with both PD and CAL >3 mm and at least one region needing crown lengthening. The diabetic subjects had to be formally diagnosed with type 2 DM by a physician (HbA1c > 6.5% and FPG > 99 mg/dl) for at least 3 years prior to the start of the study. In order to attain expressive areas of periodontal inflammation, gingival biopsies were collected from a tooth referred for extraction due to severe periodontitis (PD and CAL \geq 7 mm, BoP, mobility and/or bone loss compromising more than 50% of the root) in the subjects with periodontitis.

Clinical examinations

Plaque index (PI), BoP, PD (mm), and CAL (mm) were evaluated at six sites per tooth, except third

molars, by means of a manual periodontal probe (UNC15; Hu-Friedy, Chicago, USA) by the same examiner (TSM), who was trained and calibrated using a method described elsewhere.²⁸

Gingival tissue sampling

In subjects with no periodontitis, the gingival biopsies were sampled from teeth without signs of clinical inflammation (BoP and/or MB), but requiring crown lengthening procedures. All samples included junctional and sulcular epithelia and connective gingival tissue. The gingival tissues were stored in TRIzol (Thermo Fisher Scientific, Waltham, MA) at -80°C until RNA extraction.

Gene expression

The mRNA levels corresponding to BlyS, APRIL, RANKL, OPG, TRAP, DC-STAMP and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were assessed using Real Time PCR. The primer sequences are presented in Table 1.

RNA isolation

Total RNA from the gingival biopsies was isolated using the Trizol method (Gibco BRL, Life Technologies, Rockville, USA), according to the manufacturer's

Table 1. Gene and primer sequence.

Gene	Sequence
BlyS	F: AAC AGA CAG CCA CAG CCA AA R: TGC CCC TTT GAA TTG TGT CCT
APRIL	F: AGC ACT CTG TCC TGC ACC T R: CGG ACA CCA TAT CCT TGG GC
RANKL	5' TACACGACTCAGTATCCATGC 3' AAGGTCAACCCGTAATTGC
OPG	5' TCCGGAACAGTGAATCAACTC 3' TCTCTGCGTTTACTTTGGTGC
DC-STAMP	F: TCC TCC ATG AAC AAA CAG TTC CAA R: AGA CGT GGT TTA GGA ATG CAG CTC
TRAP	F: CCA GCG ACA AGA GGT TCC R: AGA GAC GTT GCC AAG GTG AT
GAPDH	F: ACC CAC TCC TCC ACC TTT GA R: TGT TGC TGT AGC CAA ATT CGT T

F: Forward; R: Reverse; Demographic characteristics, glycaemic and periodontal parameters (mean \pm SD); FM: full-mouth, BoP: bleeding on probing, PD: probing depth, CAL: clinical attachment level. Different letters indicate differences among groups.

recommendation. RNA samples were re-suspended in diethylpyrocarbonate-treated water and stored at -80°C. RNA concentration was established based on optical density using a micro-volume spectrophotometer (Nanodrop 1000, Nanodrop Technologies LLC, Wilmington, USA).

Reverse transcription

Total RNA was treated with DNase (Turbo DNA-frees, Ambion Inc., USA). Subsequently, 1 µg of DNase-treated RNA was used for complementary DNA (cDNA) synthesis. This reaction was performed using the First- Strand cDNA synthesis kit (Roche Diagnostic Co., USA), following the manufacturer's recommendations.

Quantitative real-time polymerase chain reaction (qPCR)

The qPCR reactions were done in the 7300 Real Time PCR (Applied Biosystem), using the SYBR Green PCR Master Mix (Thermo Fisher Scientific, Waltham, USA), according to the manufacturer's recommendations. The reaction products were calculated using the Relative Quantification tool, based on GAPDH as the reference gene. Negative controls with SYBR Green PCR Master Mix and water were used for all reactions.

