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MIP-1α and MCP-1 as salivary biomarkers in periodontal disease



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KEYWORDS

Chemokines; Periodontal diseases; Saliva; Macrophage inflammatory protein-1; Monocyte chemoattractant protein-1 **Absract** *Aim:* Chemokines released by different host cells when exposed to the components of periodontopathic bacteria induce and maintain an inflammatory response in the periodontium. The aim of the study was to estimate the salivary levels of two chemokines, macrophage inflammatory protein-1 alpha (MIP-1 α) and monocyte chemo attractant protein-1 (MCP-1) in health, gingivitis and periodontitis and to evaluate their role as reliable salivary biomarkers in discriminating gingivitis and periodontitis from health.

Methods: A cross sectional study was designed to estimate the levels of MIP-1 α and MCP-1 in whole unstimulated saliva from 75 patients who were divided into healthy (Group 1, n = 25), gingivitis (Group 2, n = 25) and chronic generalized periodontitis (Group 3, n = 25). MIP-1 α and MCP-1 levels were estimated by using ELISA and were correlated with clinical parameters. ROC curve analysis was done to determine the sensitivity and specificity of these biomarkers in distinguishing periodontal disease from health.

Results: Both the biomarkers were detected in all the saliva samples. There was a statistically significant difference in the concentration of both the analytes in Group 3 and Group 2 compared with Group 1 (p < 0.001). ROC curve analysis showed 100% sensitivity and specificity for MIP-1 α and MCP-1 in discriminating periodontitis from health. For discriminating gingivitis from health, MIP-1 α had a higher sensitivity and specificity (100% & 88% respectively) compared to MCP-1 (84.1% & 80% respectively).

Conclusion: There is a substantial increase in the concentration of both MIP-1 α and MCP-1 with increasing severity of periodontal disease. Both the analytes showed promising results as biomarkers for discriminating periodontal disease from health.

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1. Introduction

The biochemical signalling involved in the periodontal disease inflammatory disease process consists of three biological phases i.e. inflammation, connective tissue degradation, and alveolar bone turnover. An intermediate mechanism that lies

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between bacterial stimulation and tissue destruction is the production of cytokines, which stimulates inflammatory events that play an important role in leukocyte recruitment and may directly or indirectly modulate osteoclast formation (Graves, 2008). A constellation of circulating molecules in these biological phases have been detected at elevated levels in the gingival crevicular fluid and whole saliva of patients who have periodontal disease (Craig and Loberg, 2006). Among the mediators potentially involved in leukocyte migration to periodontal environment, chemokines have been investigated with special interest (Silva et al., 2007).

Chemokines are class of chemotactic cytokines that are capable of activating and promoting the migration of a variety of leukocytes and bone cells. They induce leukocyte migration and activation by binding to specific G protein coupled seven transmembrane spanning cell surface receptors (Graves and Jiang, 1995). Chemokines are found in gingival tissue and crevicular fluid and they are produced by number of cell types in the periodontium such as fibroblasts, endothelial cells, macrophages, osteoclasts, epithelial cells, polymorphonuclear leukocytes, monocytes, lymphocytes and mast cells. Some chemokines have important pro-inflammatory effects and are related to periodontal tissue destruction that involves the stimulation of bone resorption and induction of tissue damage (Hanazawa et al., 1993).

MIP-1 α and MCP-1are chemotactic chemokines secreted by a variety of cell types, including macrophages, fibroblasts, epithelial cells, and endothelial cells (Furutani et al., 1989, Gong et al., 1997, Pype et al., 1999). These play an important role in the pathogenesis of various inflammatory diseases and conditions such as periodontitis, multiple myeloma, Sjögren syndrome, and rheumatoid arthritis (Maurer and von Stebut, 2004). They perform various biological functions, such as recruiting inflammatory cells, wound healing, inhibition of stem cells, and maintaining effector immune response (Gemmell, 2001).

