Levels of common salivary protein 1 in healthy subjects and periodontal patients

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ABSTRACT

Purpose: Human saliva, as a vital part of the immune defense system, contains a number of distinct proteins and peptides. Recently human common salivary protein 1 (CSP1) has been identified as an abundant salivary protein and may play a role in promoting the binding of cariogenic bacteria to salivary pellicles. However, nothing else is known regarding the role of CSP1 in periodontology. The aim of this study was to quantify and compare CSP1 levels between healthy subjects and periodontal patients.

Methods: This controlled clinical study was conducted in periodontally healthy individuals and patients with chronic periodontitis Chonbuk National University Hospital, with Institutional Review Board approval. Whole saliva samples were collected from 36 healthy subjects and 33 chronic periodontitis patients and analyzed. Sodium dodecyl sulfate-polyacrylamide gel electrophoresis and immune blotting were conducted to ensure that anti-CSP1 monoclonal antibody (mAb) binds to CSP1 in human saliva. A sandwich enzyme-linked immunosorbent assay (ELISA) system was house-fabricated using mAb-hCSP1#14 and mAb-hCSP1#4 as a capture and a detector mAb, respectively. The CSP1 concentrations in saliva from 36 healthy subjects and 33 periodontal patients were quantified using the CSP1 sandwich ELISA system, and the results were analyzed using the Student’s t-test.

Results: Immunoblot analysis using mAb-hCSP1 as a probe confirmed that CSP1 in human saliva existed as a single band with a molecular weight of approximately 27-kDa. The quantification of CSP1 concentrations by CSP1 ELISA showed that the median values (25th to 75th percentiles) of periodontal patients and healthy subjects were 9,474 ng/mL (range, 8,434–10,139 ng/mL) and 8,598 ng/mL (range, 7,421–9,877 ng/mL), respectively. The Student’s t-test indicated the presence of a statistically significant difference between the 2 groups (P=0.024).

Conclusions: The presence of a significant difference in CSP1 levels between healthy subjects and periodontal patients suggests that CSP1 may be a potential biomarker for the detection or screening of periodontitis patients.

Keywords: Common salivary protein 1; Enzyme-linked immunosorbent assay; Monoclonal antibody; Periodontitis; Saliva
INTRODUCTION

Periodontitis is an inflammatory disease mainly caused by the growth of bacteria that attach to the surface of teeth [1]. Recently, various protein biomarkers in human saliva have been studied for the prevention and treatment of periodontal diseases [2]. Human saliva is a pivotal part of the immune defense system and contains roughly between 2,000 and 3,000 distinct salivary proteins, including alpha-amylase, lysozyme, mucins, proline-rich proteins, immunoglobulins, and many others [3,4]. Among the wide range of salivary proteins, human common salivary protein 1 (CSP1, synonyms: ZG16B, HRPE773, PRO1567) has been identified as a homolog of the demilune cell and parotid protein (Dcpp) in mice and of CSP1 in rats [5,6]. Kim et al. [7] recently reported that the pancreatic adenocarcinoma up-regulating factor gene has a high degree of homology to CSP1 and that this novel secretory protein is involved in the progression of pancreatic cancer, suggesting a strong connection between systemic disease and oral disease. In fact, some studies have provided support for the proposal that diabetes functions as a risk factor for periodontal diseases [8-10].

So far, to the best of our knowledge, only a single article regarding the function of CSP1 in the field of dental research has been published, demonstrating that CSP1 may play a role in promoting the binding of Streptococcus mutans (S. mutans) to experimental salivary pellicles formed on a hydroxyapatite surface, influencing the initial colonization of this pathogenic bacterium [11,12]. However, further functional information on CSP1 regarding periodontal pathogens remains to be elucidated. In this study, a house-fabricated sandwich enzyme-linked immunosorbent assay (ELISA) system was used as a key method for measuring CSP1 levels in human saliva. Using this method, the concentration of salivary CSP1 was compared between healthy subjects and periodontal patients. The findings of this study lead to the preliminary proposal of salivary CSP1 as a potential biomarker for the screening or monitoring of periodontal diseases.