Statistical analysis

As no previous study comparing the gene expression of BLyS and APRIL in tissues with periodontitis in diabetic and non-diabetic

patients to perform a sample power calculation, a *post hoc* analysis was performed to establish it. Considering differences of at least 1.1 in mRNA levels of BLyS between tissues with periodontitis of diabetic and non-diabetic subjects, and a standard deviation of 0.88, it was estimated that 9 subjects per group would be necessary to provide 85% power with an alpha of 0.05. The statistical analyses were performed on GraphPad Prism 6.0 (La Jolla, USA). Data were first examined for normality using the Kolmogorov-Smirnov test. Non-parametric methods were used for data that did not conform to the assumptions of normality. The mean percentages of sites with visible plaque accumulation, BoP, mean PD and CAL were computed for all teeth. Subsequently, the clinical parameters at full-mouth and sampled tooth levels as well as mRNA expression were averaged for each group. The differences among groups regarding clinical parameters, age and mRNA levels were compared using the Kruskal-Wallis and Dunn tests. Spearman's correlation was used for bivariate correlations between the expressions of the biomarkers studied. The significance level for all analyses was established at 5%.

Results

The intra-examiner variability (standard error of measurement) was 0.23mm for PD and 0.24mm for CAL. The concordance for BoP was 93% by Kappa-light test.

Table 2. Demographic characteristics, glycaemic and periodontal parameters (mean ± SD).

Variable	Healthy	Periodontitis	Periodontitis + DM
Gender (M/F)	2/8	5/12	6/13
Age (years)	34.6 ± 4.6 a	50.0 ± 10.1 b	56.6 ± 9.3 b
Years of DM	0	0	6.3 ± 5.3
HbA1c	6.0 ± 0.2 a	5.8 ± 0.2 a	7.8 ± 1.2 b
FM % of sites with plaque	11.1 ± 3.6 a	59.3 ± 24.5 b	73.8 ± 23.5 b
FM % of sites with BoP	9.0 ± 3.3 a	47.3 ± 15.2 b	40.4 ± 10.9 b
FM PD (mm)	1.7 ± 0.1 a	3.3 ± 0.9 b	3.7 ± 0.8 b
FM CAL (mm)	0.7 ± 0.9 a	4.3 ± 0.9 b	5.0 ± 1.0 b
Sampled teeth PD (mm)	2.6 ± 0.7 a	5.0 ± 1.6 b	5.7 ± 1.2 b
Sampled teeth CAL (mm)	0.8 ± 1.0 a	7.9 ± 2.2 b	7.4 ± 1.8 b

FM: full-mouth, BoP: bleeding on probing, PD: probing depth, CAL: clinical attachment level; Different letters indicate differences among groups

Gingival biopsies were collected from ten systemically healthy subjects without periodontitis, 17 systemically healthy subjects with periodontitis and 19 type-2 diabetic subjects, selected from almost 130 subjects screened. No samples were lost during laboratorial preparation.

Table 2 presents the demographic characteristics of the study population and the clinical parameters considering the sampled teeth and full-mouth levels. All clinical parameters were significantly lower in the control group, when compared to both periodontitis groups ($p < 0.05$, Table 2).

Figure presents the gene expression of BLyS, APRIL, RANKL, OPG, TRAP and DC-STAMP relative to GAPDH. The expressions of BLyS (Figure A), APRIL (Figure B), RANKL (Figure C), OPG (Figure D), RANKL/OPG (Figure E), TRAP (Figure F) and DC-STAMP (Figure G) were all significantly higher in both groups with periodontitis, when compared to the control ($p < 0.05$). In addition, individuals with periodontitis and DM exhibited higher BLyS, RANKL and TRAP expressions than the group with periodontitis alone ($p < 0.05$). APRIL, OPG, RANKL/OPG and DC-STAMP expression did not differ between periodontitis groups with or without DM ($p > 0.05$).

