




Expression of Toll-like receptors 2 and 4 and its association with matrix metalloproteinases in symptomatic and asymptomatic apical periodontitis

A. Fernández^{1,2} · A. M. Cárdenas^{1,3} · J. Astorga¹ · P. Veloso¹ · A. Alvarado² · P. Merino¹ · D. Pino² · D. Reyes-Court⁴ · M. Hernández^{1,5,6} 

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Abstract

To determine Toll-like receptors (TLR)2 and TLR4 expression levels and associate them with matrix metalloproteinases (MMPs) in asymptomatic apical periodontitis (AAP), symptomatic apical periodontitis (SAP), and healthy controls. Apical tissue/lesion samples were obtained from chronic AAP ($n = 35$) and SAP ($n = 29$), and healthy periodontal ligament (HPL, $n = 10$) with indication of tooth extraction, respectively. mRNA expression levels of TLR2, TLR4, MMP-1, MMP-2, MMP-8, and MMP-13 were determined by real-time reverse-transcription polymerase chain reaction. The data were analyzed with Kruskal-Wallis and Dunn's post hoc test ($p < 0.05$). The correlation coefficient was obtained using the Spearman correlation ($p < 0.05$). TLR2, MMP-1, MMP-2, and MMP-13 mRNA levels were the highest in SAP followed by AAP and controls ($p < 0.05$). TLR4 and MMP-8 were over expressed in AAP and SAP compared to HPL ($p < 0.05$). TLR2 positively correlated with TLR4, MMP-1, MMP-8, and MMP-13 in SAP ($p < 0.05$). TLR2 and TLR4 are overexpressed in apical lesions versus healthy periodontal ligament and correlate with collagenolytic MMPs. Particularly, TLR2 is overexpressed in SAP in association with MMP-1, MMP-8, and MMP-13. Our results suggest that the activation of TLR2 along with MMP overexpression might contribute to SAP clinical presentation and progression. TLRs, MMPs, and their interaction can explain the clinical presentations and evolution of apical periodontitis and might represent key targets for new diagnostic and treatment approaches.

Keywords Matrix metalloproteinases · Periapical periodontitis · Toll-like receptor 2 · Toll-like receptor 4

✉ M. Hernández

¹ Laboratory of Periodontal Biology, Faculty of Dentistry, Universidad de Chile, Santiago, Chile

² Faculty of Dentistry, Universidad Andres Bello, Santiago, Chile

³ Health sciences division, Faculty of Dentistry, Universidad Santo Tomás, Bucaramanga, Colombia

⁴ Department of Surgery, Faculty of Dentistry, Universidad de Chile, Santiago, Chile

⁵ Department of Oral Pathology and Medicine, Faculty of Dentistry, Universidad de Chile, Santiago, Chile

⁶ Dentistry Unit, Faculty of Health Sciences, Universidad Autónoma de Chile, Santiago, Chile

Introduction

Apical periodontitis (AP) is the result of the host's immune response to the persistent microbial infection of the dental pulp. Chronic forms of AP are characterized by the destruction of the peri-radicular tissues, particularly the alveolar bone, resulting in the formation of an osteolytic apical lesion. This response can be exacerbated and becomes symptomatic due to the imbalance between the endodontic infection and the host's immune response [1, 2], potentially representing an active form of the disease [3].

Toll-like receptors (TLRs) are central for pathogen recognition by innate immune cells during endodontic infection [4]. TLRs respond to pathogen-associated molecular patterns (PAMPs) and trigger intracellular signaling pathways that activate NF- κ B and induce the expression of pro-inflammatory genes. TLR2 and TLR4 classically recognize Gram-positive's

lipoteichoic acids and Gram-negative's lipopolysaccharide (LPS), respectively [5]. Despite their key roles, few clinical studies have explored TLRs in apical lesions, limiting to their immunolocalization. TLR2 has been identified in mononuclear plasmacytes and macrophage-like cells in refractory AP [4]. TLR2 and TLR4 showed differential expression in periapical granulomas and radicular cysts with or without sinus tract, suggesting their participation in the reactivation of periapical inflammation [6]. Accordingly, experimental studies propose that TLR2 and TLR4 might play a relevant role in the progression of AP based on their ability to stimulate receptor activator of nuclear factor kappa-B ligand (RANKL) synthesis in osteoblasts [7], to promote survival of osteoclasts [8], and to induce pro-inflammatory cytokines [9] and matrix metalloproteinases (MMPs) [10].