MATERIALS AND METHODS

Study design and clinical measurements
A controlled clinical study was conducted among healthy subjects and periodontitis patients with Institutional Review Board approval (No. 201512012) at Chonbuk National University Hospital from March 2013 to March 2014. All participants between 20 and 70 years of age were assessed for eligibility for our study after a basic medical and dental examination. Participants were excluded if they had any known systemic diseases influencing their periodontal status, such as diabetes or cardiovascular disease. Although smoking history was recorded, smokers were not excluded. All clinical parameters, including bleeding on probing, pocket depth, and clinical attachment level, were measured by a periodontist. Periodontal diagnoses were made using the criteria defined at the international workshop for a classification of periodontal diseases and conditions [13]. An analysis of the previous literature showed that 16 samples each for the controls and cases were enough to ensure statistical significance for CSP1 measurements [14]. Therefore, 36 systemically and periodontally healthy controls and 33 periodontal disease patients signed a written informed consent form and were enrolled into the present cross-sectional study (Figure 1). The clinical characteristics of the study groups are presented in Table 1.
Saliva collection and storage

All participants were advised not to eat food or drink any beverages for at least 1 hour before sample collection. Saliva samples were collected in 50 mL polypropylene conical tubes and frozen at −70°C. When all the saliva samples were obtained, they were thawed at room temperature, divided into 1.5 mL Eppendorf tubes, and centrifuged at 13,000 g at 4°C for 15 minutes [15]. The supernatant was saved and transferred to new 1.5 mL tubes, and the samples were frozen at −70°C until they were ready to be used.

Figure 1. Flow diagram of the study. This controlled study was conducted in periodontally healthy subjects and in patients with chronic periodontitis. All participants between 20 and 70 years of age were asked to take part in the study, agreed, and signed a written consent form. After providing informed consent, patients received a periodontal examination. The periodontal examinations were used to determine whether each patient belonged to the healthy group or to the group of patients with periodontitis.

CSP1, common salivary protein 1; ELISA, enzyme-linked immunosorbent assay; n, number.

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**Table 1. Demographic characteristics of the study groups**

| Characteristics                  | Healthy group | Periodontal group | P value |
|----------------------------------|---------------|-------------------|---------|
| Total                            | 36 (52.2)     | 33 (47.8)         |         |
| Gender                           |               |                   | >0.100  |
| Male                             | 21 (58.3)     | 23 (69.7)         |         |
| Female                           | 15 (41.7)     | 10 (30.3)         |         |
| Smoking                          |               |                   | 0.540   |
| Smokers                          | 8 (22.2)      | 19 (57.6)         |         |
| Nonsmokers                       | 28 (77.8)     | 14 (42.4)         |         |
| Severity of periodontitis        |               |                   | 0.024   |
| Healthy                          | 36 (100)      | 0                 |         |
| Mild                             | 0             | 10 (30.3)         |         |
| Moderate to severe               | 0             | 23 (69.7)         |         |
| Age (yr)                         | 34.30±9.70    | 47.00±12.00       |         |
| PD (mm)                          | 0.70±0.06     | 4.46±1.72         | <0.001  |
| CAL (mm)                         | 1.40±0.16     | 4.25±2.51         | <0.001  |
| BOP (%)                          | 31.76         | 66.67             | <0.001  |

Values are presented as number of patients (%) or mean±standard deviation.

PD, pocket depth; CAL, clinical attachment level; BOP, bleeding on probing.
**Sodium dodecyl sulfate-polyacrylamide gel electrophoresis and immunoblotting**

Proteins were denatured by boiling for 1.5 minutes after adding 2× sodium dodecyl sulfate (SDS) treatment buffer and then were separated on a 12% SDS-polyacrylamide gel. Proteins on the gels were either stained with Coomassie blue or electro-transferred to a polyvinylidene membrane for immunoblotting [15]. The blot was blocked with 5% nonfat powdered milk in phosphate-buffered saline buffer with 0.2% Tween 20 (PBST) and probed with biotin-conjugated monoclonal antibody (mAb) to CSP1 (mAb-hCSP1#4) for 2 hours at room temperature [13]. After the membrane was thoroughly rinsed with PBST buffer, the blot was visualized using streptavidin conjugated to horseradish peroxidase (HRP) and an enhanced chemiluminescence substrate kit (Pierce, Rockford, IL, USA). CSP1 protein was detected by exposure of the blot to Biomax-MS X-ray film (Eastman Kodak, Rochester, NY, USA) in a dark room.