Table 3 presents the correlation coefficients between the expression of the biomarkers studied. In the control group, APRIL mRNA levels showed positive correlations with DC-STAMP and RANKL ($p < 0.05$). Moreover, the mRNA levels of DC-STAMP positively correlated with RANKL ($p < 0.05$). BLyS positively correlated with APRIL while TRAP positively correlated with RANKL in subjects with periodontitis alone and in subjects with periodontitis and DM ($p < 0.05$). Furthermore, the mRNA levels of BLyS correlated positively with RANKL in the subjects with periodontitis and DM ($p < 0.05$).

Discussion

This study is the first to assess the gene expression of BLyS and APRIL, two major cytokines that support B cells differentiation and proliferation in DM-related periodontitis, and to correlate their expression with those of osteoclast differentiation and activity markers.

The main results demonstrated an overall higher expression of both BLyS and APRIL in gingival tissues from subjects with periodontitis when compared to those from subjects without periodontitis. Most importantly, significantly higher expression of BLyS was observed in gingival biopsies from patients presenting with periodontitis and DM when compared to those from subjects with periodontitis alone. Furthermore, BLyS correlated positively with RANKL in subjects with periodontitis and DM. These findings suggest a possible involvement of BLyS in the pathogenesis of DM-related periodontitis.

In the current study, BLyS and APRIL expression were upregulated in gingival tissues of subjects with periodontitis when compared to those with no periodontitis. In support of these findings, previous investigations have demonstrated elevated salivary and serum levels of BLyS and APRIL in individuals with periodontitis when compared to periodontally healthy subjects.^{23,24} Furthermore, the gingival expression of APRIL and BLyS at mRNA and protein levels were upregulated in natural and experimental periodontitis in humans and mice, respectively, when compared to healthy controls²⁵. Altogether, these findings support the roles of both APRIL and BLyS in the pathogenesis of the periodontal diseases, possibly fostering the expansion and maintenance of the B cells, found in high proportions in periodontitis lesions.

In this study, gingival tissues with periodontitis also exhibited higher expression of RANKL, OPG, TRAP and DC-STAMP than periodontally healthy tissues. RANKL-RANK signaling regulates osteoclast differentiation, activation, and survival during physiological and pathological conditions. OPG protects excessive bone resorption by binding RANKL and preventing it from binding RANK.²⁹ Therefore, in line with the findings from the present study, an increased RANKL/OPG ratio has been found in subjects with periodontitis.^{30,31} TRAP is a member of the ubiquitously expressed enzyme family of the acid phosphatases that prompts dephosphorylation of bone matrix phosphoproteins like osteopontin and bone sialoprotein,³² being a recognized cytochemical marker of osteoclasts.³³ DC-STAMP is an important protein currently considered as a master regulator of the cell fusion step of osteoclastogenesis.^{34,35}