MMPs are zinc-dependent enzymes that cleave components of the extracellular matrix. Bone resorption centrally involves the activity of collagenases (MMP-1, MMP-8, and MMP-13) and gelatinases (MMP-2 and MMP-9) that cooperatively and sequentially degrade native type I collagen and gelatin. In addition, they modulate bioactive non-matrix mediators, regulating the host's inflammatory response [11]. MMP-1, MMP-2, MMP-8, and MMP-9 immunolocalized in AL and their expression was reduced during the healing phase in experimental animal models, supporting their role in the initiation and progression of AP [12]. Many reports propose an important role for MMP-9 in human apical periodontitis, but much less is known regarding other relevant MMPs [13–16].

Previous studies support the interaction between TLRs and MMPs. The activation of TLR2 and TLR4 combined with the stimulation of mechanical force induced the secretion of MMPs by human periodontal fibroblasts via p38, JNK, and NF- κ B, and TLR2 modulated MMP-9 expression in experimentally induced AP [10, 17, 18]. Up to now, there are no studies assessing whether TLR expression and MMPs can influence the clinical presentation of AP. We aimed to determine TLR2 and TLR4 expression levels and associate them with MMPs in asymptomatic apical periodontitis (AAP), symptomatic apical periodontitis (SAP), and healthy controls. The null hypothesis is that TLR2 and TLR4 expression is not associated with MMPs or clinical symptoms in endodontic apical lesions.

Materials and methods

Study design

Cross-sectional study. The study was approved by the Ethics Committee of the Faculty of Dentistry, Universidad de Chile (no. 2016/08) and the Ethics-Scientific Committee of the Central Metropolitan Health Service (no. 2017/70). All procedures performed in studies involving human participants were

in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. Informed consent was obtained from all individual participants included in the study.

Patients and sample collection

Individuals consulting at the clinic of surgery, Faculty of Dentistry, Universidad Andrés Bello and the Public urgency hospital (HUAP), Santiago, Chile, were included if they had clinical and radiographic diagnosis of untreated AAP or SAP, and healthy periodontal ligament (HPL) from teeth having indication of extraction. The inclusion criteria were individuals ≥ 18 years, having teeth with clinical diagnosis of apical periodontitis in association with dental caries that did not respond to pulp sensitivity tests, and had a radiographic apical radiolucency ≥ 3 mm, according to previously defined criteria [3, 19]. Exclusion criteria were systemic diseases and/or antibiotic and/or anti-inflammatory consumption in the last 3 months prior to the study. SAP and AAP were diagnosed when clinical symptoms in response to percussion were present or absent, respectively, according to previously defined diagnostic terminology [20]. Apical lesions and healthy periodontal ligament were removed with sterile curettes and tweezers. Healthy periodontal ligament was extracted 2 mm below the amelocemental junction and 1 mm above the apex, washed with sterile 0.9% NaCl solution, and stored in 100 μ L of RNeasy (Qiagen, Valencia, CA, USA) at -80 °C until they were processed.

Isolation of RNA

Samples were homogenized and RNA extraction was performed using a commercial RNA commercial kit (Qiagen), following the protocol and recommendations of the manufacturer. RNA was quantified (260/280) using a spectrophotometer (Bio-Tek, Winooski, VT), and RNA integrity was evaluated in a microfluidic-based platform (Agilent 2100 Bioanalyzer; Agilent Technologies, Santa Clara, CA). Then, 1 μ g of RNA from each sample was converted to single-stranded cDNA, using a reverse transcription kit (Thermo Fisher Scientific, Carlsbad, CA).

Expression of RNA

The mRNA expression levels were determined through real-time reverse-transcription polymerase chain reaction (StepOnePlus®; Applied Biosystems, Singapore). 12.5 ng of cDNA were amplified using TLR 2, TLR4, MMP-1, MMP-2, MMP-8, and MMP-13 respective primer sets (Table 1) and KAPA SYBR® Fast qPCR Kits (KAPA Biosystems, Woburn, MA, USA). The amplification process consisted of

Table 1 Primers used in AAP, SAP, and HPL samples

Target	Sequence forward	Sequence reverse
18S	CTCAACACGGGAAACCTCAC	CGTCCACCACTAAGAACG
TLR2	CTCTCGGTGTCGGAATGTC	AGGATCAGCAGGAACAGAGC
TLR4	CCCTCCCCTGTACCCTTC	TCCCTGCCTTGAATACCTTC
MMP-1	GCTAACCTTTGATGCTATAACTACGA	GGATTTGTGCGCATGTAGA
MMP-2	TGTGTTCTTTGACAGGGAATG	AGGCTGGTCAGTGGCTTG
MMP-8	AGCAGCCAAAAGAGAACCA	CCCATTGGGTTTGGACTC
MMP-13	CCAGTCTCCGAGGAGAACA	AAAAACAGCTCCGCATCAAAAC

the following steps: 95 °C for 3 min, 40 cycles of 95 °C for 30 s, 95 °C for 15 s, 60 °C for 1 min, and 95 °C for 15 s.