Early detection of disease plays a crucial role for treatment planning and prognosis. Saliva has great potential as a diagnostic fluid and offers advantage over serum and other biological fluids by its non-invasive method of collection, smaller sample aliquots, cost effectiveness, early storage and transportation (Giannobile et al., 2011, Lamster et al., 1990). Salivary biomarkers are potentially important for determining the presence, risk and progress of periodontal disease. However, clinical translation of biomarker technology from lab to chair side requires studies that identify bio-markers associated with health, gingivitis and periodontitis (Syndergaard et al., 2014). Unlike periodontal disease, gingivitis has lacked evidence of salivary biomarkers with strong discriminatory capacity, a feature critical for the translational utility of salivary biomarkers to chair side diagnostics (Ebersole et al., 2015). Because of the relationship of gingivitis with periodontitis and the unidentified biologic processes that lead to the transition between these two inflammatory conditions, diagnostic tools for early detection are desirable (Syndergaard et al., 2014). Thus, the present study aimed to estimate the levels of salivary MIP-1a and MCP-1 in healthy individuals, patients with gingivitis and chronic periodontitis and to correlate their levels with the clinical parameters of periodontal disease. Also, the study aimed to evaluate the discriminatory capacity of MIP-1a and MCP-1 to distinguish periodontal disease from health.

2.1. Study population

This cross-sectional study was approved by Institutional Ethical Committee, Vydehi Institute of Dental Sciences and Research Centre, Bangalore, India. Patients aged between 18 and 50 years, who reported to the outpatient unit of Department of Periodontics, between September 2015 and June 2016 and were willing to participate in study and give informed consent were considered for the study. Patients using tobacco and tobacco related products or alcohol, patients having any systemic diseases, pregnant and lactating women, those who underwent periodontal treatment in the last 6 months, patients who have taken antibiotics or anti-inflammatory drugs within one month prior to the study, patients who had any oral mucosal inflammatory conditions, xerostomia or salivary gland disorders and those who were having < 20 teeth were excluded from the study. 75 adults (40 males and 35 females) conforming to the inclusion criteria were included in the study. Plaque Index (PI) (Loe, 1967), Gingival Index (GI) (Loe, 1967), presence or absence of bleeding on probing (BOP) (Lang et al., 1986), probing pocket depth (PPD) and clinical attachment loss (CAL) (Ramfjord, 1974) were recorded using a mouth mirror and UNC 15 probe.¹

Based on the clinical findings, patients were categorized into three groups. Group 1 consisted of 25 individuals with clinically healthy periodontium, with Mean GI \leq 1, BOP <20% of sites, no sites with PPD \geq 3 mm and no CAL, Group 2 consisted of 25 individuals who showed clinical signs of gingival inflammation, Mean GI > 1, BOP > 20% of sites, no sites with PPD \geq 3 mm and CAL \geq 2 mm. Group 3 included patients who were diagnosed with chronic periodontitis. Diagnosis of chronic periodontitis was made according to the criteria given by 1999 International Workshop for classification of Periodontal Diseases and Conditions (Armitage, 1999). 25 patients who showed signs of clinical inflammation, mean GI > 1, CAL > 3 mm and PPD \geq 5 mm in at least 30% of sites were included in the study.

2.2. Saliva sample collection

Unstimulated whole saliva was collected according to a previously described method (Miller et al., 2006). Individuals were recalled the following day for saliva sample collection to avoid contamination by blood during periodontal examination. 5 ml of unstimulated whole expectorated saliva was collected from each patient between 9 and 11 am while seated in upright position. Patients were asked to avoid all oral hygiene measures, eating, drinking, or gum chewing for 1 h before collection. Patients rinsed their mouth with water, and then expectorated whole saliva into sterile tubes. Collected samples were stored at -80 °C prior to analysis.

2.3. MIP-1a and MCP-1 assay

The levels of MIP-1 α and MCP-1 in saliva were estimated using an enzyme-linked immunoassay kit.² All analyses were

¹ Hu-Friedy PCP UNC 15 Qulix, Chicago, IL.

² EMD Millipore, Billerica, MA.