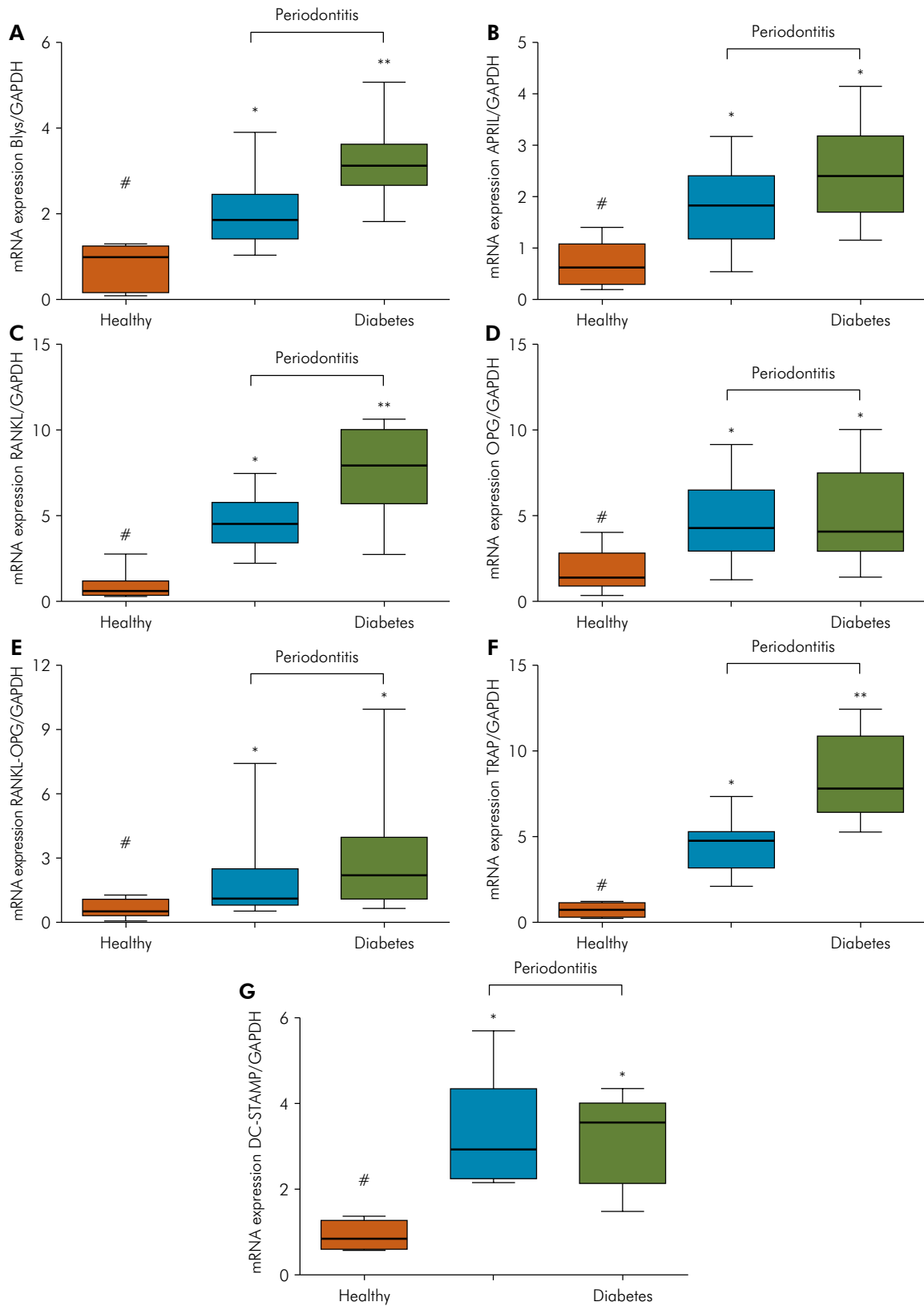


Figure. Relative expression of genes. qPCR analysis of BlyS (A), APRIL (B), RANKL (C), OPG (D), RANKL/OPG (E), TRAP (F) and DC-STAMP (G) expression on mRNA extracted from gingival tissue of healthy and periodontitis patients, associated or not with diabetes mellitus (DM). Different symbols indicate statistical significance amongst the groups ($p < 0.05$).

Table 3. Correlation coefficients among all biomarkers at mRNA levels.

Variable	BlyS	APRIL	RANKL	OPG	TRAP	DC-STAMP
Control						
BlyS	1.00	-0.36	0.25	-0.47	-0.06	0.40
APRIL	-	1.00	0.82*	0.66	0.05	0.97*
RANKL	-	-	1.00	0.36	0.03	0.98*
OPG	-	-	-	1.00	-0.08	0.77
TRAP	-	-	-	-	1.00	-0.07
DC-STAMP	-	-	-	-	-	1.00
Periodontitis						
BlyS	1.00	0.70*	0.22	0.03	0.10	0.22
APRIL	-	1.00	0.26	-0.11	0.02	0.09
RANKL	-	-	1.00	0.25	0.80*	0.41
OPG	-	-	-	1.00	0.01	-0.25
TRAP	-	-	-	-	1.00	0.06
DC-STAMP	-	-	-	-	-	1.00
Periodontitis/DM						
BlyS	1.00	0.75*	0.51*	-0.01	0.47	0.45
APRIL	-	1.00	0.32	-0.02	0.30	0.91
RANKL	-	-	1.00	0.22	0.72*	0.59
OPG	-	-	-	1.00	-0.46	-0.51
TRAP	-	-	-	-	1.00	-0.25
DC-STAMP	-	-	-	-	-	1.00

