

Host-derived diagnostic markers related to soft tissue destruction and bone degradation in periodontitis

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Buduneli N, Kinane DF: Host-derived diagnostic markers related to soft tissue destruction and bone degradation in periodontitis. J Clin Periodontol 2011; 38 (Suppl. 11): 85–105. doi: 10.1111/j.1600-051X.2010.01670.x.

Abstract

Background: A major challenge in clinical periodontics is to find a reliable molecular marker of periodontal tissue destruction with high sensitivity, specificity and utility.

Objectives: The aim of this systematic review is to evaluate available literature on 'the utility of molecular markers of soft and hard periodontal tissue destruction'.

Materials and Methods: Based on the focused question, 'What is the utility of molecular markers of soft and hard periodontal tissue destruction', an electronic and manual search was conducted for human studies presenting clinical data for the potential of molecular markers of tissue destruction in biofluids; gingival crevicular fluid (GCF), saliva, and serum.

Results: Papers fulfilling the inclusion criteria were selected. All relevant data from the selected papers were extracted and recorded in separate tables for molecules in GCF, saliva, and serum.

Conclusion: Within the defined limits of the Problem/Population, Intervention, Comparison, Outcome, the present analysis reveals that (a) no single or combination of markers exists that can disclose periodontal tissue destruction adequately; (b) while the most fruitful source of biomarkers for periodontal destruction appears to be in molecules tightly related to bone and soft tissue destruction, this remains to be objectively demonstrated. Currently, clinical measurements are still the most reliable.

Key words: biomarkers; diagnosis; gingival crevicular fluid; periodontal disease; saliva; serum

Accepted for publication 7 November 2010

Periodontitis is one of the most common oral diseases and is characterized by gingival inflammation and alveolar bone resorption (Savage et al. 2009). More than 500 different bacterial spe-

cies are able to colonize the oral biofilm and up to 150 different species of bacteria are possible in any individual's subgingival plaque. According to a report by the World Health Organisation, severe periodontitis leading to tooth loss was found in 5–15% of most populations worldwide (Armitage 2004). Hence, it can be considered among the prevalent and important global health problems in terms of quality of life.

A diagnostic tool should provide pertinent information to aid differential diagnosis, screening, presence, location, severity or staging and prognosis of a disease. At present, periodontitis is diagnosed almost entirely on the basis of an

array of clinical measurements including probing depth (PD), clinical attachment level (CAL), bleeding on probing (BOP), plaque index (PI) recordings and radiographical findings. However, PD and CAL measurements by periodontal probes and radiographic bone levels, provide information about past periodontal tissue destruction and do not elucidate the current state of the disease activity nor predict the future. Therefore, one of the major challenges in the field of periodontology is to discover a method of predicting the future of periodontal disease or at least to declare the current state of disease activity. Thus, objective and ideal diagnostic methods

Conflict of interest and source of funding statement

The authors declare that they have no conflict of interests.

This work was supported in part by Ege University (NB) and by the United States Public Health Service National Institute of Health, NIDCR Grant DE017384 to D. F. K. This supplement was supported by an unrestricted grant from Colgate.

for periodontal diagnosis are still being sought. The ideal periodontal diagnostic method should be able firstly to screen susceptible subjects in the general population, secondly to differentiate active and inactive sites, thirdly to predict future tissue destruction in particular individuals and sites, and finally to monitor the response to periodontal therapy.

Periodontitis is described as a multifactorial irreversible and cumulative condition, initiated and propagated by bacteria and host factors (Kinane 2001). Given the complex nature of periodontitis, it is unlikely that one single clinical or laboratory examination can address all issues concerning diagnosis, classification, and prognosis (Van der Velden 2005, Xiang et al. 2010). Moreover, molecular markers of bone resorption have advantages as well as disadvantages as they relate to specificity for bone, ease of detection, pre-analytic stability, and availability of sensitive and specific assays for detection (Koka et al. 2006). Proteins derived from inflamed host tissue and pathogenic bacteria have the potential of being used as markers of periodontitis (Özmeriç 2004).

Numerous molecules in the oral fluids namely; gingival crevicular fluid (GCF) and saliva, as well as molecules in the blood circulation; serum or plasma have been investigated so far in an attempt to provide a sensitive and specific marker for periodontal tissue destruction. GCF and saliva are particularly promising as they can easily be obtained non-invasively and with minimal discomfort to the patient and consist of both locally synthesized and systemically derived molecules. Thus, evaluation of various biologically specific proteins or markers in oral fluids by using immunologic or biochemical methods may provide information on the events going on in the periodontal micro-environment (Kinney et al. 2007).

The immune and inflammatory responses are critical to understanding the pathogenesis of periodontal diseases and they are orchestrated by a number of host-related factors, either intrinsic or induced (Taubman et al. 2005). Under normal physiologic conditions, there is a balance between bone formation and bone resorption. Bone homeostasis is maintained as long as this balance is preserved. When the structural integrity and/or calcium metabolism is altered, this balance is lost towards either increased bone formation or towards bone resorption. Inflammatory conditions like periodontal diseases and rheumatoid

arthritis, and metabolic conditions like osteoporosis are examples of an altered balance between bone formation and bone resorption.

The threshold for periodontitis progression is based upon extensively documented evidence within the periodontal literature. Many epidemiological studies apply a threshold for loss of attachment of changes of 3 mm as the definition of disease progression. The threshold is set at the level of two teeth to minimize the risk of including cases of progression arising because of reasons other than periodontitis and/or measurement error. Presence of ≥ 2 teeth demonstrating a longitudinal loss of proximal attachment of ≥ 3 mm has been proposed as the criteria for case definition of periodontitis progression (Tonetti & Claffey 2005). In situations where serial proximal attachment level measurements are not available, longitudinal radiographic bone loss of ≥ 2 mm at ≥ 2 teeth may be used as a substitute.

The criteria for definition of progressing sites are as stated by the working group in the Fifth European Workshop in Periodontology, in 2005 as follows:

- The exposure of interest, along with relevant covariates including age, is used in a multi-factorial model, using the "case" as the dependent variable. Adjusted odds ratios and/or relative risk estimates as well as 95% confidence limits should be reported.
- The impact of the putative risk factor on the extent of periodontitis progression (percentage of teeth affected by disease progression) is examined to investigate a dose-response effect, using a new multivariate model.
- Additional models may be developed as needed to test specific hypotheses.

Clinical examination and the clinical periodontal measurements can provide information on the disease activity, only if these measurements are repeated at two time-points. However, knowing the disease activity state might be critical to clinical decision making at one single time point and clinical evaluation cannot fulfil this requirement. This deficiency has spawned dozens of studies with as yet no definitive answer.

There are also well-accepted risk factors such as diabetes and smoking

to be considered when determining the diagnosis as well as prognosis of a particular case. Cigarette smoking represents a risk factor for progression of periodontitis, the effect of which may be dose related. Indeed, apart from the plaque biofilm, it is the major environmental risk factor. Heavy smokers should be considered as high-risk individuals for disease progression. The clinical implications for this are that smokers should be identified during patient examination and efforts should be made to modify this behavioural risk factor. Furthermore, smoking or molecules related to smoking such as blood cotinine induced by smoking should be considered as important risk markers of periodontal disease that are relevant to the assessment of prognosis (Calsina et al. 2002, Tang et al. 2009).

The aim of the present review was to provide a systematic review of the state of evidence on whether current periodontal tissue destruction can be disclosed accurately utilizing chemical analysis of various molecules in the biological fluids.

Materials and Methods

Search strategy

A literature search of the last thirty years was performed using the ISI and PubMed database from 1980 to 15 June 2010, with the following search strategy: ("periodontitis" OR "periodontal disease") AND ("progression" OR "activity") AND ("saliva" OR "gingival crevicular fluid" OR "serum" OR "plasma"). The search was limited to the English language. In vitro studies on cell cultures, experimental studies on animal models, polymorphism studies, studies particularly investigating possible role of various therapeutic agents such as subantimicrobial dose doxycycline, anti-inflammatory agents, or dietary supplements, as well as studies comprising only patients with particular systemic diseases such as diabetes or rheumatoid arthritis and studies focused only on smoking were excluded from the present review. Titles and abstracts were screened and full text of publications was obtained for the selected articles. All levels of evidence were included. In addition, the reference lists of review papers were hand searched. To be eligible for inclusion in this systematic review, studies had to meet the following criteria: (1) original

investigations; (2) studies conducted within a human population; (3) studies having a systemically healthy chronic periodontitis group; (4) studies correlating the biochemical findings in biofluids with clinical periodontal parameters: PD, CAL, and BOP; (5) preferentially follow-up and/or intervention studies, but cross-sectional studies were also included in the review.

Outcome variables

The primary outcome variables investigated were PD, CAL, and incidence of BOP.