Analysis of results

Relative quantification of gene expression levels of TLR2, TLR4, MMP-1, MMP-2, MMP-8, and MMP-13 in AAP, SAP, and HPL controls was analyzed by $2^{-\Delta\Delta CT}$ method. The normalization of gene expression was made through the 18S rRNA, and HPL was used as a fold change reference.

The Shapiro-Wilk test was used to determine the distribution of the data. Inferential analyses were performed with chi-square test, Mann-Whitney or Kruskal-Wallis, and Dunn's post hoc tests. The correlation coefficient between TLR and MMPs was obtained using the Spearman's correlation analysis. The level of significance was defined at $p < 0.05$. The statistical analysis was performed using STATA 12® (StataCorp LP, TX, USA). The figures were performed in GraphPad Prism 6 (GraphPad Software, Inc., San Diego, CA, USA). No formal sample-size calculation was performed due to the exploratory nature of this study.

Results

The study sample consisted of 74 individuals. Age, gender, tobacco smoking, and educational level for each diagnostic group are shown in Table 2.

Table 2 Demographic parameters and smoking habit of study individuals. AAP, asymptomatic apical periodontitis; SAP, symptomatic apical periodontitis; HPL, healthy periodontal ligament

Parameters	HPL	AAP	SAP	<i>p</i>
Age (years, median (IQR))	22 (13)	47 (20)	34 (18)	0.0001*
Females (<i>n</i> , %)	7 (63.6%)	20 (54.1%)	15 (44.8%)	0.533
Smokers (<i>n</i> , %)	3 (27.3%)	14 (37.8%)	13 (44.8%)	0.585
Educational level (median)	Full high school	Full high school	Full high school	0.051

* $p < 0.05$

TLR-mRNA expression levels in AAP, SAP, and control groups are presented in Fig. 1. The expression levels of TLR2 were the highest in SAP, followed by AAP and controls, with significant differences among the three groups ($p < 0.05$). Significantly higher expression levels of TLR4 mRNA were observed in SAP and AAP, compared to healthy controls ($p < 0.05$). No significant differences were found between SAP and AAP.

MMP mRNA expression levels in AAP, SAP, and control groups are presented in Fig. 2. MMP-1, MMP-2, and MMP-13 were the highest in SAP followed by AAP and controls, with statistically significant differences among the three groups ($p < 0.05$). MMP-8 was significantly higher in SAP and in AAP versus healthy controls ($p < 0.05$). No differences were found between SAP and AAP.

Table 3 shows the correlation matrix between TLR and MMP gene expression. In AAP, a positive correlation was found between TLR2 and TLR4. TLR2 positively correlated with MMP-13, whereas TLR4 positively correlated with MMP-8 and MMP-13 ($p < 0.05$). In SAP, positive correlations were found between TLR2 and TLR4. Additionally, positive correlations were identified between TLR2 and MMP-1, MMP-8, and MMP-13, whereas TLR4 positively correlated with MMP-8 and MMP-13 ($p < 0.05$).

Finally, we explored whether smoking influenced TLR and MMP. Table 4 shows their expression levels between non-smokers and smokers in AAP and SAP. Among them, MMP-2 was significantly lower in smokers compared to AAP ($p < 0.05$), and a similar tendency was found in SAP, but non-significant.