DC-STAMP-knockout mice presented significantly lower bone resorption in a ligature-induced periodontitis model compared to wild type mice, while local administration of anti-DC-STAMP-mAb downregulated the ligature-induced bone loss and the number of multinucleated TRAP⁺ cells, by downregulating cell fusion of osteoclast precursors cells in mice.^{36,37} Overexpression of the genes involved in the multiple steps leading to osteoclastogenesis observed in the tissues with periodontitis were therefore expected findings, which support the bone resorption process during the pathogenesis of periodontitis.

The most unique finding of the current study is that the expression of BlyS was higher in subjects with periodontitis and DM than in subjects with periodontitis alone, while these groups did not differ significantly in terms of APRIL expression. The role of BlyS and APRIL have been well documented in the context of B cells, as both cytokines are described as key regulators of survival and/or expansion of B-cell subsets.³⁸ However, a previous study from

our group demonstrated a tendency toward low levels of B plasma cells in gingival tissues with periodontitis from type 2 diabetic subjects, when compared to those from subjects without DM.⁷ It is important to point out that BlyS and APRIL exert their functions by interacting with their receptors (BAFF-R), the transmembrane activator and cyclophilin ligand interactor (TACI) and the B-cell maturation antigen (BCMA), which are expressed not only in B-cell subsets but also in different types of immune cells including monocytes, dendritic cells and T cells. Therefore, evidence has shown that BlyS and APRIL are also related to T-cell differentiation and maintenance and to the production of pro-inflammatory cytokines.^{22,39,40} BlyS is particularly recognized as a significant co-stimulator of T-cell function, besides regulating multiple B-cell-related functions. Human T cells secrete pro-inflammatory cytokines, such as interferon (IFN)- γ and interleukin (IL)-2, under the stimulation of recombinant or endogenous BlyS.^{41,42} BlyS is also a critical cytokine for the development of glucose intolerance.⁴³

Noteworthy, besides the expression of BLyS, RANKL and TRAP were also overexpressed in patients with periodontitis and DM, when compared to those with periodontitis alone. Furthermore, the expression of BLyS correlated positively with the expression of RANKL in the diabetic subjects with periodontitis.

Previous study demonstrated that ~50% of T cells and ~90% of B cells in tissues with periodontitis express RANKL.^{17,19} Despite the evidence suggesting that B cells express more RANKL and support more osteoclastogenesis than T in periodontitis,^{17,19} further studies are needed to elucidate the influence of DM on the interaction among BLyS, B cells, T cells and RANKL expression. Our preliminary data are just able to suggest that increased expression of BLyS, RANKL and TRAP in tissues of type 2 diabetic subjects might contribute to the pathogenesis of periodontitis in these subjects. However, whether B cells and/or T cells are activated and maintained with increased BLyS expression under DM challenge needs further investigation.

The main strength of this study is to show for the first time a possible role of BLyS on the pathogenesis of DM-related periodontitis. The small sample size,

particularly in the control group, was anticipated to be a limitation of the present study; however, all genes studied differed in expression between control and periodontitis groups with statistical significance. Another limitation of this study was the lack of a group of patients with diabetes but without periodontitis. This group should be included in further studies in order to determine the influence of DM only on the expression of the studied molecules. Determining which cells are essential BLyS targets and whether, and to what extent, this cytokine really confer greater susceptibility to periodontal breakdown in diabetic patients require significant further investigation.

In conclusion, BLyS is overexpressed in periodontitis tissues of subjects with type 2 DM, which may be critical to the pathogenesis DM-related periodontitis.

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