Results

Clinical methods for periodontal diagnosis

Measuring PD and CAL by a calibrated periodontal probe and assessing gingival inflammation by gingival index (GI) (Löe & Silness 1963) and/or a bleeding index such as BOP within a certain time after probing or papilla bleeding index (PBI) (Saxer & Mühlemann 1975) are still the most reliable clinical periodontal parameters for periodontal diagnosis. Furthermore, PI (Silness & Löe 1964) provides information on the major local aetiological factor; microbial dental plaque. Presence of BOP is still the most reliable clinical finding indicating periodontal disease activity. However, the absence of bleeding is much more valuable in terms of being a highly specific negative predictor of periodontal disease activity (Lang et al. 1986, 1990). This helps us exclude healthy patients but of course does not help us truly focus on the really at risk subjects as if we were to address all patients with BOP we would be overwhelmed with false positives as this approach would be far too sensitive to be useful as a screening technique. Sensitivity is the ability of a diagnostic test to identify the target molecule when this is truly present. Specificity is the probability of a diagnostic test being negative when the target molecule is truly absent. Misleading test results occur when the test is positive and the disease is absent (false positive) or when the test is negative and the disease is present (false negative). An ideal diagnostic test should have sensitivity and specificity values approaching 100%. But unfortunately this is never the case.

Source for samples of the different diagnostic tests

GCF components

GCF is a transudate originating from the gingival plexus of blood vessels in the gingival connective tissue, close to the epithelium lining of the dentogingival space. It also collects resident host cells and microorganisms in the microbial dental plaque as well as their cellular products. GCF provides an accurate representation of tissue and serum concentrations of inflammatory mediators (Giannobile 1997, Champagne et al. 2003, Armitage 2004). Up to now, at least 90 different components in GCF have been evaluated as possible biomarkers for diagnosis of periodontal disease (Loos & Tjoa 2005). These markers can be divided into three major groups: host-derived enzymes and their inhibitors, inflammatory mediators and host-response modifiers, and byproducts of tissue breakdown (Lamster & Ahlo 2007).

A substantial number of studies throughout the 1980s and the 1990s explored the predictive ability of GCF components for identification of progressive periodontal lesions (Heitz-Mayfield 2005). While individual GCF components produced positive predictive values that were superior to individual clinical measures (Chapple et al. 1999), these studies focused largely on the prediction of periodontitis at the site level rather than the identification of high-risk groups and individuals. Data from the selected studies investigating potential biomarkers in GCF in relation to periodontal disease are outlined in Table 1. An early multi-centre study by Lamster et al. (1995) did examine the predictive value of β -glucuronidase (β G) at the patient level, in a population of predominantly recall patients, and demonstrated that subjects with persistently elevated levels of GCF β G at baseline, 2-week and 3-month recalls had between 7 and 14 times (dependent upon the algorithm used) increased risk ratio for periodontitis progression.

Salivary components

Saliva is a mirror of the body that can be used to monitor the systemic as well as the oral health status. Whole saliva contains constituents from exocrine glands secreting into the oral cavity, GCF and dietary and oral plaque components. Saliva is readily available and

easily collected without specialized equipment or personnel. Several mediators of chronic inflammation and tissue destruction have been detected in whole saliva of periodontitis patients as described in relatively recent reviews (Kaufman & Lamster 2000, Kinane & Chestnutt 2000, Lamster et al. 2003). In addition, as whole saliva represents a pooled sample with contributions from all periodontal sites, analysis of biomarkers in saliva may provide an overall assessment of disease status as opposed to site-specific GCF analysis (Miller et al. 2006).

Recently, the use of whole saliva as a means of evaluating host-derived products (e.g. salivary gland product, gingival crevice fluid, host enzymes) as well as exogenous components (e.g. oral microorganisms and microbial products) has been suggested as a potential diagnostic marker for disease susceptibility (Şahingür & Cohen 2004). The source of saliva, types of saliva samples, as well as saliva collection methods has been described in detail by Şahingür & Cohen (2004). Development of tests based on the detection of neutrophil defects, genetic markers or the detection and measurement of antibodies specific for periodontal pathogens may be useful in the future. However, there is currently insufficient evidence available for the predictive value of diagnostic tests assessing the host's susceptibility to future periodontitis progression.

Salivary enzymes originate from three major sources; the actual salivary secretions, the host cells found in GCF, and finally disposed bacterial cells from dental plaque and mucosal surfaces. Data from the selected studies evaluating the potential utility of salivary components as diagnostic markers for periodontal tissue destruction are presented in Table 2.

Blood components

Various studies have evaluated the molecular markers of tissue destruction in serum or plasma: these manifestations of periodontal diseases are mainly sought to clarify the possible interactions between periodontitis and various systemic diseases and/conditions such as cardiovascular diseases (CVDs), pregnancy complications, diabetes mellitus, and rheumatoid arthritis. Serum or plasma provides information about the inflammatory stimulus and/or response generated in circulation towards the

Table 1. Studies evaluating possible biomarkers in gingival crevicular fluid (GCF) samples

Reference	Study design	Study groups	Follow-up period	Clinical parameter	Biological sample	Biological parameter	Results
Akalun et al. (2005)	Cross-sectional	26 CP, 18 controls		Routine clinical parameters	GCF, biopsy	SOD	Increased in gingiva, no difference in GCF
Akalun et al. (2007)	Cross-sectional	36 CP, 28 healthy			Serum, saliva, GCF	Lipid peroxidation, malondialdehyde, total antioxidant status	LPO and TOS may play an important role in periodontitis
Baltacıoğlu et al. (2008)	Cross-sectional	33 CP, 24 healthy			Serum, GCF	Total protein, protein carbonylation	Elevated protein carbonylation may be a sign of oxidative stress in CP
Buduneli et al. (2005)	Cross-sectional	20 gingivitis, 20 CP, 20 healthy controls.			GCF, serum	t-PA, u-PA, PAI-1, PAI-2	GCF PAI-2 concentrations were higher in CP, gingivitis than healthy controls. Periodontal disease rather than smoking seems to affect PA system
Chapple et al. (2007)	Prospective	18 CP	3 months after SRP		GCF, plasma	TAOC	GCF TAOC was lower in CP, increased after SRP
Chen et al. (2009)	Cross-sectional	25 CP, 24 healthy			GCF, serum	PAF	Higher PAF in GCF, serum in CP
Dezerega et al. (2010)		15 progression, 18 CP, 10 healthy			GCF	MCP-3	
Dutzan et al. (2009)	Cross-sectional	106 moderate to advanced CP			GCF	IFN- γ	Higher in CP, higher in active sites than inactive sites, may be a diagnostic marker
Emingil et al. (2006b)	Cross-sectional	18 GAgP, 29 CP, 20 gingivitis, 20 healthy controls.		PD, CAL, BOP, PI	GCF	Laminin-5 gamma2-chain	Higher in active sites, may be an indicator for progression
Fitzsimmons et al. (2010)	Cross-sectional, population based	430 periodontitis, 509 healthy controls			GCF	IL-1 β , CRP	CP had elevated laminin-5 gamma2-chain levels
Gapski et al. (2009)	Multi-centre, prospective, controlled trial	CP SDD+surgery Placebo+surgery	12 months	1° outcome: CAL 2° outcome: PD, BOP, ICTP	GCF	ICTP	Higher levels in periodontitis, GCF levels of IL-1 β , CRP may indicate higher susceptibility
Garg et al. (2009)	Intervention	20 healthy 20 gingivitis 20 CP (also after SRP)	6–8 weeks after SRP	GI, PD, CAL	GCF	Cathepsin K	SDD decreased ICTP effectively Provided better clinical outcomes
Golub et al. (2008)	Prospective, controlled	CP, Postmenopausal women SDD: 64 Placebo: 64	2 years		GCF	ICTP, MMP-8, MMP-1, MMP-13, IL-1 β	Cathepsin K increased in CP, decreased with SRP
Grant et al. (2010)	Intervention	20 healthy, 20 CP	Before and 3 months after SRP		GCF	Reduced glutathione, oxidized glutathione	SDD decreased collagenase activity, ICTP
Holzhausen et al. (2010)	Cross-sectional, and intervention	40 moderate CP 40 severe CP 40 healthy controls	At baseline and also 1 month after SRP		GCF	Trypsin-like activity, TNF- α , IL-1 α , IL-6, IL-8, PAR ₂	GSH, GSSG lower in CP GSH: GSSG ratio increased after SRP
Ikezawa-Suzuki et al. (2008)	Intervention	35 CP	At baseline and after SRP		GCF	TNF- α , TNF receptor types 1 and 2	PAR ₂ was higher in CP than controls. IL-1 α , -6, -8 TNF- α , total proteolytic activity were all higher in CP Ratio of TNF R2: R1 increased after treatment, may be related to clinical periodontal data