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Category	Group 1 ($n = 25$)	Group 2 ($n = 25$)	Group 3 (n = 25)	P value			
Age in years	* 24.8 ± 3.1	30 ± 7.6	46.2 ± 9.9	0.000^{\dagger}			
Gender (M/F	F) 11/14	12/13	17/8	0.190 [‡]			
PI*	$0.44~\pm~0.48$	0.67 ± 0.44	1.87 ± 0.38	0.000^{\dagger}			
GI [*]	0.26 ± 0.23	1.37 ± 0.27	1.72 ± 0.35	0.000^{\dagger}			
% BOP*	12.04 ± 3.76	25.48 ± 3.42	31.64 ± 3.8	0.000^{\dagger}			
PPD^*	1.58 ± 0.58	2.37 ± 0.25	5.5 ± 0.66	0.000^{\dagger}			
CAL*	-	$1.43~\pm~0.30$	3.46 ± 0.34	0.000^{\dagger}			

 Table 1
 Descriptive statistics of demographics and clinical parameters of the study population.

* Mean \pm Standard Deviation.

[†] Significant P value using One-way ANOVA test.

[‡] Non-significant P value using Chi-square test.

Table 2 Scheffe's multiple comparison test for MIP-1 α and MCP-1 and the ANCOVA test result after controlling age of the subjects.

Comparison		Mean difference	P-value	ANCOVA test result		
Mean	Mean			Source	F-value	P-value
MIP-1α						
Group 1	Group 2	-11.3320*	.000	Age	0.315	0.576
	Group 3	-28.1440^{*}	.000	Group	70.207	0.000^{*}
Group 2	Group 3	-16.8120^{*}	.000			
MCP-1						
Group 1	Group 2	-176.1320^{*}	.000	Age	1.623	0.207
	Group 3	-550.1240^{*}	.000	Group	75.321	0.000^{*}
Group 2	Group 3	-373.9920^{*}	.000	·		
*						

^{*} P value statistically significant.

performed in duplicate within six months of sample procurement. Standards were included on all runs and all results were reported within the linearity of the assays. Readings were made at 450 nm with a micro plate spectrophotometer.³

2.4. Statistical analyses

All data were analysed using SPSS (version 18.0, Chicago, IL, USA). Descriptive and inferential statistical analysis has been carried out in the present study. Results on continuous measurements are presented as Mean \pm SD and results on categorical measurements are presented in Number (%). P <0.05 was considered statistically significant. Analysis of Variance (ANOVA) was carried out to find the significance of study parameters on categorical scale between two or more groups. Scheffe's multiple comparison test was applied to know the difference between the groups. Since the three groups had a statistically significant difference with respect to age, to rule out the possibilities of age acting as a confounding factor on the levels of the chemokines, ANCOVA test was done. Also, Bonferroni multiple comparison test was applied to compare the adjusted mean values. Pearson correlation co-efficient test was used to find the degree of correlation between the clinical parameters and salivary MIP-1a and MCP-1 concentration in all the three groups. Partial correlation coefficient has been calculated after the controlling the effect of the age of the subjects on the levels of the biomarkers studied. ROC curve analysis was done to determine the sensitivity and specificity of these biomarkers in distinguishing periodontal disease from health.

3. Results

Descriptive statistics of the study population is shown in Table 1. The significant P value for age between the groups revealed that all the three groups have been statistically different. In order to get more details of the variation between the three groups, repeated contrast test was applied. The result indicated the chronic periodontitis patients belonged to a higher age range when compared to individuals in other two groups. All three groups have been similar with respect to the gender of the individuals.

The clinical parameters used to differentiate the three groups (gingival index, bleeding on probing, probing pocket depth and clinical attachment level) were found to be significantly higher in group 3 relative to group 1 and 2 therefore conforming to the inclusion criteria (Table 1).

3.1. Salivary analyte levels

Scheffe's multiple comparison test showed that the mean values of MIP-1 α and MCP-1 levels in saliva was higher in Group 3 when compared to Group 1 and 2 (Table 2). The difference in levels of the analytes was found to be statistically significant. Since age can act as a confounding factor, ANCOVA test was applied to see whether the difference in

³ SpectraMAX 340, Molecular Devices, Sunnyvale, CA, USA.

the mean level of the chemokines has occurred due to periodontal inflammation or due to age of the subjects. The non-significant P value for age of the subjects revealed that age has no specific role in the difference of mean levels of MIP-1 α and MCP-1 and the difference in the mean values of the groups are due to the clinical conditions of the subjects only (Table 2). Further, Bonferroni multiple comparison test has been applied to compare the adjusted mean values. The result indicated that all the three comparisons were statistically different (Table 3).