**Standard curve for ELISA**

A standard curve for CSP1 levels in human saliva using a sandwich ELISA system was obtained with different concentrations of recombinant hCSP1 (0–14,000 ng/mL). In the standard curve, the optical density value at 450 nm was plotted on the y-axis against the hCSP1 concentration on the x-axis. In order to detect hCSP1 as a standard protein, mAb-hCSP1#14 and mAb-hCSP1#4 were used as the capture mAb and the detector mAb, respectively. A reliable Pearson correlation coefficient value ($r^2=0.983$) was observed between the 2 parameters, indicating good linearity throughout the entire measurement range (data not shown).

**Sandwich ELISA**

Sandwich ELISA was used in this study to measure the concentration of CSP1 in saliva. In brief, 50 µL of capture mAb (mAb-hCSP1#14, 2 µg/mL) was applied to the ELISA plate, and the plate was incubated for 1 hour at 37°C and washed 3 times with PBST. After the plate was treated with 50 µL of bovine serum albumin blocking solution (10 mg/mL), 50 µL of the serum was added to the wells, and the plate was incubated for 1 hour at 37°C. After washing the plates with PBST, 50 µL of the biotin-conjugated detector mAb (mAb-hCSP1#4, 2 µg/mL) was added to the wells and incubated for 1 hour at 37°C prior to being probed with an avidin-HRP solution (10 mg/mL). Following a final rinse with PBST, the color reaction for the detection of antigen-mAb complex was initiated with the addition of 50 µL of 3,3',5,5'-tetramethylbenzidine substrate solution (10 mg/mL) for 15 minutes and then stopped by adding 50 µL of 10 µmol/mL H$_2$SO$_4$. The absorbance was measured at 450 nm in an automatic ELISA reader (Model 550, Bio-Rad, Hercules, CA, USA).

**Data analysis**

To determine the normality of the distribution of the sample, the D’Agostino and Pearson omnibus normality test was performed using PRISM version 6 (GraphPad Software Inc., La Jolla, CA, USA). PASW Statistics for Windows version 18.0 (SPSS Inc., Chicago, IL, USA) and Microsoft Excel 2010 (Microsoft Corp., Redmond, WA, USA) were used to analyze and compare the test results. $P$-values <0.05 were considered to indicate statistical significance. Pearson correlation coefficients ($r^2$) were used to evaluate correlations between 2 parameters.
RESULTS

Clinical characteristics and diagnosis
Among the 106 volunteers whose basic medical and dental conditions were examined, 37 individuals did not meet the proposed inclusion criteria and were not eligible for this controlled clinical study. The exclusion criteria included acquired immunodeficiency syndrome, diabetes, high blood pressure, pregnancy, breastfeeding, and use of antibiotics over the last 3 months. After comprehensive periodontal examinations, 69 individuals were allocated into 2 groups: 36 healthy subjects and 33 periodontal patients (Figure 1). Among the 33 periodontal disease patients, 10 patients were diagnosed with mild periodontitis and 23 patients with moderate to severe periodontitis according to the criteria defined at the international workshop for a classification of periodontal diseases and conditions (Table 1).

Immunoblotting
In order to test the binding affinity and quality of anti-hCSP1 mAbs, human saliva proteins were resolved using SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and electro-transferred to nitrocellulose paper. The blot was then probed with mAb-hCSP1#4 for an immune reaction. As shown in lanes 1 and 2 of the western blot in Figure 2, the mAb detected a 46-kDa band corresponding to glutathione-S-transferase (GST)-tagged recombinant human CSP1 and a single band at approximately 27-kDa in the human saliva sample (Figure 2). When the blot was probed with mAb-hCSP1#14, a similar result was obtained (data not shown).