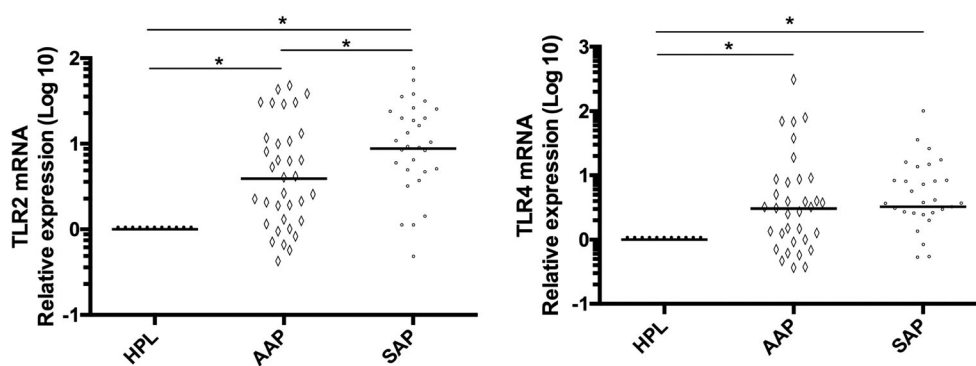


Fig 1 Relative gene expression of TLR2 and TLR4 in AAP, SAP, and HPL controls. Gene expression was normalized relative to 18S rRNA and expressed as the difference between $2^{-\Delta\Delta CT}$. Values on y-axis are shown in

log₁₀ scale. AAP, asymptomatic apical periodontitis; SAP, symptomatic apical periodontitis; HPL, healthy periodontal ligament. Bars represent the medians and asterisk $p < 0.05$

Discussion

The exacerbation of the clinical symptoms in SAP has been proposed to represent a progressive stage of apical lesions [3], which might be explained by immune activation in response to the bacterial challenge inside the root canal or apical lesion [21]. In the present study, we show that the symptomatic stage of AP-associated lesion may be conditioned by the participation of TLR in association with MMPs.

TLR2 and TLR4 have previously been identified in human apical tissues. Our results demonstrate that TLR2 and TLR4 were significantly over expressed in SAP and AAP versus HPL. However, only TLR2 expression levels were significantly higher in SAP compared to AAP. In line with our results, a previous study reported higher localization of

TLR2 in symptomatic than asymptomatic periapical granulomas or inflammatory periapical cyst in refractory lesions through immunohistochemical analysis [4]. A more recent study reported higher immunodetection of TLR2 and TLR4 in inflammatory cysts with sinus tracts compared to those with no sinus tracts [6], supporting a role for TLR in different clinical forms of acute exacerbation of AP. Our results also showed a positive correlation in the expression of TLR2 and TLR4 in both, AAP, and SAP. Experimentally induced AP in TLR2-knockout mice showed larger periapical lesions in association with a more pro-inflammatory phenotype induced by TLR4 over activation, suggesting a cooperative cross-talk between TLR2 and TLR4. In other study, TLR2 agonists up-regulated osteoclastic genes, resulting in higher osteoclast formation and bone loss in wild type mice and osteoblast cell

Fig 2 Relative gene expression of MMP-1, MMP-2, MMP-8, and MMP-13 in AAP, SAP, and HPL controls. Gene expression was normalized relative to 18S rRNA expressed as the difference between $2^{-\Delta\Delta CT}$. Values on y-axis are shown in log₁₀ scale. AAP, asymptomatic apical periodontitis; SAP, symptomatic apical periodontitis; HPL, healthy periodontal ligament. Bars represent the medians and asterisk $p < 0.05$

