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ABSTRACT

Understanding the composition and function of the acquired enamel pellicle (AEP) has been a major goal in oral biology. The aim of this study was to test the hypothesis that intact histatins are part of the *in vivo* AEP and that histatins after adsorption to HA have effects on *in vitro* enamel demineralization. This is the first study demonstrating the presence of intact histatins *in vivo* in the AEP. The *in vitro* experiments show that all naturally occurring histatins in the AEP have the potential to provide some level of protection against acid injury.

KEY WORDS: histatin, proteomics, enamel pellicle, demineralization, saliva.

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Evidence of Intact Histatins in the *in vivo* Acquired Enamel Pellicle

INTRODUCTION

Histatins are members of a family of low-molecular-weight salivary pro-teins secreted by the major and minor salivary glands (Baum *et al.*, 1977; Oppenheim *et al.*, 1986; Siqueira *et al.*, 2008). Like other salivary proteins, histatins appear to be multifunctional, and their major property is their antifungal activity against the opportunistic yeast, *Candida albicans* (Oppenheim *et al.,* 1988). Besides antifungal properties, antibacterial effects have been attributed to histatins, based on their killing and growth-inhibitory activity against several species of oral bacteria (Pollock *et al.*, 1984; Xu *et al.*, 1991). Based on *in vitro* hydroxyapatite adsorption studies, histatin 1, histatin 3, and histatin 5 have also been considered to be protein components of the acquired enamel pellicle (AEP). Recent investigations with state-of-the-art proteomics focusing on the protein composition of *in vivo* AEP have revealed the presence of protein fragments originating from these histatins (Siqueira *et al.*, 2007b; Siqueira and Oppenheim, 2009). These protein fragments are generated due to the high proteolytic activity present in oral fluid (Helmerhorst *et al*., 2006). Despite the harsh environment of the oral cavity, it is apparent that some of the secreted histatin 1, histatin 3, and histatin 5 could survive and thus could become, after their adsorption to the enamel surface, constituents of the composition and structure of the *in vivo-*formed AEP. The aim of this study was to investigate whether intact histatins can be detected in *in vivo-*formed AEP. In addition, we assessed, with *in vitro* experiments, whether histatins after adsorption to human enamel surfaces could exert a functional effect in protection against demineralization.

Materials & Methods

Human Participants

Saliva and AEP samples were obtained from healthy male and female volunteers, ranging in age from 24 to 40 yrs. Saliva, human teeth, and AEP collection protocols were approved by the Institutional Review Board of Boston University Medical Center, and informed consent was obtained from each person participating in the study.

Collections of Glandular Secretion and Whole Saliva

Stimulated parotid secretion from both glands was collected with the use of 3% citric acid and a Carlson-Crittenden device. Whole-saliva collection was carried out under masticatory stimulation with parafilm. The samples were kept on ice during the collection procedure. The mean flow rates for parotid secretion and whole saliva were 0.9 mL/min and 1.2 mL/min, respectively. Immediately after collection, whole-saliva samples were centrifuged at 14,000 x *g* for 20 min at 4°C. Whole-saliva supernatants were separated from the pellet and used immediately for electrophoretic analysis. The total protein concentration was measured by the bicinchoninic acid (BCA) assay (Pierce Chemical, Rockford, IL, USA) and bovine serum albumin protein standard. The mean protein concentration in parotid secretion was 0.84 mg/mL, while that in supernatant whole saliva was 1.5 mg/mL.

Harvesting of Human AEP

AEP material was harvested from the enamel surface as described previously (Siqueira and Oppenheim, 2009). Briefly, *in vivo* AEP formed naturally on the enamel surface over a twohour period was collected by means of a collection strip with the dimensions of 0.5 x 1.0 cm (Electrode wick filter paper; BIO-RAD, Hercules, CA, USA) pre-soaked in 3% citric acid. After collection, the collection strips containing AEP were placed in a microcentrifuge tube and kept frozen at -20°C until used for elution and electrophoresis.

Cationic PAGE

Cationic PAGE was performed as described previously (Oppenheim *et al.*, 1986). To minimize losses of AEP proteins, we applied a modified sample-handling method as described (Siqueira *et al.*, 2007a). Briefly, a 35-µL quantity of sample buffer was added to a microcentrifuge tube containing 4 collection strips. The tubes were vortexed for 5 sec at maximum speed, and the 4 strips were placed directly in the wells of the stacking gel, thus avoiding any intermediate elution steps. A constant voltage of 120 V was used for the development of electrophoretic separation, and the gel was stained with silver stain.

