Saliva-Binding Region of *Streptococcus mutans* Surface Protein Antigen

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A 190-kDa surface protein antigen (PAc) of Streptococcus mutans binds to human salivary components. For detection of specific binding of the PAc protein to human salivary components, a simple sandwich assay was used. Microtiter plates precoated with recombinant PAc (rPAc), PAc fragments, or S. mutans whole cells were allowed to react with human whole saliva and then were incubated with biotinylated rPAc. The biotinylated rPAc bound to salivary components was detected by use of alkaline phosphatase-conjugated streptavidin and p-nitrophenylphosphate. In this assay, the binding of whole cells of S. mutans and purified rPAc to salivary components was confirmed. For determination of a saliva-binding region of the PAc molecule, 14 truncated PAc fragments were constructed by use of the polymerase chain reaction and an expression vector, pAX4a+. The binding of these truncated PAc fragments to human salivary components was determined by the sandwich assay. Among the truncated PAc fragments, fragments corresponding to residues 39 to 864 and residues 39 to 1000 of PAc showed a high ability to bind to salivary components. Shorter recombinant fragments corresponding to residues 39 to 217, residues 200 to 481, residues 470 to 749, and residues 688 to 864 did not exhibit any binding ability. The fragment that corresponds to a proline-rich repeating region (residues 828 to 1000) bound directly to the PAc protein. These results suggest that residues 39 to 864 of the PAc molecule are important in the binding of the surface protein to human salivary components, and the proline-rich repeating region of the PAc protein may contribute to spontaneous self-aggregation of the PAc protein.

Dental caries is one of the most common diseases in humans. Among numerous members of the oral flora, *Streptococcus mutans* is considered to be a major causative agent of human dental caries (11, 20). A 190-kDa protein antigen, PAc, which is also known as I/II (28), B (30), P1 (8), or MSL-1 (5), has been given much attention as a possible dental caries vaccine. Subcutaneous immunization of monkeys with purified PAc (I/II or B) (18, 19, 31) and local, passive immunization of monkeys (17) and humans (21) with anti-PAc (I/II) monoclonal antibodies (MAbs) were found to confer significant protection against dental caries or inhibition of *S. mutans* colonization.

The molecular structure of PAc (P1) of S. mutans serotype c was clarified by cloning (15, 24) and sequencing (12, 25) of the gene coding for this protein. The PAc protein possesses two internal repeating amino acid sequences; one is rich in alanine and is located in the N-terminal region (A region), and the other is rich in proline and is located in the central region (P region). The biological function of the PAc protein remains unclear. However, there is suggestive evidence that PAc contributes to the salivary agglutinin-mediated aggregation and adherence of S. mutans. It is well known that whole cells of S. mutans aggregate in the presence of human whole saliva or salivary agglutinin (5, 7, 9). Koga et al. (13) and Lee et al. (16) have constructed isogenic mutants of S. mutans deficient in PAc and have revealed that the PAc-defective mutants do not aggregate in the presence of whole saliva or salivary agglutinin. Ogier et al. (22, 23) have cloned and sequenced the gene coding for a saliva receptor protein of S. mutans serotype f and have found that the molecular struc-

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ture of the saliva receptor protein is very similar to that of the PAc protein. In addition, Russell and Mansson-Rahemtulla (29) have shown that salivary components with molecular weights of 28,000 and 38,000 are capable of binding to the PAc (I/II) protein. Brady et al. (3) and Demuth et al. (5) have reported that PAc (P1, MSL-1) inhibits the salivary agglutinin-induced aggregation and adherence of S. *mutans*. These findings indicate that the PAc protein is capable of binding to salivary components.

In the present study, we used a sandwich assay to detect the binding of the PAc protein to salivary components in 96-well microtiter plates. Furthermore, a set of truncated PAc fragments was prepared by expressing subcloned fragments of *pac*, generated by the polymerase chain reaction (PCR), in an *Escherichia coli* expression vector. By use of these fragments, a saliva-interacting region of the PAc molecule was identified. In addition, fragments corresponding to the P region of PAc were shown to bind directly to the PAc protein.