Jepsen et al. (2003)	Cross-sectional	54 teeth with CP, 11 experimental gingivitis	GCF	lysylpyridinoline, hydroxylslypyridinoline	Increased hydroxylslypyridinoline, Lysylpyridinoline may be indicators of ongoing destruction
Kardeşler et al. (2008)	Cross-sectional	17 DM + periodontal disease, 17 healthy + periodontal disease, 17 healthy controls	GCF	PGE ₂ , IL-1 β , t-PA, PAI-2	PGE ₂ , t-PA were higher in DM, and periodontal disease groups than in healthy controls
Kurtiş et al. (1999)		24 CP, 24 healthy	GCF	IL-6	Higher in CP, no correlation with clinical data
Lamster et al. (1994)			GCF, saliva, serum	β -glucuronidase	GCF levels correlated with CAL
Lin et al. (2005)	Cross-sectional	14 patients (62 sites divided into 4 groups by PD, BOP).	GCF	Oncostatin M, IL-6	Total amounts but not concentrations correlated with disease severity.
Mogi & Orogoto (2007)	Cross-sectional	66 CP; 20 mild, 24 moderate, 22 severe	GCF	Cathepsin K	Increased in periodontitis <i>versus</i> control subjects, (+) correlation between cathepsin K and RANKL suggesting that both contribute to osteoclastic bone destruction
Mogi et al. (2004)	Cross-sectional	19 healthy controls	GCF	Cathepsin K, RANKL	Both increased in CP
Nakamura-Minami et al. (2003)	Intervention	21 CP	GCF	α -1 protease inhibitor, secretory leukocyte protease inhibitor	Both seem to be related with healing after SRP
Oringer et al. (2002)	Intervention	48 CP; SRP+placebo, SRP+minocycline	GCF	ICTP, IL-1	ICTP, IL-1 correlated with clinical data, may be disease markers
Palys et al. 1998	Cross-sectional	7 healthy and 8 gingivitis 21 CP	GCF subgingival plaque	ICTP, 40 subgingival taxa	Positive relation between ICTP and putative pathogens
Perinetti et al. (2008)	Intervention	15, 60 days after SRP	GCF	Alkaline phosphatase	Decreased on day 15, increased on day 60 after SRP
Pradeep et al. (2009a, b)	Intervention	20 healthy, 20 gingivitis, 20 CP patients also 6–8 weeks after SRP	GCF	MCP-1	Increased gradually with severity of periodontal disease, decreased after SRP, correlated with clinical periodontal data; may be a biomarker
Pradeep et al. (2009a, b)	Intervention	20 healthy, 20 gingivitis, 20 CP patients also 6–8 weeks after SRP	GCF	IL-18, MCP-1	Higher levels in CP, +correlation between IL-18 and MCP-1; IL-18 may induce MCP-1, promote inflammation
Rescala et al. (2010)	Cross-sectional	20 GCP, 17 GAgP, 10 gingivitis	GCF	IL-1 β , IL-2, IL-4, IL-8, elastase, IFN- γ	IL-1 β , elastase were higher in deep sites than shallow sites in periodontitis groups. No significant difference between CP and AgP in GCF levels of biomarkers
Rosin et al. (1999)	Cross-sectional	140 subjects	Saliva, GCF	Peroxidase, lysozyme	No significant correlation between clinical periodontal indices and salivary data
Srinath et al. (2010)	Cross-sectional	15 healthy, 15 gingivitis, 15 CP	GCF, Russell periodontal index	Melatonin	Lowest in CP, may have a protective role in periodontal disease
Suma et al. (2009)	Cross-sectional	Healthy Gingivitis CP	GCF, saliva	Lysozyme activity	Lysozyme activity correlated with PD
Teles et al. (2010)	Cross-sectional	20 healthy, 20 CP		IL-1 β , IL-8, MMP-8, 40 bacterial taxa	+ correlation between clinical indices, GCF cytokines, orange and red complexes

Table 1. (Contd.)

Reference	Study design	Study groups	Follow-up period	Clinical parameter	Biological sample	Biological parameter	Results
Thorat Manojkumar et al. (2010)	Intervention	20 healthy, 20 gingivitis, 20 CP	CP patients also 8 weeks after SRP	PD, CAL	GCF; subgingival plaque	Oncostatin M	Healthy sites of CP had higher cytokine levels than healthy group
Toker et al. (2008)	Intervention	15 healthy, 15 generalized AgP	6 weeks after SRP	PD, CAL, PI, BOP, GI	GCF	IL-1 β , IL-1ra, IL-10	Highest in CP, +correlation with PD, CAL; can be a biomarker in periodontitis
Tsalikis et al. (2001)	Intervention	12 advanced periodontitis	4 weeks after SRP	PD, CAL, BOP	GCF	Aspartate amino transferase	IL-1 β is high and decreased after SRP
Türkoglu et al. (2009)	Cross-sectional			PD, CAL, PI, BOP, PBI	GCF	IL-18, cathelicidin LL-37	AST decreased after SRP
Tüter et al. (2007)	Cross-sectional	20 CP, 17 healthy			Serum, GCF	hsCRP	Cathelicidin LL-37 increased in CP, may play a role in destruction
Yetkin Ay et al. (2009)	Cross-sectional	40 CP, 20 healthy		GI, PD, CAL, BOP, PI	GCF	IL-11, IL-17	No difference between groups
Yin et al. (2000)	Intervention	Periodontitis, gingivitis, healthy	14 days after SRP		GCF	t-PA, PAI-2	Healthy group had highest IL-11: IL-17 ratio
Yoshinari et al. (2004)	Intervention	7 CP	Before and after SRP	Routine clinical parameters	GCF, gingiva	IL-1 α , IL-1 β , IL-1ra	Decreased after SRP, may be diagnostic markers
Zheng et al. (2006)	Cross-sectional	21 CP, 19 gingivitis, 20 healthy controls			GCF, serum	PAF	IL-1 is effective for evaluating the gingival inflammation

CP, chronic periodontitis; GCF, gingival crevicular fluid; SOD, superoxide dismutase; LPO, lipid peroxidation; TOS, total oxidant status; t-PA, the tissue/blood vessel type plasminogen activator; u-PA, urokinase type plasminogen activator; PAI, plasminogen activator inhibitor; PA, plasminogen activators; TAOC, total antioxidant capacity; SRP, scaling and root planing; PAF, platelet activating factor; MCP-3, monocyte chemoattractant protein-3; IFN- γ , interferon- γ ; GAgP, generalized aggressive periodontitis; PD, probing depth; CAL, clinical attachment level; BOP, bleeding on probing; PI, plaque index; IL, interleukin; CRP, C-reactive protein; SDD, subantimicrobial dose doxycycline; ICTP, carboxy-terminal telopeptide of type I collagen; GI, gingival index; MMP, matrix metalloproteinase; GSH, reduced glutathione; GSSG, oxidized glutathione; PAR₂, protease activated receptor 2; TNF, tumour necrosis factor; DM, diabetes mellitus; PGE₂, prostaglandin E₂; RANKL, receptor activator of nuclear factor κ B ligand; MCP-1, monocyte chemoattractant protein-1; TACE, transcatheper arterial chemoembolization; SBI, sulcus bleeding index; IL-1ra, interleukin-1 receptor antagonist; AST, aspartate amino transferase; PBI, papilla bleeding index; hsCRP, high-sensitive C-reactive protein.

Table 2. Studies evaluating possible biomarkers in saliva samples

Reference	Study design	Study groups	Follow-up period	Clinical parameter	Biological sample	Biological parameter	Results
Aenaimanan et al. (2009)	Cross-sectional	30 localized CP, 30 generalized CP, 20 controls		PD, CAL, BOP	Saliva, subgingival plaque	Alanine aminopeptidase	Dipeptidyl peptidase IV but not alanine aminopeptidase activity is associated with CP, presence of <i>P. gingivalis</i>
Aurer et al. 1999	Cross-sectional	20 RPP, AP, 8 PD patients, 16 controls			Saliva	<i>P. gingivalis</i> IL-6	IL-6 increased in RPP, CRP decreased in AP
Baron et al. (1999)	Cross-sectional	37 patients, variable PD			Saliva	Cystatins	Total cystatin activity decreased in patients
Cutando et al. (2006)	Cross-sectional	175 volunteers		CPI	Whole saliva	Total cystatin activity	Inverse relation between salivary melatonin and PD severity
De la Pena et al. (2007)	Cross-sectional	17 PD, 20 controls		Number of teeth, PD	Saliva	Melatonin	Increased in periodontitis, correlated with PD, may be a biomarker
Diab-Ladki et al. 2003	Cross-sectional	69 Periodontal disease patients			Saliva	Lactate dehydrogenase	TAC decreased, an imbalance between oxidants and antioxidants
Garito et al. (1995)	Cross-sectional	22 CP, 22 healthy (half smoker, half non-smoker)		PD	Saliva	TAC, total protein content	Salivary PAF correlated with PD severity
Ghallab & Shaker (2010)	Intervention	18 CP, 18 controls	At baseline, 1 month after SRP	PI, GI, PD, CAL only in CP	Saliva	PAF	Highest in smoker CP, decreased after SRP in both smokers and non-smokers
Gheren et al. (2008)	Prospective	84 CP, 81 controls	30 days after perio. therapy	PD, CAL, PI, GI	Saliva	sCD44	Salivary arginase activity increased in CP, decreased after therapy: a candidate salivary marker of periodontal status
Gürsoy et al. (2009)	Cross-sectional	Severe periodontitis			Saliva	Salivary arginase activity	Only IL-1 β can differentiate periodontitis
Haigh et al. (2010)	Cross-sectional, intervention	380 subjects		PD, CAL, GI	Saliva	Elastase, lactate dehydrogenase	S100 proteins highly related with PD severity
Lamster et al. (2003)	Cross-sectional	89 mild, moderate, severe CP			Saliva	IL-6, IL-1 β , TNF- α	Periodontal clinical parameters correlated with salivary β glucuronidase
Koss et al. 2009	Cross-sectional	29 controls			Whole saliva	Proteomics	Peroxidase, hydroxyproline increased, sIgA diminished in CP
Nomura et al. (2006)	Cross-sectional	187 subjects			Saliva	β glucuronidase	Salivary lactate dehydrogenase, AST, blood urea nitrogen have potential to screen PD
Özmeric et al. (2000)	Cross-sectional	20 CP, 15 controls		PD, CAL, PI, GI	Saliva	Total protein, arginase	Salivary arginase increased in CP
Su et al. (2009)	Cross-sectional	58 periodontitis 234 controls		CPTN	Whole saliva	ROS, TAC	ROS increased in periodontitis, TAC correlates with disease severity
Takane et al. (2002)	Cross-sectional	78 CP, 17 healthy			Saliva	8-OHdG	Higher in CP, can be a diagnostic marker
Teles et al. (2009)	Cross-sectional	74 CP, 44 healthy			Saliva	G-MCSF, IL-1 β , -2, -4, -5, -6, -8, -10	No difference between groups. IL-8 correlated with PD, BOP, these cytokines cannot discriminate CP and healthy
Totan et al. (2006)	Cross-sectional	Periodontitis, controls		PD, BOP	Saliva	IFN- γ , TNF- α	Salivary AST, ALP increased with pockets, BOP, suppuration
Wilezyska-Borawska et al. (2006)	Cross-sectional	26 PD, 20 healthy controls		GI, PBI, PI, PD, CAL, number of teeth	Whole saliva	AST, ALP, aminopeptidases, glucuronidases	Salivary HGF may be related with severity of PD
Yoshie et al. (2007)	Intervention	49 CP	4 weeks after SRP	PD, CAL, BOP	Saliva	Hepatocyte growth factor	All decreased after SRP, they can be useful markers for inflammation, destruction