Protein Extraction

Protein zones of interest were removed by means of a razor blade, and each gel band was further cut into small pieces. Proteins were extracted from the gel pieces as described (Cohen and Chait, 1997), with a solution containing formic acid/ water/2-propanol in a 1:3:2 v/v/v ratio. An aliquot of 200 µL of this elution medium was added to the microcentrifuge tube containing the gel fragments, and the tube was subjected to shaking for 20 hrs at 4°C. Gel pieces and supernatant were separated by centrifugation, and the pellet was extracted twice after one-hour periods of shaking. The resulting 600 µL of supernatant were dried in a speed vac, and this residue was utilized for mass spectrometric analysis.

Mass Spectrometric Analysis

To determine the molecular weight of the protein bands harvested from Cationic PAGE, we used a Voyager DE-Pro MALDI-TOF mass spectrometer (Applied Biosystems, Foster City, CA, USA) in the reflector mode, 25 kV accelerating voltage, delayed extraction, position ion mode, and close external calibration. The matrix used consisted of alpha-cyano-4 hydroxy-cinnamic acid.

Enamel Demineralization Assay

To evaluate the effects of full-length histatins on the *in vitro* demineralization of human enamel, we isolated native histatin 1, histatin 3, and histatin 5 from human glandular secretions and obtained synthetic histatin 1, lacking phosphate, and synthetic histatin 3, phosphorylated at residue 2, commercially (American Peptide Company, Sunnyvale, CA, USA). Protein solutions were prepared in 50 mM NaCl, pH 7.0, at a protein concentration of 1 mg/mL.

Human permanent first molars were selected and cleaned with a 5-micron alumina suspension with the use of a prophy cup and rinsed with distilled water. The crowns were sectioned into 2 parts, each of which was cut perpendicular to the surface with a diamond-coated band saw, yielding 4 sections of approximately 300 μ m thickness. Subsequently, these sections were ground to a uniform thickness of 180 µm. All surfaces of each enamel section, except a 2-mm-long zone of the natural surface enamel (lingual or buccal surface), were then coated with nail varnish. A gold finder grid used for electron microscopy was buried in the nail varnish close to the uncoated area to help orient the microradiographic images on the computer before and after the specific treatment.

The demineralization was carried out with a total of 30 enamel sections, which were randomly divided into 6 groups $(n = 5 \text{ per group})$. Each specimen was exposed to 100 μ L of protein solution (100 μ g specific protein/100 μ L) for a period of 2 hrs at 37°C with gentle agitation, and the control group specimens were exposed to 100 µL of 50 mM NaCl containing no protein. After this period, the specimens were washed in distilled water. Subsequently, the specimens were immersed in individual tubes containing 2 mL of demineralization solution $(2.2 \text{ mM } CaCl₂;$ 2.2 mM NaH₂PO₄; 0.05 M acetic acid; pH 4.5) for a period of 12 days at 37ºC. Immediately after the demineralization period, the specimens were extensively washed in distilled water and dried with filter paper. We used 2 mL of demineralization solution to determine the calcium and phosphate concentrations released from the enamel specimens during the demineralization procedure, and we further used the enamel specimens to evaluate mineral loss and lesion depth by comparing the microradiographs before and after exposure to acidic conditions.

Calcium and Phosphate Analyses

The calcium content of the samples was analyzed by atomic absorption spectrophotometry (Model 3300, Perkin Elmer Analytical Instruments, Norwalk, CT, USA) at a wavelength of 422.7 nm, with the addition of 1% LaCl₃ to prevent phosphate interference. The phosphate concentration was measured spectrophotometrically as described previously (Margolis and Moreno, 1985), with a UV-visible spectrophotometer (Cary 50 Scan, Varian, Lake Forest, CA, USA) for determination of the O.D. at a wavelength of 390 nm. All samples were analyzed in triplicate.

Figure 1. Cationic PAGE with samples applied to the stacking gel trough after the strip was dipped into the protein solution. Lane 1, parotid secretion; Lane 2, whole saliva; Lane 3, AEP material from 8 collection strips; Lane 4, 0.125 µg each of native histatins 1, 3, and 5.