3.2. Relationship between salivary analyte levels and clinical parameters of periodontal disease

Pearson correlation coefficient value between the clinical parameters and MIP-1 α level and MCP-1 level demonstrated a significant positive correlation (Table 4). Also, Partial correlation coefficient has been calculated after controlling the effect of the age of the individuals. All the partial correlation coefficient values were lesser than their corresponding Pearson correlation coefficient values, however, the significant P value shows that these variables correlate with MIP-1 α and MCP-1 values even after controlling the effect of age of the individuals (Table 4).

Figs. 1 and 2 shows the results of receiver operating characteristic (ROC) analysis which was done to find out whether these two chemokines could be used as biomarkers for gingivitis and periodontitis and which among the two is a better predictor for disease progression. The P value was found to be highly significant for both MIP-1 α and MCP-1 in gingivitis

4. Discussion

This cross-sectional study evaluated the levels of two chemokines, MIP-1a and MCP-1 in whole unstimulated saliva of 75 adults who were categorized into three groups- healthy controls, patients having gingivitis and chronic generalized periodontitis. At the time of designing this study, literature search did not reveal any studies comparing healthy, gingivitis and periodontitis patients for the two biomarkers used, and hence the third group of gingivitis was included. Ebersole et al. in 2015, conducted a similar study for evaluating IL-1 β , IL-16, MIP-1 α , and MMP-8 in whole saliva in periodontal health, gingivitis and periodontitis. The results of this study had a close resemblance to the present study in that the levels of MIP-1 α was significantly increased in the periodontitis group compared to both gingivitis and healthy groups. Few other studies have also reported similar results for MIP-1 α in gingivitis (Syndergaard et al., 2014) and periodontitis (Al Sabbagh et al., 2012). In contrary to all the previously reported studies, Emingil et al. in 2005 reported that generalized aggressive periodontitis, chronic periodontitis, gingivitis and healthy groups had comparable GCF MIP-1a levels and concluded that the MIP-1a levels in GCF do not seem to play a discriminatory role in periodontitis. They explained that the low MIP- 1α levels in periodontitis group could be because of the lack of macrophages as well as subsets lymphocytes with specific receptors for MIP-1a.

	Estimated marginal value		95% Confidence value		Bonferroni multiple comparison test result			
	Mean	SE*	Lower	Upper	Comparison	n	Mean difference	P value
MIP-1a								
Group 1	7.89	1.27	5.35	10.44				
Group 2	18.99	1.08	16.82	21.15	Group 1	Group 2	-11.091	.000
Group 3	35.05	1.47	32.11	37.98	_	Group 3	-27.151	.000
_					Group 2	Group 3	-16.061	.000
MCP-1								
Group 1	141.57	27.01	87.69	195.45				
Group 2	329.30	23.03	283.38	375.22	Group 1	Group 2	-187.728^{*}	.000
Group 3	739.42	31.15	677.30	801.53		Group 3	-597.848^{*}	.000
,					Group 2	Group 3	-410.119^{*}	.000

* Standard error.

Table 4 Correlation coefficient value between the clinical parameters and MIP-1a level and MCP-1 level.

Clinical parameters	MIP-1a			MCP-1	MCP-1		
	r-Value	r-Value*	p-Value	r-Value	r-Value*	p-Value	
PI	0.769	0.555	< 0.001	0.706	0.470	< 0.001	
GI	0.819	0.694	< 0.001	0.703	0.507	< 0.001	
BOP	0.792	0.624	< 0.001	0.750	0.569	< 0.001	
PPD	0.880	0.739	< 0.001	0.859	0.730	< 0.001	
CAL	0.879	0.730	< 0.001	0.862	0.737	< 0.001	

* Partial correlation coefficient value after adjusting for age.



Fig. 1 Receiver operating characteristic (ROC) analysis showing MIP-1a in (A) gingivitis and (B) periodontitis.



Fig. 2 Receiver operating characteristic (ROC) analysis showing MCP-1 in (A) gingivitis and (B) periodontitis.