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MATERIALS AND METHODS

Bacterial strains. S. mutans MT8148 was used as a representative strain of S. mutans serotype c. S. mutans TK18 is a recombinant strain that produces a large amount of PAc (13). S. mutans PAcEm-2 is a PAc-defective mutant (24).

rPAc and PAc fragments. Recombinant PAc (rPAc) protein was purified from the culture supernatants of transformant *S. mutans* TK18 by ammonium sulfate precipitation and chromatography on DEAE-cellulose (13). Truncated PAc and its polypeptide fragments were prepared as fusion proteins with β -galactosidase by use of an expression plas-

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TABLE 1. PCR primers used in the present study

Primer	Sequence ^a
pac 1L	5'-AAGGTACC-GATGAAACGACCACTAC-3'
pac 1R	5'-AAGTCGAC-TTGAACGGCTGCTAAAT-3'
pac 2L	5'-AAGGTACC-ACAGCTTATGAAGCTAA-3'
pac 2R	5'-AAGTCGAC-TTCGTATGCCTTTAACT-3'
pac 3L	5'-AAGGTACC-CCAGTTAAGTTAAAGGC-3'
pac 3R	5'-AAGTCGAC-ATACATAGTCCAGCGAG-3'
pac 4L	5'-AA <u>GGTACC</u> -GATAATGCCCTTCTCTC-3'
pac 4R	5'-AAGTCGAC-ATAATTTGGAGCTACTG-3'
pac 5L	5'-AAGGTACC-CCTAAAGTTACTAAGGA-3'
pac 5R	5'-AAGTCGAC-CTGAACAGCTAGTTTAA-3'
pac 6L	5'-AAGGTACC-GTTCATTTCCATTACTT-3'
pac 6R	5'-AAGTCGAC-GGCTGCTTCAAGATTAG-3'
pac 7L	5'-AAGGTACC-GGTGTTAGTGTGGATAA-3'
pac 7R	5'-AAGTCGAC-TGTTTCTTGAACAGAGC-3'

^a Restriction site sequences are underlined.

mid vector, pAX4a+ (Medac, Hamburg, Germany). Truncated *pac* gene fragments were amplified by the PCR (32). Primers used in the PCR are listed in Table 1. The amplified DNAs were digested with *KpnI* and *SalI* and ligated to the *KpnI-SalI*-digested plasmid vector. The ligated DNAs were transformed into *E. coli* NM522. Fusion proteins were extracted from whole cells of the transformants by sonication and purified by *p*-aminophenyl-1-thio- β -D-galactopyranoside (APTG) affinity chromatography in accordance with the instructions of the manufacturer (Medac). The amino acid positions in the PAc molecule of these rPAc fragments are summarized in Fig. 1. Fusion proteins were analyzed by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) (14).

Saliva. Paraffin-stimulated whole saliva was collected from five healthy donors into an ice-chilled plastic tube and clarified by centrifugation for 10 min at $12,000 \times g$. Whole saliva that was obtained from one donor (donor A; see Fig. 6) and that exhibited a relatively high ability to bind to the PAc protein was mainly used in the present study.