Mineral Loss Determination

The loss of mineral content was assessed by quantitative transverse microradiography (Margolis *et al.*, 1999). Enamel section microradiographs were obtained with the use of a Faxitron x-ray system (Hewlett Packard, McMinnville, OR, USA) and highspeed holographic film before and after demineralization treatment. Images were analyzed by means of a video image system and software developed by Ed Marcus Laboratories (Newton, MA, USA). The gold grid attached to the cut surface of each section and the outer enamel edge was used to align the microradiographs obtained for each section prior to and after exposure to demineralization solution. The area density between the mineral content profile acquired before demineralization and the corresponding density obtained after demineralization were integrated, and the differences were established, yielding the numerical value of ΔZ.

Statistical Analyses

Statistical procedures were performed with the software package Minitab 13.1. After checking for normal distribution, we subjected the data to analysis of variance (ANOVA) followed by Tukey's test. The level of significance was set at a value of $p < 0.05$.

RESULTS

The silver-stained cationic PAGE of AEP formed over 2 hrs showed 3 major bands in the middle to lower regions of the AEP lane (Fig. 1, lane 3), while the lane corresponding to whole saliva showed the absence of these 3 major bands (Fig. 1, lane 2). The bands from lane 3 exhibited electrophoretic mobilities similar to those of standard native histatin 1, histatin 3, and histatin 5 (Fig. 1, lane 4), as well as those of native histatin 1, histatin 3, and histatin 5 present in parotid secretion (Fig. 1, lane 1). For further exploration of the identity of these proteins, the 3

Table. Micrographic Density Measurements of Enamel Sections after Exposure to Demineralization Conditions with Prior Coating with Specific Histatins and Control Solution (n = *5 per* group)

		ΔZ (vol % x μ m) ^{1,2} Lesion Depth (μ m) ²
Histatin 1	$1.520 + 530^{\circ}$	$.55 + 11^{\circ}$
Unphosphorylated histatin 1	$3326 \pm 646^{\circ}$	$130 + 23$ °
Historin ₃	$3737 + 931^{\circ}$	$86 + 13^{b}$
Phosphorylated histatin 3	1627 ± 555 ^o	$69 + 20^{b}$
Histatin ₅	4128 ± 718 ^b	$94 + 12^{b}$
Control	7354 ±1258°	$1.50 + 14$ ^c

 ΔZ = numerical value for mineral loss due to demineralization.

Different superscripts indicate statistical difference, and the same superscripts indicate no statistical difference, according to Tukey's test among proteins and control.

major bands from lane 3 were excised from the gel, and the contained proteins were eluted and subjected to MALDI-TOF analysis. The mass spectrometry results obtained are shown in Fig. 2. The band with the slowest electrophoretic mobility contained a protein with an m/z of 4933.60, consistent with the molecular-weight (Mr) value of histatin 1. The middle band showed a predominant protein with an m/z of 4062.21, consistent with the M^r value of 4062.5 Da of histatin 3, and the fastestmoving band contained a protein with an m/z of 3036.45, consistent with the M^r value of 3036.7 Da of histatin 5.

To gain insight into the biological functions of these histatins when adsorbed onto the enamel surface, we evaluated their effects on the demineralization of enamel *in vitro*. To evaluate the role of covalently linked phosphate, we included 2 synthetic proteins (histatin 1 lacking phosphate and synthetic histatin 3 phosphorylated at residue 2) in the analyses. The amounts of phosphate dissolved after demineralization at pH 4.5 are shown in Fig. 3A. The control group, not treated with any protein, showed the most drastic phosphate loss. All specimens treated with histatins demonstrated reduction in phosphate loss. Histatin 1 and phosphorylated histatin 3 treated specimens both demonstrated the highest inhibition of phosphate loss. Unphosphorylated histatins showed an intermediate degree of phosphate loss when compared with control and phosphorylated histatins.

Similar results were obtained in relation to the release of calcium. Histatin 1 and phosphorylated histatin 3 treated specimens showed the lowest calcium loss. Sections exposed to unphosphorylated histatins (synthetic histatin 1, histatin 3, and histatin 5) demonstrated more calcium release, but their calcium loss was significantly lower than that observed with the control group (Fig. 3B).