CP, chronic periodontitis; PD, probing depth; CAL, clinical attachment level; BOP, bleeding on probing; RPP, rapidly progressive periodontitis; AP, adult periodontitis; IL, interleukin; CRP, C-reactive protein; CPI, community periodontal index; TAC, total antioxidant capacity; PAF, platelet activating factor; PI, plaque index; sCD44, soluble CD44; SRP, scaling and root planing; TNF, tumour necrosis factor; sIgA, secretory immunoglobulin A; AST, aspartate amino transferase; CPTN, community periodontal index of treatment needs; ROS, reactive oxygen species; IFN- γ , interferon- γ ; ALP, alkaline phosphatase; PBI, papilla bleeding index; HGF, human gingival fibroblasts; LDH, lactate dehydrogenase.

periodontal pathogens that colonize in the subgingival area (Pussinen et al. 2007). Increased circulating levels of cytokines were reported in chronic periodontitis patients compared with the clinically healthy control subjects (Mooney & Kinane 1994). Data from the selected studies evaluating the potential utility of serum components as diagnostic markers for periodontal tissue destruction are presented in Table 3. Researchers have mainly focused on C-reactive protein (CRP), fibrinogen, and serum amyloid A (SAA) concentrations in serum in relation to periodontitis.

Molecular markers

The ratio of two molecules has gained attention for being indicative of the regulation of normal bone resorption and deposition activities that occur during bone remodelling, that is the ratio of receptor activator of NF- κ B ligand (RANKL) to osteoprotegerin (OPG) (Cochran 2008, Leibbrandt & Penninger 2008). RANKL, receptor activator of NF- κ B (RANK), and OPG interactions are important in coordinating osteoclastogenesis and alveolar bone resorption (Suda et al. 1999). RANKL is expressed by osteoblasts/stromal cells (Yasuda et al. 1998), fibroblasts (Quinn et al. 2000), and activated T cells (Horwood et al. 1999). It binds directly to RANK on the surface of preosteoclasts and osteoclasts. RANKL stimulates both the differentiation of osteoclast progenitors and activity of mature osteoclasts (Lacey et al. 1998). This ligand can be found as a cell membrane-bound variant (mRANKL) or in a primary soluble (secreted) form, which has been described in activated T cells (Kong et al. 1999). OPG is the naturally occurring inhibitor of osteoclast differentiation. It is a soluble molecule that binds to RANKL with high affinity and blocks RANKL from interacting with RANK (Lacey et al. 1998). As part of an inflammatory response, pro-inflammatory cytokines, such as interleukin-1 β (IL-1 β), -6, -11, -17, and tumour necrosis factor- α (TNF- α) can induce osteoclastogenesis by increasing the expression of RANKL while decreasing OPG production in osteoblasts/stromal cells (Nakashima et al. 2000). On the contrary, anti-inflammatory mediators such as IL-13 and interferon- γ (IFN- γ) may lower RANKL expression and/or increase OPG expression to inhibit osteoclastogenesis (Nakashima et al.

2000). Moreover, in a cross-sectional study, Sakellari et al. (2008) reported that soluble RANKL levels in GCF samples of the chronic periodontitis patients were higher than the periodontally healthy subjects, which also correlated with *Treponema denticola* and *Porphyromonas gingivalis* levels in the subgingival plaque samples of the same periodontitis patients. However, they failed to show a correlation between GCF RANKL levels and clinical periodontal measurements, namely PD, gingival recession and BOP either cross-sectionally or longitudinally. The concentrations of RANKL and OPG show great variation from study to study, but the ratio of RANKL/OPG has a consistent tendency to increase from periodontal health to periodontitis and to decrease following non-surgical periodontal treatment (Table 4) (Crotti et al. 2003, Mogi et al. 2004, Kawai et al. 2006, Lu et al. 2006, Bostancı et al. 2007, Wara-Aswapati et al. 2007, Buduneli et al. 2008, 2009, Dereka et al. 2010). Thus the ratio of RANKL/OPG shows promise as a discloser of periodontal disease activity.

Alkaline phosphatase (ALP) is a catalysing enzyme that accelerates the removal of phosphate groups in various molecules. ALP was not supported to have a predictive value for periodontal breakdown, but it may serve as a marker in periodontal treatment planning and monitoring (McCauley & Nohutçu 2002, Taba et al. 2005). Aspartate amino transferase (AST) was found to be increased in GCF of severe periodontitis patients decreasing after initial periodontal treatment (Tsalikis et al. 2001).

GCF oncostatin M levels were highest in chronic periodontitis patients when compared with gingivitis patients and healthy control subjects (Thorat Manojkumar et al. 2010). Serum oncostatin M levels were also associated with PD and CAL in chronic periodontitis patients (Pradeep et al. 2010). The positive correlations between oncostatin M levels and PD as well as CAL suggest that it may be a useful biomarker in periodontitis.

Cathepsin B is a proteinase which is synthesized mainly by macrophages in GCF and levels of cathepsin B may distinguish gingivitis from periodontitis (Kennett et al. 1997). Furthermore, GCF levels of cathepsin B correlate significantly with clinical parameters before and also after periodontal treatment, suggesting a use for this enzyme in

treatment planning and monitoring (Chen et al. 1998).

Cathepsin K, a cysteine protease, is capable of hydrolysing extracellular bone matrix proteins, is highly expressed in osteoclasts, and it is a well-known marker of osteoclast activity (Motyckova et al. 2001). Eley & Cox (1996a,b) have suggested that GCF cathepsin B level may indeed serve as a predictor of attachment loss. Mogi & Otogoto (2007) investigated GCF levels of cathepsin K and RANKL in 20 mild, 24 moderate, and 22 severe chronic periodontitis patients in comparison with 19 healthy control subjects. Cathepsin K was below the detection limit in the healthy control group, whereas it was detectable in all periodontitis samples. In a recent intervention study, Garg et al. (2009) reported that GCF levels of cathepsin K was higher in chronic periodontitis patients than gingivitis patients and healthy controls, whereas it was decreased significantly after scaling and root-planing. The high levels of cathepsin K in periodontitis samples together with the positive correlation between cathepsin K and RANKL levels suggest that both of them contribute to osteoclastic bone resorption in periodontitis, although no significant correlation could be found between the biomarkers and clinical periodontal measurements.

Carboxyterminal telopeptide pyridinoline cross-links of type I collagen (ICTP) is released into the periodontal tissues as a consequence of collagen degradation and alveolar bone resorption (Seibel 2003). Type I collagen composes 90% of the organic matrix of bone and is the most abundant collagen in osseous tissue (Narayanan & Page 1983). Studies assessing the role of ICTP levels in GCF or peri-implant crevicular fluid as a diagnostic marker of periodontal disease activity have had promising results so far (Oringer et al. 1998, 2002). ICTP was suggested to predict future bone loss, to correlate with clinical parameters and putative periodontal pathogens and also to reduce following periodontal therapy, thus leading to an accurate assessment of tissue breakdown (Giannobile 1999). Because of its specificity and sensitivity for bone resorption, ICTP represents a potentially valuable diagnostic marker for periodontal disease (Giannobile et al. 2003).