Measurement of CSP1 in saliva by ELISA
Saliva CSP1 concentrations of 36 healthy subjects and 33 periodontal patients were quantified based on the standard curve generated by the house-fabricated sandwich ELISA system. Figure 3 shows the profiles of saliva CSP1 concentration in the healthy group and periodontal group. The profiles demonstrated that the saliva of periodontitis patients had higher CSP1 levels than that of healthy subjects. The median values (25th to 75th percentiles) of the

![Figure 2. Detection of CSP1 in human saliva with a mAb. CB-stained sodium dodecyl sulfate-polyacrylamide gel electrophoresis demonstrates molecular size markers (lane M), purified GST-tagged recombinant hCSP1 as a positive control (lane 1), and human saliva (lane 2). The WB is an immunoblot analysis of the CB probed with primary mAb-hCSP1#4. mAb recognized a single protein band at approximately 27-kDa in the blot. CSP1, common salivary protein 1; mAb, monoclonal antibody; CB, Coomassie blue; GST, glutathione S-transferase; WB, western blot.](https://doi.org/10.5051/jpis.2016.46.5.320)
Healthy group and periodontal group were 8,598 ng/mL (range, 7,421–9,877 ng/mL) and 9,474 ng/mL (range, 8,434–10,139 ng/mL), respectively. The Student’s t-test results showed statistically significant differences between the 2 groups (P=0.024). The analysis of CSP1 levels in the 2 groups is also summarized in Table 2.

DISCUSSION

Saliva is thought to act as an indicator of oral disease, including periodontitis. However, our knowledges of whether and which salivary proteins are affected by the processes of periodontal disease remain limited. In this study, the subjects were divided into 2 groups, periodontally healthy individuals and periodontitis patients, after a comprehensive periodontal examination (Figure 1, Table 1). Then, saliva levels of human CSP1, a major salivary protein, were measured in both healthy individuals and periodontitis patients. CSP1 levels were measured using a custom-fabricated sandwich ELISA system, and were compared and analyzed (Table 2, Figure 3). As a capture antibody and a detector antibody, mAb-hCSP1#14 and mAb-hCSP1#4 were tested for their binding capacity by immunoblotting with human saliva (Figure 2) and evaluated using the Pearson correlation coefficient with the standard curve.

Table 2. CSP1 levels in the healthy and periodontal groups

| Characteristics               | Healthy group | Periodontal group |
|-------------------------------|---------------|-------------------|
| No. of subjects               | 36            | 33                |
| Mean (ng/mL)                  | 8,598         | 9,474             |
| Standard deviation            | 1,932         | 1,290             |
| Standard error of the mean    | 322.1         | 224.6             |
| Median (ng/mL)                | 9,474         | 8,598             |
| 25th to 75th percentiles (ng/mL) | 8,434–10,139  | 7,421–9,877       |

CSP1, common salivary protein 1.
SDS-PAGE and western blot analysis showed that the molecular weight of recombinant hCSP1 tagged with GST was approximately 46-kDa, as indicated in lane 1 of Figure 2. Since the molecular weight of GST is known to be 26-kDa, the molecular weight of CSP1 was estimated to be approximately 20-kDa. However, mAb-hCSP1#4 recognized only a single band at approximately 27-kDa in human saliva, as shown in lane 2 of the western blot in Figure 2, which is somewhat larger than was predicted from the CSP1 gene sequence. This result is consistent with previously reported data [11,15]. Ambatipudi et al. [11] explained this discrepancy by noting that the human CSP1 sequence contains one potential N-glycosylation site, and thus human CSP1 may exist in a glycosylated form in saliva. Dcpp in mice, a homolog of human CSP1, has 3 forms encoded by different genes [5], but rat CSP1 is encoded by only 1 gene, as in humans. In the case of rats, interestingly, 3 CSP1 isoforms with molecular weights of 16-kDa, 20-kDa, and 22-kDa, were identified in the salivary glands, suggesting that different isoforms of CSP1 may have been derived from N-glycosylation or phosphorylation of the CSP1 protein [6].