The average values of densitometric mineral loss and lesion depth after acidic treatment with demineralization solution are presented in the Table. As observed with ionic calcium and phosphate release measurements, enamel sections exposed to histatin 1 and synthetic phosphorylated histatin 3 showed a level of protection significantly higher than that observed for the other groups. Mineral losses with specimens treated with unphosphorylated histatin 1, histatin 3, and histatin 5 were significant lower than those of the control group ($p < 0.05$).

Figure 2. Mass spectrum of protein eluted from an excised gel piece containing the protein band of *in vivo* AEP exhibiting the same electrophoretic mobility as histatin 1 standard **(A).** Mass spectrum of protein eluted from an excised gel piece containing the protein band of *in vivo* AEP exhibiting the same electrophoretic mobility as histatin 3 standard **(B).** Mass spectrum of protein eluted from an excised gel piece containing the protein band of *in vivo* AEP exhibiting the same electrophoretic mobility as histatin 5 standard **(C).**

DISCUSSION

This is the first study demonstrating the presence of intact histatins in *in vivo-*formed AEP. This observation is of interest in view of the fact that the life-span of full-length histatins in the whole-saliva environment is very short (Helmerhorst *et al.*, 2006). This is due to the high sensitivity of these proteins to

Figure 3. Phosphate **(A)** and calcium **(B)** released from human enamel sections first exposed to specific histatin preparations or control buffer followed by exposure to demineralizing solution. Ordinates: Ion composition of demineralizing solution after 12 days' exposure. Abscissae: Columns indicating protein preparations and controls used in enamel window treatment for 2 hrs (n = 5 *per* group). Different superscripts indicate statistical difference, and the same superscripts indicate no statistical difference, according to Tukey's test among proteins and control. Bars represent standard deviation of the mean.

whole-saliva-associated proteases (Castagnola *et al.*, 2004; Hardt *et al.*, 2005). It thus appears that binding of intact histatins to the enamel surface occurs prior to their proteolytic degradation, and that binding to the mineral exerts a protective effect against further enzymatic proteolytic degradation.

From a functional point of view, the identification of intact histatin 1, histatin 3, and histatin 5 in the *in vivo* AEP is of considerable interest, considering the various biological properties of histatins described so far. Histatins are components of the non-immune or innate oral host defense system, able to kill pathogenic yeasts (Helmerhorst *et al.*, 2005) and exhibit microbicidal activity against a variety of oral bacteria (Gusman *et al.*, 2001). Their presence in an intact form in the AEP would suggest that they could exhibit at least some of their known biological functions and could provide additional biological properties specific for the role AEP plays *in vivo*. The *in vitro* results obtained in this study showed that all 3 naturally occurring histatins in AEP have the potential to provide some level of protection against acid injury.

It should be pointed out that the adsorption behavior of a single protein to hydroxyapatite may show differences when compared with its adsorption in the presence of more than one polypeptide, as must be the case in the *in vivo* formation of AEP. Studies focusing on such differences with histatins have been carried out in the presence of other phosphoproteins, and the use of binary and ternary systems has indeed shown differences in the adsorption behavior of histatins (Yin *et al.*, 2003, 2006). No studies have as yet been carried out to address the functional effect of histatin in such multi-component protein films. The single-component functional consequences after adsorption represent the first step in gaining insights into the possible role of the 3 major histatins in enamel protection. A clear functional difference was found in this study between phosphorylated and unphosphorylated histatins. Analysis of the data provides evidence that the presence of phosphoserine at position 2, whether present in native histatin or the synthetic histatin 3, showed the highest degree of protection against acid demineralization. This is consistent with previous adsorption data which demonstrated that native histatin 1 shows a higher hydroxyapatite adsorption affinity than recombinant histatin 1, lacking covalently linked phosphate at position 2 (Driscoll *et al.*, 1995). The protection with phosphorylated histatins was higher than that observed with the protection gained with the unphosphorylated histatins. The latter group still reduced demineralization when compared with the control. These findings indicate that the phosphorylation of serine residue 2 is not the only determining factor in enamel/hydroxyapatite binding, and other residues must be contributing to the selective adsorption of histatins to hydroxyapatite. The precise mechanisms of how phosphorylated and unphosphorylated histatins reduce mineral dissolution remain to be elucidated. The ultimate goal is to understand how the entire population of specific polypeptides constituting the *in vivo*formed AEP affects the mineral homeostasis of enamel surfaces.

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