Osteocalcin (OC) is a calcium-binding protein of bone and the most abundant non-collagenous protein of the mineralized tissue (Lian & Gundberg

Table 3. Studies evaluating possible biomarkers in serum/plasma samples

Reference	Study design	Study groups	Follow-up period	Clinical parameter	Biological sample	Biological parameter	Results
Al-Ghamdi & Anil (2007)	Cross-sectional Follow-up	30 smoker CP, 30 non-smoker CP, 30 healthy	6 years	PD, CAL	Serum	Immunoglobulins	Smoking decreases Ig content
Amarasena et al. (2008)	Intervention	266 elderly	Progression: CAL increase > 3mm Before, 4-weeks after SRP		Serum	Albumin, Ca, IgG, A, M	Serum Ca correlated with PD, may be a risk factor for progression
Behle et al. (2009)	Intervention	30 CP			Plasma	19 biomarkers, multiplex	Overall reduction in systemic inflammation, great personal variation
De Queiroz et al. (2008)	Cross-sectional	17 CP, 8 healthy			Serum	24 cytokines with multiplex	RANTES differed between groups
Duarte et al. (2010)	Intervention	14 healthy controls, 14 CP, 14 GAgP	At baseline and 6 months after SRP		Serum	IL-17, IL-4, IL-23, IFN- γ , TNF- α	IL-17, TNF- α levels were higher in GAgP than CP, healthy both at baseline and after SRP
Funugen et al. (2008)	Cross-sectional	158 Japanese elders			Serum	Adiponectin, resistin, IL-6, TNF- α	Increased serum cortisol is associated with BOP
Glas et al. (2008)	Cross-sectional	105 CP, 122 healthy			Plasma	Surfactant protein D	Increased in CP; may be a biomarker for CP
Guenisch et al. (2009)	Cross-sectional	425 non-smoker, systemically healthy adults		Number of teeth with PD > 4–6 mm	Serum	IL-6, TNF- α	Serum IL-6 is associated with periodontal disease severity
Ishisaka et al. (2008)	Cross-sectional	467 adults		PD, CAL, BOP	Serum	Cortisol	Serum cortisol associated with PD, CAL
Iwasaki et al. (2008)	Follow-up	600 elderly	4 years	CAL	Serum	Albumin	Serum albumin may be a risk predictor for progression
Liu et al. (2010)	Cross-sectional	40 CP, 40 healthy		PD, CAL, BOP	Serum	CRP, lipid profile	CRP higher in CP, HDL lower in CP. correlated with clinical parameters
Marcacini et al. (2009)	Intervention	25 CP, 20 healthy	3 months after SRP	PD, BOP, CAL	Serum, plasma	hsCRP, IL-6, CD40 ligand, MCP-1, sP-selectin, sVCAM-1, sICAM-1	SRP decreased CRP and IL-6, others were not affected
Nakajima et al. (2010)	Intervention	78 CP, 40 healthy	At baseline, after SRP		Serum	hsCRP, IL-6, TNF- α	CRP, IL-6 higher and TNF lower in CP, IL-6, CRP decreased after SRP
Nibali et al. (2007)	Cross-sectional	302 CP, 183 healthy			Serum	Inflammatory, metabolic markers	CP may be related with systemic inflammation, dysmetabolic state
Nicu et al. (2009)	Cross-sectional	105 CP, 57 healthy			Serum	CRP, sCD14	sCD14 increased in CP
Offenbacher et al. (2009)	RCT	151 SRP, 152 community care	6 months	PD, CAL, BOP, subgingival calculus	Serum	CRP	Serum CRP was not affected by SRP
Pradeep et al. (2010)	Intervention	20 healthy, 20 gingivitis, 20 CP	CP patients also 6–8 weeks after SRP	PD, CAL, GI	Serum	Oncostatin M	Highest in CP, decreased after SRP, not detectable in gingivitis, healthy. Correlated with PD, CAL
Raunio et al. (2007)	Cross-sectional	52 subjects		PD, CAL, BL	Mouth-wash, serum	IL-6	Serum IL-6 increased with periodontal disease severity
Raunio et al. (2009)	Cross-sectional	56 CP, 28 healthy			Serum	sCD14	Higher in CP, correlated with periodontal disease extent
Renvert et al. (2009)	RCT	28 CP test (anti-inflammatory therapy), 29 CP placebo		PD, BOP	Serum	hsCRP, IL-6, IL-1 β , IL-8, IL-12, TNF- α , IFN- γ	CRP, IFN- γ , IL-6 decreased, others did not change
Trindade et al. (2008)	Cross-sectional	26 healthy			Serum	Antibody levels against <i>P. gingivalis</i>	Serum Ig titres correlated with clinical periodontal diagnosis
Yoshihara et al. (2009)	Cross-sectional	148 subjects at age 77		CAL	Serum	Osteocalcin, bone-specific alkaline phosphatase	Negative correlation between CAL > 6 mm and osteocalcin

BL, bone level; CP, chronic periodontitis; CAL, clinical attachment level; PD, probing depth; Ig, immunoglobulin; RANTES, regulated upon activation normal T-cell expressed and secreted; GAgP, generalized aggressive periodontitis; SRP, scaling and root planing; IL, interleukin; IFN- γ , interferon- γ ; TNF, tumour necrosis factor; BOP, bleeding on probing; CRP, C-reactive protein; HDL, high-density lipoprotein; MCP, monocyte chemoattractant protein; sVCAM, soluble vascular cell adhesion molecule; sICAM, soluble intercellular adhesion molecule.

Table 4. Studies evaluating RANKL, OPG, ICTP in gingival crevicular fluid (GCF), saliva or serum samples

Reference	Study design	Study groups	Follow-up period	Clinical parameter	Biological sample	Biological parameter	Results
Bostanci et al. (2007)	Cross-sectional	21 healthy, 22 gingivitis, 28 CP, 25 GAgP, 11 CP immunosuppressed			GCF	RANKL, OPG	RANKL levels increased, OPG decreased in periodontitis <i>versus</i> gingivitis, healthy. RANKL/OPG ratio may predict disease occurrence
Bostanci et al. (2008)	Cross-sectional	21 healthy, 22 gingivitis, 28 CP, 25 GAgP, 11 CP immunosuppressed			GCF	TACE	GCF TACE levels were higher in periodontitis, TACE showed (+) correlation with PD, CAL, GCF RANKL concentration
Buduneli et al. (2008)	Cross-sectional	67 untreated CP, 44 maintenance patients			Saliva	sRANKL, OPG	Salivary sRANKL, OPG may be affected by smoking and showed significant differences between treated <i>versus</i> untreated CP
Buduneli et al. (2009)	Intervention	10 smoker, 10 non-smoker CP	4 weeks after SRP		GCF	IL-17, sRANKL, OPG	OPG decreased, IL-17 increased, sRANKL always similar
Frodge et al. (2008)	Cross-sectional	35 moderate to severe CP, 39 healthy controls			Saliva	TNF- α , ICTP, RANKL	TNF- α detected in all subjects, ICTP, RANKL were detected only in a minority. TNF- α was higher in CP than controls. Subjects with salivary TNF- α levels above a threshold had more sites with BOP, PD > 4 mm, CAL > 2 mm. Salivary TNF- α may be a biomarker for periodontal destruction
Golub et al. (2008)	Prospective, controlled	CP, Post-menopausal women SDD: 64 Placebo: 64	2 years		GCF	ICTP, MMP-8, MMP-1, MMP-13, IL-1 β	SDD decreased collagenase activity, ICTP
Golub et al. (2010)	Prospective	128 postmenopausal women with CP SDD or placebo in maintenance	2 years		Serum	Osteocalcin, ICTP, CTX, bone-specific alkaline phosphatase	SDD decreased significantly serum biomarkers of bone resorption
Gürlek et al. (2009)	Cross-sectional	34 smokers, 22 non-smokers, 11 ex-smokers			Saliva	ICTP, OC, cotinine	ICTP was similar in study groups, OC was lower in smoker than the other groups, correlated negatively with pack-years
Mogi et al. (2004)	Cross-sectional				GCF	Cathepsin K, RANKL	Both increased in CP
Oringer et al. (2002)	Intervention	48 CP; SRP+placebo, SRP+minocycline	Before, 1,3,6 months after SRP		GCF	ICTP, IL-1	ICTP, IL-1 correlated with clinical data, may be disease markers
Özçaka et al. (2010)	Cross-sectional	58 CP, 47 healthy controls			plasma	ICTP, osteocalcin	CP and healthy control groups were similar in plasma ICTP, OC levels
Palys et al. (1998)	Cross-sectional	7 healthy, 8 gingivitis 21 CP		BOP, PD, CAL, PI	GCF subgingival plaque	ICTP, 40 subgingival taxa	Positive relation between ICTP and putative pathogens
Sakellari et al. (2008)	Cross-sectional	35 CP, 38 periodontally healthy		PD, recession, BOP	GCF, subgingival plaque	Free sRANKL, <i>Aggregatibacter actinomycetemcomitans</i> , <i>Tannerella forsythensis</i> , <i>Treponema denticola</i>	sRANKL was higher in CP, correlated with <i>Treponema denticola</i> , <i>P. gingivalis</i> but not with clinical parameters
Silva et al. (2008)	Longitudinal	56 CP moderate to severe, followed until progression			GCF	RANKL, MCP-1, TNF- α , IL-1 β , MMP-13	Higher RANKL, IL-1 β , MMP-13 activity in active sites

CP, chronic periodontitis; GAgP, generalized aggressive periodontitis; GCF, gingival crevicular fluid; RANKL, receptor activator of nuclear factor κ B ligand; OPG, osteoprotegerin; sRANKL, soluble receptor activator of nuclear factor κ B ligand; IL, interleukin; TNF, tumour necrosis factor; ICTP, carboxy-terminal telopeptide of type I collagen; BOP, bleeding on probing; PD, probing depth; CAL, clinical attachment level; SDD, subantimicrobial dose doxycycline; MMP, matrix metalloproteinase; CTX, carboxy-terminal collagen crosslinks; OC, osteocalcin; SRP, scaling and root planing; PI, plaque index; MCP, monocyte chemoattractant protein.