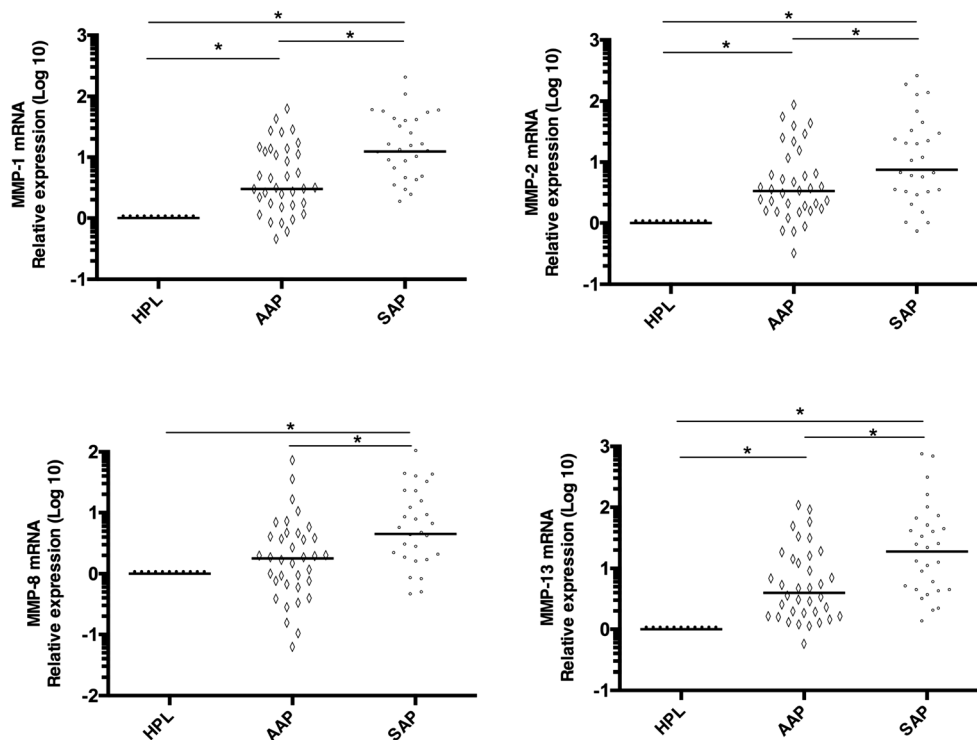


Table 3 Correlation matrix between TLR and MMP gene expression in AAP and SAP

	AAP		SAP	
	TLR2	TLR4	TLR2	TLR4
TLR2	1	–	1	–
TLR4	0.592*	1	0.569*	1
MMP-1	0.341	0.346	0.472*	0.264
MMP-2	0.275	0.080	0.242	0.332
MMP-8	0.164	0.439*	0.632*	0.524*
MMP-13	0.504*	0.469*	0.389*	0.455*

AAP, asymptomatic apical periodontitis; SAP, symptomatic apical periodontitis

Spearman's correlation coefficient (*r*)

**p* < 0.05

cultures, but not in TLR2-deficient mice [7], supporting that TLR2 also represents a key regulator of alveolar bone resorption [22, 23]. Overall, both TLR2 and TLR4 might interact in the settlement of the apical lesion during AAP, whereas the immunologically active stage in SAP could be triggered predominantly by TLR2 engagement.

A classic drawback from our study design is the young age of the control group, because orthodontics is a common indication particularly in young people. However, it has been previously validated in the literature based on the fact that healthy periodontal ligament represents the closest histophysiological counterpart for AP in humans [24, 25].

The interaction between periodontal inflammation and persistent bacterial infection upregulates MMPs, which contributes to the progressive breakdown of apical tissues. Collagenases (MMP-1, MMP-8, and MMP-13) process native collagen without unwinding the triple helical assembly of the molecule, generating 3/4 fragments, whereas gelatinases (MMP-2 and MMP-9) further process denatured collagen in a cooperative manner [11, 26, 27]. In this study, we showed that MMP-1, MMP-2 and MMP-13 were significantly over expressed in SAP followed by AAP and controls. We also found significantly higher levels of

MMP-8 mRNA in both AP forms compared to HPL. The results support that these MMPs participate in the pathogenesis of AP, but particularly MMP-1, MMP-2, and MMP-13 showed to be relevant in the active stage and/or clinical symptoms of apical chronic inflammatory lesions.

Clinical and experimental animal studies have also demonstrated high levels of MMP-2 in association with acute forms of AP. Previously, MMP-2 was detected in primary acute apical abscess and secondary apical abscess. Conversely, MMP2 forms were not detected in SAP with sinus tract [28]. Also, higher mRNA expression of MMP-2, MMP-7, and MMP-9 were reported in chronic apical abscesses compared to AAP [29], as well as higher MMP-2 and MMP-9 immunodetection in symptomatic apical lesions versus asymptomatic ones [16, 30]. All these antecedents in line with our results support a role for MMP-2 in the acute exacerbations and symptoms of chronic apical lesions. MMP-2 has been reported to have a central role in bone resorption in experimental models and has also been proposed to contribute to neuropathic pain symptoms during tissue injury [31, 32].

MMP-1 has been classically involved in physiologic matrix turnover [33], though a previous study reported higher MMP-1 mRNA in periapical lesions compared to healthy tissues [34]. Moreover, MMP-1 mRNA levels were over expressed in active lesions, defined as those having a higher RANKL/OPG ratio, along with the osteolytic cytokines, TNF- α , IFN- γ , IL-21, and IL-17A, in accordance with our results [34, 35].