The salivary concentrations of MCP-1was found to be higher in group 3 when compared to group 1 and 2. Of the available literature, the only report on the levels of MCP-1 in saliva was the study done by Gupta et al. (2013). The results of the present study are in accordance with this study, which compared the MCP-1 levels in saliva, GCF and serum in individuals with chronic periodontitis. Same authors published a study in 2015 in which the MCP-1 levels were correlated with soluble CD-40 ligand, a cardiac marker in both GCF and serum. Correlation studies revealed that both these markers demonstrated a strong correlation among each other both in GCF and serum before and after periodontal therapy. Various other recent studies also reported higher GCF levels of MCP-1 in periodontitis patients when compared to healthy individuals (Hanazawa et al., 1993, Kurtis et al., 2005, Pradeep et al., 2009a, 2009b). Comparable results were also obtained with MCP-1 levels in serum and gingival tissues also (Gemmell et al., 2001, Tonetti et al., 1994). Tonetti et al. demonstrated mRNA for MCP-1 in tissue biopsies taken from diseased periodontal sites. The presence of MCP-1 was detected in the basal

layers of oral epithelium as well as in the inflammatory infiltrate in the diseased periodontal tissues.

The present study showed a positive statistical correlation between the levels of MIP-1a and MCP-1 with all the clinical parameters evaluated. Other studies which evaluated these two biomarkers in saliva, GCF or serum have shown similar correlation with clinical parameters of periodontal disease (Emingil et al., 2005, Pradeep et al., 2009b, Sexton et al., 2011). The elevated levels of MIP-1a and MCP-1 and its statistically significant positive correlation with clinical parameters suggests their role in the pathogenesis of inflammatory periodontal disease. MIP-1 α is the most abundantly expressed chemokine in periodontitis tissues, with its expression localized in the connective tissue subjacent to the pocket epithelium of inflamed gingival tissues (De Lima et al., 2016). Both MIP-1a and MCP-1 are involved in the migration of macrophages to periodontal tissues (Gemmell et al., 2001). Products derived from macrophages such as IL-1 and TNF- α are pro-inflammatory cytokines which are known to induce bone resorption. Thus, these chemoattractants play a significant role in maintenance of chronic inflammatory reaction and the bone loss characteristic of periodontal disease.

Numerous studies have been conducted across the globe to identify biomarkers for distinguishing the individuals with periodontal disease from periodontally healthy individuals. But clinical relevance of these studies can be considered only if the sensitivity and specificity of the biomarker to discriminate the appropriate disease are evaluated (Fine et al., 2009). The results of our present study revealed that MIP-1a was the better biomarker in terms of sensitivity and specificity in differentiating gingivitis from health. Whereas in terms of differentiating periodontitis, both MIP-1a and MCP-1 had 100% sensitivity and specificity. Fine et al. in 2009 reported a sensitivity of 90.3% and specificity of 85.7% for salivary MIP-1a for the diagnosis of aggressive periodontitis. Later, the same authors reported the sensitivity and specificity of MIP-1 α in GCF to be 93.33% and 98.73% respectively (Fine et al., 2014). The systematic review and meta-analysis published by De Lima et al. in 2016, which reviewed the clinical utility of various salivary biomarkers (MMP 8, MMP 9, IL-1β, IL-6, IL-10, PGE2, OPG, TNF α , MIP-1 α , calprotectin, albumin, ICTP), identified that only MIP-1 α had a high accuracy of identifying or excluding the occurrence of periodontal disease in adults.

5. Conclusion

The results of the present study suggest that salivary levels of both MIP-1 α and MCP-1 could have clinical utility as a screening tool for discriminating the various stages of periodontal disease. Future longitudinal studies are recommended to determine whether these two biomarkers could be of help in predicting periodontal breakdown in susceptible individuals.

6. Conflict of interest and source of funding statement

The authors declare that they do not have a conflict of interest. This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

7. Ethical statement

This cross-sectional study was approved by Institutional Ethical Committee, Vydehi Institute of Dental Sciences and Research Center, Bangalore, India. Informed consent was obtained from all participants prior to start of the study.

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