1988). Osteoblasts, odontoblasts, and chondrocytes produce OC (Gallop et al. 1980). Serum level of OC is considered as a marker of bone formation (Christenson 1997, Giannobile et al. 2003). Lower serum levels of OC were reported in periodontitis patients compared with healthy subjects suggesting lower osteoblastic activity and bone formation ability (Shi et al. 1996). Yoshihara et al. (2009) reported that serum OC levels correlated with CAL in elderly Japanese subjects. On the other hand, indifferent plasma levels of ICTP and OC were reported in a recent study comparatively evaluating chronic periodontitis patients and healthy control subjects (Özçaka et al. 2010). There is no consensus, yet on the potential usage of OC for assessment of periodontal diagnosis and prognosis.

Osteonectin is a single-chain polypeptide that binds strongly to hydroxyapatite and other extracellular matrix molecules like collagens. It may be a sensitive marker for detection of periodontal disease status. However, no clinical study reporting osteonectin levels in biological fluids in relation with periodontitis could be found in the present literature search.

Osteopontin (OPN), a glycosylated phosphoprotein, is a bone matrix component produced by osteoblasts, osteoclasts, and macrophages as a multifunctional cytokine. It is a single-chain polypeptide which is concentrated at clear zone attachment areas of plasma membrane where osteoclasts are attached to the underlying mineral surface that is, the clear zone attachment areas of the plasma membrane (Rodan 1995). Kido et al. (2001) showed the presence of OPN in GCF and also showed that GCF OPN levels increased parallel to the increase in PD. Sharma & Pradeep (2006) reported that GCF OPN concentrations increased proportionally with the progression of disease and decreased significantly following non-surgical periodontal treatment. The same authors reported that chronic periodontitis patients exhibited the highest GCF and plasma OPN levels which decreased significantly following initial periodontal treatment (Sharma & Pradeep 2007). The GCF OPN levels correlated with plasma levels and clinical attachment loss suggesting a role in the pathogenesis of periodontitis. Thus, OPN seems to be a promising biomarker for periodontal disease progression.

Matrix metalloproteinases (MMPs) are considered as modifiers of host

response and it has been suggested that their role and involvement should not be interpreted solely as surrogate promoters of tissue destruction, but also as defensive or protective factors against inflammation as a whole (Sorsa et al. 2004, 2006). MMPs represent a structurally related but genetically distinct superfamily of proteases acting not only in physiological development and tissue remodelling but also in pathological tissue destruction (Sorsa et al. 2004). MMPs can be divided into five major groups: collagenases (MMP-1, -8, -13), gelatinases (MMP-2, MMP-9), stromelysins (MMP-3, -10, -11), membrane-type MMPs (MMP-14, -15, -16, -17), and others (Sorsa et al. 2004). MMPs can collectively degrade almost all components of extracellular matrix and basement membrane and their excess activity lead to periodontal tissue destruction. The interplay of cell-cell and cell-matrix interactions involving the production of enzymes, activators, inhibitors, cytokines, and growth factors regulate not only connective tissue remodelling but also connective tissue matrix destruction (Reynolds & Meikle 1997). MMPs produced by resident cells including fibroblasts, macrophages, neutrophils, and epithelial cells, are controlled by tissue inhibitors of MMPs (TIMPs). MMPs can also process bioactive non-matrix substrates such as cytokines, chemokines, growth factors, and immune modulators thereby mediating anti-inflammatory and pro-inflammatory processes (Sorsa et al. 2006, Kuula et al. 2009). Upon bacterial insult, triggered leucocytes migrate to the site of inflammation and release MMP-8 and MMP-9, which are activated locally (Sorsa et al. 2006). TIMPs regulate the activities of these enzymes and TIMP-1 is more effective on interstitial collagenases. An imbalance between MMPs and TIMPs result in the pathological tissue destruction observed in periodontitis (Table 5) (Aiba et al. 1996, Bıyıkoglu et al. 2009, Kardeşler et al. 2010).

Salivary TIMP-1 concentration has been reported to be lower in periodontitis patients than the healthy controls, while collagenase activity was higher, which exhibited reciprocal changes after initial periodontal therapy (Hayakawa et al. 1994). Furthermore, saliva samples from healthy subjects consisted mainly procollagenase whereas active collagenase predominated in diseased subjects. Greater collagenase activity has been found also in GCF samples

of periodontitis patients than the healthy subjects (Villega et al. 1987). In a prospective study on chronic periodontitis patients comparing active *versus* inactive sites, significantly higher GCF levels of MMP-13 were reported in active sites, suggesting that it could serve as a marker of disease progression (Hernandez et al. 2006).

Ramseier et al. (2009) reported from a recent cross-sectional study that salivary concentrations of MMP-8, -9, and OPG combined with red complex bacteria predicted periodontal disease. In another recent study (Hernandez-Rios et al. 2009), active sites in moderate to advanced chronic periodontitis patients followed for 2 months were defined and GCF levels of MMP-9 and -13 were suggested as useful biomarkers for periodontitis progression. Another cross-sectional study reported that GCF MMP-8, -9 levels correlated with disease activity in chronic periodontitis patients (Beklen et al. 2006). Supporting data came from an intervention study comprising 28 chronic periodontitis patients and 22 controls (Marcaccini et al. 2009). The authors reported higher MMP-3, -8, -9, and gelatinolytic activities in plasma of chronic periodontitis patients which decreased significantly 3 months after non-surgical periodontal treatment. Similar data on GCF MMP-8 levels were also reported by Chen et al. (2000). Kinane et al. (2003) also reported that GCF MMP-8 levels decreased significantly 3 months after non-surgical periodontal therapy in 20 chronic periodontitis patients. Various treatment modalities have been investigated to determine their potential to control the activities of MMPs in periodontal tissue destruction (Buduneli et al. 2002, 2007, Golub et al. 2008, Vardar-Şengül et al. 2008).

GCF MMP-8 levels in shallow crevices were reported to be associated with CAL and thus it was suggested as a prognostic marker (Passoja et al. 2008). Higher MMP-8 levels in saliva and higher MMP-9 levels in GCF of chronic periodontitis patients were detected in comparison with gingivitis patients and healthy control subjects (Rai et al. 2008). Significant positive correlations were found between periodontal disease severity and GCF MMP-8, -9 activities together with negative correlations with TIMP-1, -2 levels (Pozo et al. 2005). Accordingly, GCF MMP-8 activation was found to be the highest in chronic periodontitis patients compared with healthy subjects and

Table 5. Studies evaluating matrix metalloproteinases in GCF, saliva and serum/plasma samples

Reference	Study design	Study groups	Follow-up period	Clinical parameter	Biological sample	Biological parameter	Results
Alfant et al. (2008)	Cross-sectional	44 children with/without AgP CP		PD	GCF	MMP-1, -2, -3, -8, -9, -12, -13	MMP levels higher in AgP than the others, especially in diseased sites
Alpagot et al. (2001)	Prospective	Healthy children 40 CP	6 months	GI, PI; BOP, PD, CAL, supuration	GCF	MMP-3, TIMP-1	No correlation with PD High GCF MMP-3, TIMP-1 may indicate high risk for progression
Beklen et al. (2006)	Cross-sectional	CP			GCF	MMP-3, -8, -9	GCF MMP-8, -9 levels correlated with disease activity
Bildt et al. (2008)	Cross-sectional	8 healthy, 12 CP			GCF	MMPs, TIMPs	MMPs higher, TIMPs lower in CP, with high individual variation
Emingil et al. (2006c)	Cross-sectional	35 GAgP, 29 CP, 20 gingivitis, 21 healthy controls.		PD, CAL, BOP, PI	GCF	MMP-25, MMP-26	MMP-25, -26 levels increased in both periodontitis groups and associated with severity of periodontal inflammation, suggesting that they can have a role in disease progression
Emingil et al. (2006a)	Cross-sectional	20 GAgP, 20 CP, 20 gingivitis, 20 healthy controls		PD, CAL, BOP, PI	GCF	MMP-7, TIMP-1, EMMPRIN	MMP-7 levels were similar in groups. All patient groups had higher EMMPRIN, TIMP-1. Periodontitis groups had higher TIMP-1 than gingivitis. Increased EMMPRIN, TIMP-1 related with severity of periodontal inflammation
Gürsoy et al. (2010)	Cross-sectional	84 CP, 81 controls			Saliva	MMP-8, -14 TIMP-1, ICTP	MMP-8, TIMP-1, ICTP increased in CP and can differentiate periodontitis
Heikkinen et al. (2010)	Cross-sectional	258 boys, 243 girls all adolescents, smoker <i>versus</i> non-smokers			Saliva	MMP-8, elastase	Median MMP-8 and elastase were lower in smokers, boys seem to be more susceptible than girls
Hernandez et al. (2006)	Prospective	21 CP Active and inactive sites			GCF	MMP-13	Both biomarkers correlated with BOP Active sites had higher MMP-13; a marker of disease progression
Hernandez-Rios et al. (2009)	Prospective, active sites defined in 2 months	Moderate-advanced CP		PD, CAL, BOP	GCF	MMP-9, MMP-13, TIMP-1	MMP-13 activity significantly elevated in active sites, high levels of ICTP, associated with progression of periodontal breakdown. MMP-9, -13 may be useful biomarkers for periodontitis progression
Kardeşler et al. (2010)	Cross-sectional	12 DM gingivitis, 12 DM periodontitis, 12 H gingivitis, 13 H periodontitis, 24 healthy controls. 20 CP, before and after SRP			GCF	MMP-8, MMP-13, TIMP-1	MMP-8 total amounts were higher in periodontitis, DM-G groups. MMP-13, TIMP-1 total amounts were similar in study groups
Kinane et al. (2003)	Intervention	11 CP, 10 gingivitis, 8 healthy controls	3 months	PD, CAL	GCF	MMP-8	MMP-8 decreased with SRP
Mäntylä et al. (2003)	Intervention	CP	Before and after SRP	Routine clinical parameters	GCF	MMP-8	Chair-side MMP-8 can differentiate CP from gingivitis, healthy, and monitor treatment of CP
Mäntylä et al. (2006)	Prospective	Smoker <i>versus</i> NS	12 months		GCF	MMP-8	Persistent elevation in MMP-8 may indicate high risk, poor response to therapy
Marcaccini et al. (2009)	Intervention	28 CP, 22 controls	before and 3 months after SRP		Plasma	MMP-2, -3, -8, -9, TIMP-1, -2, total gelatinolytic activity	Higher MMP-3, -8, -9, gelatinolytic activity in CP. MMP-8, -9, gelatinolytic activity decreased after therapy