In the present study, the t-test was used to examine whether the mean differences between the 2 groups were statistically significant because the values of CSP1 concentrations exhibited a normal distribution. The means (standard deviations) of saliva CSP1 concentrations in 36 healthy subjects and 33 periodontal patients were 8,598 ng/mL (1,932 ng/mL) and 9,474 ng/mL (1,290 ng/mL), respectively (Table 2, Figure 3), showing that the difference in CSP1 levels between the 2 groups was statistically significant (P=0.024). No statistically significant intragroup differences in CSP1 levels were found, for example, smoking vs. non-smoking (P=0.400) or mild vs. moderate to severe periodontitis (P=0.800) in periodontal patients or smoking vs. non-smoking (P=0.200) in healthy subjects. At this point, the reason why CSP1 levels were higher in periodontal patients than in normal healthy subjects remains unknown.

One possible explanation for the presence of a higher level of CSP1 in periodontal patients than in healthy subjects can be deduced from the presence of a putative jacalin-related lectin (JRL) domain in human CSP1. Members of the JRL protein family are known to bind to glycoproteins, are ubiquitously expressed throughout the plant and animal kingdoms, and carry out functions such as cell agglutination and antimicrobial activity [5,16]. Thus, CSP1 may play a role in modulating the composition of oral microflora, similarly to other salivary proteins such as lysozyme and lactoperoxidase, by binding pathogenic bacteria [17] and causing aggregation and direct killing. Although rat CSP1 includes a JRL domain, it did not show any detectable effects on cell agglutination and antimicrobial activity, but rather led to enhanced bacterial adherence when S. mutans were exposed to rat CSP1 in saliva [12]. Human CSP1, similarly to amylase, may encourage pathogenic bacteria to bind to the tooth surface and act as bacterial anchor sites of initial pathogen tooth colonizers [18], suggesting that the presence of other salivary proteins may mediate the attachment of CSP1 to bacterial surfaces or between the bacteria and salivary pellicles, possibly by forming nanostructure protein complexes. Thus, another explanation for the higher level of CSP1 in periodontal patients than in healthy subjects may be related to the higher stress level associated with chronic diseases. As several recent studies have shown that that CSP1 was highly expressed in stressed conditions, including periodontitis, pancreatic cancer [7], and diabetes mellitus [14], chronic stress may act as a risk factor for higher CSP1 levels in chronic periodontitis. Whether chronic stress directly affects the level of CSP1 remains an intriguing question to be resolved.

Most importantly, the present study showed that periodontitis patients had a significantly higher level of salivary CSP1 than healthy subjects. Our findings may suggest that CSP1 could
be used as a potential biomarker for the detection or screening of periodontal patients. Due to the advantages of saliva, including non-invasiveness, safety, and simplicity, early diagnosis and treatment of periodontal disease may become more accessible [19]. As a future strategy, a much larger randomized controlled cohort study [20] is required.

In conclusion, human CSP1 was identified in saliva as a single 27-kDa band by probing with mAb-hCSP1#4 and immunoblotting. The mean (standard deviation) CSP1 levels in healthy controls and periodontal patients were 8,598 ng/mL (1,932 ng/mL) and 9,474 ng/mL (1,290 ng/mL), respectively, and a statistically significant difference in CSP1 levels between the 2 groups (P=0.024) was observed using a sandwich ELISA system. These results indicate that CSP1 may be used in the future as a potential biomarker for the detection and screening of periodontal patients.