Regarding MMP-8 and MMP-13, no previous reports are available in human symptomatic apical lesions. Still, our results are supported in part by previous reports from our research group. MMP-8 levels were elevated in gingival crevicular fluid (GCF) of teeth with AAP, SAP, and chronic marginal periodontitis compared to healthy controls [15, 36, 37]. MMP-13 on the other hand, was reported to increase in GCF from active periodontal lesions [38], whereas both MMP-8 and MMP-13, were involved in the development of apical lesions in experimentally induced AP [39]. They can also modulate bone resorption through osteoclast activation and differentiation, besides their classical role of periodontal tissue

Table 4 TLR and MMP gene expression between smokers and no smokers in AAP and SAP

	AAP		<i>p</i>	SAP		<i>p</i>
	NS (<i>n</i> = 23)	S (<i>n</i> = 13)		NS (<i>n</i> = 16)	S (<i>n</i> = 13)	
TLR2	3.25 (27.24)	3.91 (4.15)	0.83	7.80 (12.92)	11.96 (18.16)	0.38
TLR4	2.40 (7.58)	3.36 (3.00)	0.58	3.08 (7.425)	3.27 (4.30)	0.46
MMP-1	3.02 (12.06)	2.51 (4.39)	0.10	11.50 (33.08)	14.19 (26.66)	0.77
MMP-2	5.05 (18.68)	1.83 (1.44)	0.03*	18.22 (71.98)	5.50 (10.44)	0.35
MMP-8	1.61 (4.06)	1.93 (2.95)	0.41	7.67 (18.23)	3.21 (3.59)	0.29
MMP-13	3.44 (5.29)	5.29 (30.12)	0.66	25.41 (53.48)	7.73 (30.99)	0.26

AAP, asymptomatic apical periodontitis; SAP, symptomatic apical periodontitis; NS, no smokers; S, smokers. Results are expressed as medians and interquartile range (IQR). **p* < 0.05

breakdown. In vivo and in vitro studies support that MMP-13 is required for the differentiation of pre-osteoclasts, osteoclasts activation, and osteolysis. Several mechanisms have been reported, which include the activation of osteoclast-secreted pro-MMP-9, cleaving of galectin-3, a known inhibitor of osteoclastogenesis, and by regulating the RANKL/OPG axis, among others [11].

In the current study, moderate positive correlations were found for TLR2, TLR4, and the studied MMPs in SAP and to a lesser extent in AAP, suggesting that TLR signaling, among others, induces MMP expression in association with the clinical presentation of AP. Particularly, our results demonstrated positive correlations between TLR2, TLR4, and collagenolytic MMPs (MMP-1, MMP-8, and MMP-13) in SAP. It is likely to hypothesize that PAMPs derived from exacerbated bacterial infection and/or DAMP released in SAP induce TLR overexpression and activation, which in turn induce collagenolytic MMP synthesis, aggravating periapical tissue breakdown. Specially, MMP-8 and MMP-13 have been recognized as key MMPs associated with periodontal tissue destruction [11, 33]. In support of our proposal, TLR2 and also TLR4 have been reported to activate the synthesis of MMP-1, MMP-8, MMP-9, and MMP-13 via p38, JNK, and NF- κ B/AP-1 signaling in human fibroblasts and monocytic/macrophage cell lines [9, 18, 40, 41]; TLR2 activation and consequent recruitment of MyD88 was also reported to modulate the secretion of MMP-2 and MMP-9 in experimentally induced apical lesions [10]. MyD88 is an essential adaptor protein that has a key role in TLR signaling and triggers downstream cascades that activate NF- κ B/AP-1, which regulates the expression of MMPs [42–45].

It is well-defined that smoking impacts on the progression and outcome of marginal periodontitis [46]; however, this is not yet clear in apical periodontitis [47, 48]. In this regard, a previous study reported that the frequency and severity of apical periodontitis were not affected by the smoking habits of individuals [47]. Though smoking habit was similarly distributed between groups in the present study, we explored whether smoking might influence TLR or MMP expression within AAP and SAP groups. Only MMP-2 was significantly lower in smokers compared to non-smokers AAP ($p < 0.05$), and a similar tendency was found in SAP, but non-significant. Supporting our results, a previous study reported lower of MMP-2 expression levels in smokers than non-smokers in (marginal) periodontitis patients [49]. Nonetheless further studies designed to assess the influence of tobacco on the levels of key inflammatory markers are needed.

Conclusions

TLR2 and TLR4 are overexpressed in apical lesions versus healthy periodontal ligament and correlate with collagenolytic

MMPs. Particularly, TLR2 is overexpressed in SAP in association with MMP-1, MMP-8, and MMP-13. Our results suggest that the activation of TLR2 along with MMP overexpression might contribute to SAP clinical presentation and progression.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

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

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