Miller et al. (2006)	Cross-sectional	28 moderate-severe CP, 29 healthy controls	Saliva	IL-1 β , MMP-8, OPG	Elevated IL-1 β , MMP-8 but not OPG significantly increased risk of periodontal disease. Their salivary levels may be biomarkers for periodontal disease
Passoja et al. (2008)	Cross-sectional	48 CP patients	GCF, serum	MMP-8	MMP-8 in shallow crevices associated with CAL, can be a prognostic marker
Pozo et al. (2005)	Intervention	13CP, 11 healthy	GCF	MMP-8, -9, TIMP-1, -2	Significant + correlations between periodontal disease severity and MMP activity, - correlations with TIMP-1, -2
Rai et al. (2008)	Cross-sectional	15 healthy 18 gingivitis 20 CP	GCF, saliva	MMP-2, -8, -9	Salivary MMP-8, GCF MMP-9 were higher in CP GCF MMP-2 was lower in CP
Ramseier et al. (2009)	Cross-sectional	50 healthy/gingivitis 50 periodontitis	Whole saliva	MMP-8, -9, OPG	High correlation with PD, BOP MMP-8, -9, OPG combined with red complex predicted periodontal disease
Xu et al. (2008)	Cross-sectional	Healthy, gingivitis, CP	GCF	MMP-8 isoenzymes (fibroblast and PMN-type)	CP exhibited highest activation of GCF MMP-8 isoenzymes

AgP, aggressive periodontitis; CP, chronic periodontitis; PD, probing depth; MMP, matrix metalloproteinase; GI, gingival index; PI, plaque index; BOP, bleeding on probing; CAL, clinical attachment level; TIMP, tissue inhibitors of metalloproteinases; GAgP, generalized aggressive periodontitis; ICTP, carboxy-terminal telopeptide of type I collagen; DM, diabetes mellitus; SRP, scaling and root planing; OPG, osteoprotegerin.

gingivitis patients (Xu et al. 2008). In cross-sectional studies, elevated GCF levels of MMP-7, TIMP-1 (Emingil et al. 2006c), laminin-5 gamma2-chain (Emingil et al. 2006a), MMP-25, -26 (Emingil et al. 2006b) were reported in chronic periodontitis patients compared to gingivitis patients and healthy control subjects. High GCF MMP-13, TIMP-1 levels were suggested to indicate high risk for periodontal disease progression (Alpagot et al. 2001). Similarly, persistent elevation of MMP-8 in GCF samples was regarded as indicating high risk and poor response to periodontal therapy (Mäntylä et al. 2006). Very recently, Gürsoy et al. (2010) also confirmed that salivary MMP-8 and TIMP-1 are increased in chronic periodontitis patients differentiating periodontitis patients from the controls. Furthermore, a chair-side MMP-8 test was indicated to effectively differentiate chronic periodontitis from gingivitis and clinically healthy sites and thus, to monitor treatment of chronic periodontitis patients (Mäntylä et al. 2003).

Neutrophil elastase (NE) is one of the most destructive enzymes with the capability of degrading almost all extracellular matrix components as well as plasma proteins and activating pro-MMPs and inactivating TIMP-1 (Meyle et al. 1992, Eley & Cox 1996a, b, Sorsa et al. 2006, Geraghty et al. 2007). A high concentration of NE is stored in azurophilic granules of PMNs, providing an important step in host defence. When activated, NE can be released rapidly into the extracellular space and cause local tissue damage (Kawabata et al. 2002). Endogenous proteinase inhibitors are important to protect tissues from unregulated proteolysis. Once released in circulation, NE is rapidly inactivated by conjugating with protease inhibitors. Heikkinen et al. (2010) reported that salivary concentrations of MMP-8 and NE correlated with BOP in a cross-sectional study comprising 501 adolescents.

Azurophilic granules of PMNs also contain the enzyme myeloperoxidase (MPO), which can generate a reactive oxidant species including hypochlorous acid (HOCl). MPO is produced by neutrophils whose oxidant products are capable of modifying low-density lipoprotein cholesterol and has been shown to be present in human atheromas and unstable plaques (Nicholls & Hazen 2005). MPO is released into the extracellular environment following neutro-

phil stimulation and/or degranulation (Buchmann et al. 2002). MPO can oxidatively activate MMP-8 and -9 and inactivate TIMP-1 (Sorsa et al. 2006). Decreased plasma MPO levels following periodontal treatment of severe periodontitis patients suggests that MPO may play a major role in pathogenesis of destructive periodontal diseases (Behle et al. 2009). Thus, MPO and NE can potentiate the destructive MMP cascades and indeed MPO has been regarded as a promising marker of periodontal disease activity (Arbes et al. 1999, Yamalik et al. 2000, Wei et al. 2004).

Eley & Cox (1995) have investigated whether GCF levels of dipeptidyl peptidase (DPP) II or IV levels, total activity and concentration could predict progressive attachment loss and suggested that both GCF DPP II and IV may be predictors of periodontal attachment loss. In another early study, Cox & Eley (1992) have also analysed cathepsin B/L-, elastase-, tryptase-, trypsin-, and DPP IV-like activities in GCF samples obtained from chronic periodontitis patients before and also after non-surgical periodontal treatment. GCF protease levels were suggested to reflect the clinical status of periodontal lesions and may thus be of value in monitoring disease activity.

Prostaglandin E_2 (PGE_2), a metabolite of the cyclooxygenase (COX) pathway, is an arachidonic acid metabolite and considered a potent mediator of alveolar bone loss in periodontitis (Offenbacher et al. 1984, 1993). PGE_2 is known to have an activity on fibroblasts and osteoclasts to induce the synthesis of MMPs, IL-1 β and other cytokines. PGE_2 has been detected in higher levels in gingival tissue and GCF proportional to the severity of periodontal disease. IL-1 β is a central mediator of inflammation and connective tissue destruction in rheumatoid arthritis (Raymond et al. 2006). IL-1 β increases matrix degradation also by inducing the production of PGE_2 in synovial cells, as well by its role as a mediator of bone and cartilage destruction (Cutolo 2004). TNF- α is a less potent stimulator of PGE_2 production compared to IL-1 β , but these two cytokines synergistically enhance PGE_2 production (Yücel-Lindberg et al. 1999). Several of the cytokines stimulating RANKL expression and bone resorption also enhance the expression of COX-2 and prostaglandin production (Lerner 2006). PGE_2 is also

an efficient stimulator of RANKL expression in osteoblasts (Li et al. 2002) as well as on osteoclast progenitor cells (Ono et al. 2005). At persistent high concentrations, PGE_2 stimulates the master osteoclast activator; RANKL expression in stromal cells/osteoblasts and eventually causes enhanced bone resorption. Offenbacher et al. (1986) demonstrated that PGE_2 levels in GCF of patients exhibiting periodontal diseases are significantly higher than those in periodontally healthy subjects, and furthermore, that GCF PGE_2 concentrations are effective for predicting periodontitis progression, i.e. attachment loss, with a high degree of sensitivity and specificity (0.76 and 0.96, respectively). Evidence indicating the role of prostaglandins in periodontal tissue destruction also comes from human studies evaluating potential effects of non-steroidal anti-inflammatory drugs in periodontal treatment (Buduneli et al. 2002, 2010, Vardar et al. 2003).