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REFERENCES

1. Savage A, Eaton KA, Moles DR, Needleman I. A systematic review of definitions of periodontitis and methods that have been used to identify this disease. J Clin Periodontol 2009;36:458-67. P UBMED | C R O S S R E F
2. Baser U, Oztekin G, Ademoglu E, Isik G, Yalcin F. Is the severity of periodontitis related to gingival crevicular fluid and serum high-sensitivity C-reactive protein concentrations? Clin Lab 2014;60:1653-8. P UBMED
3. Wilmarth PA, Riviere MA, Rustvold DL, Lauten JD, Madden TE, David LL. Two-dimensional liquid chromatography study of the human whole saliva proteome. J Proteome Res 2004;3:1017-23. P UBMED | C R O S S R E F
4. Yao Y, Berg EA, Costello CE, Troxler RF, Oppenheim FG. Identification of protein components in human acquired enamel pellicle and whole saliva using novel proteomics approaches. J Biol Chem 2003;278:5800-8. P UBMED | C R O S S R E F
5. Mullins JJ, Mullins LJ, Dunbar DR, Brammar WJ, Gross KW, Morley SD. Identification of a human ortholog of the mouse Dcpp gene locus, encoding a novel member of the CSP-1/Dcpp salivary protein family. Physiol Genomics 2006;28:129-40. P UBMED | C R O S S R E F
6. Girard LR, Castle AM, Hand AR, Castle JD, Mirels L. Characterization of common salivary protein 1, a product of rat submandibular, sublingual, and parotid glands. J Biol Chem 1993;268:26592-601. P UBMED
7. Kim SA, Lee Y, Jung DE, Park KH, Park JY, Gang J, et al. Pancreatic adenocarcinoma up-regulated factor (PAUF), a novel up-regulated secretory protein in pancreatic ductal adenocarcinoma. Cancer Sci 2009;100:828-36. P UBMED | C R O S S R E F
8. D’Aiuto F, Parkar M, Andreou G, Suvan J, Brett PM, Ready D, et al. Periodontitis and systemic inflammation: control of the local infection is associated with a reduction in serum inflammatory markers. J Dent Res 2004;83:156-60. P UBMED | C R O S S R E F
9. Mealey BL, Oates TW. American Academy of Periodontology. Diabetes mellitus and periodontal diseases. J Periodontol 2006;77:1289-303. P UBMED | C R O S S R E F
10. Wu YY, Xiao E, Graves DT. Diabetes mellitus related bone metabolism and periodontal disease. Int J Oral Sci 2015;7:63-72. P UBMED | C R O S S R E F

http://jpis.org https://doi.org/10.5051/jpis.2016.46.5.320 327
11. Ambatipudi KS, Hagen FK, Delahunty CM, Han X, Shafi R, Hryhorenko J, et al. Human common salivary protein 1 (CSP-1) promotes binding of Streptococcus mutans to experimental salivary pellicle and glucans formed on hydroxyapatite surface. J Proteome Res 2010;9:6605-14.

12. Koo H, Vacca Smith AM, Bowen WH, Rosalen PL, Cury JA, Park YK. Effects of Apis mellifera propolis on the activities of streptococcal glucosyltransferases in solution and adsorbed onto saliva-coated hydroxyapatite. Caries Res 2000;34:418-26.

13. Armitage GC. Development of a classification system for periodontal diseases and conditions. Ann Periodontol 1999;4:1-6.

14. Wang H, Heo SM, Jin HY, Choi EY, Oh SW. Common salivary protein 1 in serum of diabetes patients. J Clin Lab Anal. Forthcoming 2016.

15. Towbin H, Staehelin T, Gordon J. Electrophoretic transfer of proteins from polyacrylamide gels to nitrocellulose sheets: procedure and some applications. Proc Natl Acad Sci U S A 1979;76:4350-4.

16. Raval S, Gowda SB, Singh DD, Chandra NR. A database analysis of jacalin-like lectins: sequence-structure-function relationships. Glycobiology 2004;14:1247-63.

17. Marsh PD. Host defense and microbial homeostasis: role of microbial interactions. J Dent Res 1989;68:1567-75.

18. Nobbs AH, Lamont RJ, Jenkinson HF. Streptococcus adherence and colonization. Microbiol Mol Biol Rev 2009;73:407-50.

19. Kaufman E, Lamster IB. The diagnostic applications of saliva--a review. Crit Rev Oral Biol Med 2002;13:197-212.

20. Hansen TV, Simonsen MK, Nielsen FC, Hundrup YA. Collection of blood, saliva, and buccal cell samples in a pilot study on the Danish nurse cohort: comparison of the response rate and quality of genomic DNA. Cancer Epidemiol Biomarkers Prev 2007;16:2072-6.