Pro-inflammatory cytokines also play a significant role in the pathogenesis of periodontal diseases in terms of both soft and hard tissue destruction. Besides IL-1 β and TNF- α , IL-6 is also hypothesized to stimulate bone resorption (Ishimi et al. 1990). TNF- α , IL-1 β , -6, -8, -18 have been found to be abundantly expressed in human gingiva, and increased levels have been reported in GCF of periodontitis patients (Preiss & Meyle 1994, Boch et al. 2001, Graves & Cochran 2003, Toker et al. 2008, Pradeep et al. 2009a, b, Fitzsimmons et al. 2010, Teles et al. 2010). Increased GCF levels of IL-1 β and NE have been reported with increasing level of gingival inflammation also in experimental gingivitis studies and indeed elastase activity has been suggested as an excellent quantitative measure of gingival inflammation (Hermann et al. 2001, Gonz  les et al. 2001). On the contrary, mean salivary levels of granulocyte-macrophage colony-stimulating factor, IL-1 β , -2, -4, -5, -6, -8, -10, interferon gamma (IFN- γ), and TNF- α failed to discriminate between periodontal health and disease (Teles et al. 2009). Neutralization of IL-1 and TNF- α by soluble receptors has been reported to decrease osteoclast formation and bone loss in experimental periodontitis (Assuma et al. 1998). Silva et al. (2008) followed 56 moderate to severe chronic periodontitis patients until progression of periodontal destruction was detected and they reported significantly higher GCF levels

of RANKL, IL-1 β , and MMP-13 in active sites than the inactive sites. Very recently, Rescala et al. (2010) conducted a cross-sectional study comprising 20 chronic periodontitis patients, 17 generalized aggressive periodontitis patients and 10 gingivitis patients. They reported that IL-1 β and elastase levels in the GCF samples were higher in deep sites compared with the shallow sites in both of the periodontitis groups and suggested that these biomarkers may indicate periodontal tissue destruction. Furthermore, Frodge et al. (2008) evaluated salivary concentrations of TNF- α , RANKL, and ICTP in 35 subjects with moderate to severe chronic periodontitis in comparison with 39 healthy controls. The authors reported that salivary TNF- α levels were significantly elevated in chronic periodontitis patients suggesting the utility of this biomarker in a panel of salivary parameters that could facilitate the screening, diagnosis, and management of periodontal disease. In another recent study, Yetkin Ay et al. (2009) investigated GCF IL-11 and IL-17 levels in 40 chronic periodontitis patients and 20 healthy controls. It was reported that the IL-11/IL-17 ratio was significantly higher in the healthy control group than the periodontitis patients, whereas shallow sites in the periodontitis patients exhibited higher ratios than the deep sites in the same patients. Thus, an imbalance in the pro- and anti-inflammatory cytokines may be responsible for periodontal breakdown through immune responses.

CRP is a circulating acute-phase protein acting in an innate immune response. Significantly higher serum high-sensitivity CRP (hs-CRP) concentrations were found in periodontitis patients than the healthy controls in various case-control studies (Ebersole et al. 1997, Fredriksson et al. 1999, Loos et al. 2000, Glurich et al. 2002, Craig et al. 2003, Liu et al. 2010). On the other hand, T  ter et al. (2007) found no difference in serum CRP levels between the chronic periodontitis and healthy control groups, and serum CRP levels were not associated with PD. Moreover, periodontal treatment modalities resulted in decreases in serum CRP concentrations (Ebersole et al. 1997, Nakashima et al. 2000, Mattila et al. 2002, Swoboda et al. 2008, Marcaccini et al. 2009, Katagiri et al. 2009, Renvert et al. 2009, Golub et al. 2010). However, Offenbacher et al. (2009) reported that serum CRP concentration was not affected by scaling and root planning.

Serum amyloid A (SAA) is another acute-phase protein, which is associated with high-density lipoprotein (HDL). Elevated SAA concentrations in blood circulation have been found in subjects with both periodontitis and CVDs compared with controls without either disease (Glurich et al. 2002). A positive correlation between SAA level in serum and number of purulent periodontal pockets has been reported (Pussinen et al. 2004). Vuletic et al. (2009) reported that serum SAA concentration decreased after full-mouth extraction in periodontitis patients.

Increased serum levels of IL-1 β , TNF, OC, soluble intercellular adhesion molecule (sICAM), IL-6, MMP-9 have been reported in experimental studies in animal models as well as in clinical studies in humans (Raunio et al. 2007, Guentsch et al. 2009, Nakajima et al. 2010). However, Özçaka et al. (2010) failed to find significant differences between chronic periodontitis patients and healthy control subjects in terms of plasma OC and ICTP concentrations. In an intervention study, Marcaccini et al. (2009) reported that chronic periodontitis patients had higher plasma concentrations of MMP-3, -8, -9, and gelatinolytic activity, whereas non-surgical periodontal treatment decreased their levels significantly.

Serum cortisol concentration was reported to be associated with clinical periodontal parameters; PD and CAL (Ishisaka et al. 2008) and with BOP (Furugen et al. 2008). Furthermore, surfactant protein D was suggested to be a biomarker for chronic periodontitis, as it was found to be increased in 105 chronic periodontitis patients in comparison with 122 healthy control subjects (Glas et al. 2008). In a follow-up study, Amarasena et al. (2008) suggested that serum calcium level may be a risk factor for periodontal disease progression, as it correlated with PD. Higher platelet activating factor (PAF) levels were found in both GCF and serum of chronic periodontitis patients than gingivitis patients and healthy control subjects (Zheng et al. 2006, Chen et al. 2009). These elevated levels together with the significant correlations with clinical periodontal measurements suggested that PAF may have a role in the pathogenesis of periodontal destruction.

Conclusions

There exists extensive evidence that molecules in the saliva and in the GCF

correlate with tissue inflammation and bone destruction. Among those showing most promise is the ratio of RANKL/OPG, which is a discloser of periodontal disease activity but its utility in predicting future disease is questionable. However, the sensitivity and specificity of these molecules as predictors of future disease, i.e. prognosis, has not been reliably demonstrated. Even after the advent of highly sophisticated methods and almost 30 years of research on finding a consistent and definitive biochemical marker for periodontal disease activity and prognosis, no adequate marker has been successfully identified as yet. The natural history and nature of periodontal disease is one highly complicating factor in this situation as periodontal disease progresses episodically and it is very difficult to define the quiescent and active periods. Moreover, periodontal disease is a multi-factorial disease with several host- and microorganism-related factors acting consecutively. Last but not least, there is a wide range of individual variability in the response to the microbial triggers as well as to the therapeutic measures. Another issue making it difficult to come to a conclusion on the biochemical parameters is the diversity in biological sample collection, storage, and analysis methods used in different studies evaluating the same parameters. We should not be surprised that finding markers for periodontal disease is so elusive: other chronic conditions such as CVDs are similarly complex and despite even greater expenditure and manpower being applied, recent reports reveal that the previously lauded vascular risk factors such as CRP and N-BNP, are in fact not useful (Melander et al. 2009). Interestingly CRP and other risk factors in CVD are more useful as negative predictive factors similar to the use of BOP as a negative predictive factor in periodontal disease (Lang et al. 1986). At present, we do not have a clear understanding of the pathology or the molecular events occurring in the periodontal microenvironment during the tissue destruction process. Certain clinical measures such as clinical attachment loss remain the strongest predictor of future attachment loss and absence of certain clinical inflammatory signs such as BOP are excellent negative predictors of periodontal inflammation. Thus we can conclude that much work remains to identify molecules with clinical utility for estimating current and future

destructive periodontal disease activity. We provide the following supportable conclusions, many of which are not novel.

- Clinical measures such as PD, attachment level and BOP are essential for the diagnosis of prevalent periodontal tissue destruction.
- BOP remains to be the most reliable clinical finding correlated with periodontal disease activity.
- An increase in PD over time is associated with loss of attachment and loss of tooth supporting bone. Current PD is indicative of future attachment loss (Van der Velden et al. 2006). Thus, clinical periodontal measurements, mainly PD and BOP continue to be the most reliable parameters not only for the diagnosis but also for determining the prognosis of periodontal diseases, although they do not provide 100% accuracy.
- Investigating the levels of RANKL, OPG, and a range of host bone destruction markers and inflammation related molecules in GCF and saliva may prove to be reliable information on the state of periodontal disease activity however, currently cannot predict future disease activity.
- With the current picture, it is quite clear that highly specific and sensitive biomarkers for diagnosis and monitoring of periodontal diseases are still needed for early and better detection of periodontal tissue destruction.

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Clinical Relevance

Scientific rationale for the study: Objective and ideal diagnostic methods for periodontal diagnosis of the present as well as future tissue destruction are still being sought.

Principal findings: Clinical periodontal measurements, mainly PD

and BOP, continue to be the most reliable parameters not only for the diagnosis but also for determining the prognosis of periodontal diseases. A range of host bone destruction markers and inflammation related molecules in GCF, saliva and blood

are not reliable to predict future tissue destruction or disease activity. *Practical implications:* It is quite clear that highly specific, sensitive biomarkers for monitoring of periodontal diseases are still needed for better detection of periodontal tissue destruction.

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