Sandwich assay. Ninety-six-well flat-bottom microtiter plates (Sumitomo Bakelite Co., Tokyo, Japan) were coated with rPAc $(1 \mu g)$, fusion PAc fragments $(5 \mu g)$, or whole cells of S. mutans (10 µg) in 100 µl of 50 mM carbonate buffer, (pH 9.6) overnight at 4°C. After three washes with phosphate-buffered saline containing 1 mM CaCl₂ (Ca-PBS), 1% (wt/vol) bovine serum albumin (BSA) in Ca-PBS was added, and the mixture was incubated overnight at 4°C. Excess BSA was removed by three washes with Ca-PBS, and then human whole saliva diluted with Ca-PBS was added (100 µl per well). After overnight incubation at 4°C, the plates were washed three times with Ca-PBS. Biotinylated rPAc (5 μ g/ml) in Ca-PBS containing 1% BSA was added (100 μ l per well), and the mixture was incubated overnight at 4°C. After three washes with Ca-PBS, alkaline phosphatase-conjugated streptavidin (BRL) diluted 1:1,000 with Ca-PBS containing 1% BSA was added (100 µl per well). The plates were incubated for 2 h at 20°C and washed with Ca-PBS, and the substrate, p-nitrophenylphosphate (1 mg/ml) in 0.1 M diethanolamine buffer (pH 9.6), was added (100 µl per well). After 1 h of incubation with the substrate, the optical density at 405 nm (OD_{405}) was measured with a microplate reader (MPR A4; Tosoh Co., Tokyo, Japan).

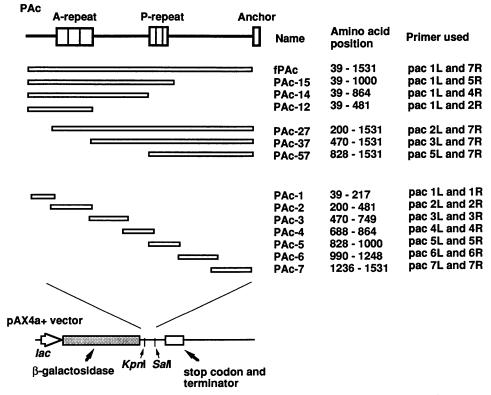


FIG. 1. Cloning and expression of PCR-amplified truncated *pac* gene fragments. PCR-amplified gene fragments were ligated to the *KpnI-SaII*-digested pAX4a+ vector. The nucleotide sequences of the primers are listed in Table 1. *E. coli* transformants harboring the chimeric plasmids expressed truncated PAc fragments as β -galactosidase fusion proteins.

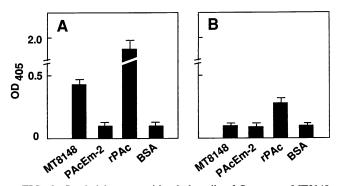


FIG. 2. Sandwich assay with whole cells of *S. mutans* MT8148 and PAcEm-2 and purified rPAc. Microtiter plates were coated with whole cells of *S. mutans* MT8148 (0.1 mg/ml), whole cells of PAc-defective *S. mutans* PAcEm-2 (0.1 mg/ml), purified rPAc (10 μ g/ml), or BSA (0.1 mg/ml). Human saliva from donor A was added to the wells of coated plates, and then biotinylated rPAc was added (A). For examination of the direct interaction between the coating substance and biotinylated rPAc, the treatment with human saliva was omitted (B). The OD₄₀₅ was measured after 1 h of incubation with the substrate. Each value represents the mean ± the standard deviation for triplicate assays.

Glucose, galactose, mannose, fucose, lactose, glucosamine, galactosamine, N-acetylglucosamine, N-acetylgalactosamine, and N-acetylneuraminic acid were tested for their ability to inhibit the binding of biotinylated rPAc to salivary components. Biotinylated rPAc (5 μ g/ml in Ca-PBS containing 1% BSA) was mixed with the sugar being tested (0.2 M), and the mixture was added to wells of plates and incubated overnight at 4°C as described above.

RESULTS

Sandwich assay. A sandwich assay was used to examine the interaction between the PAc protein and salivary components. Binding of whole cells of PAc-producing *S. mutans* MT8148 to salivary components was detected by this procedure. On the other hand, a PAc-defective mutant, *S. mutans* PAcEm-2, showed a negative reaction, indicating that binding in this assay is PAc specific (Fig. 2). A panel of sugars (glucose, galactose, mannose, fucose, lactose, glucosamine, galactosamine, *N*-acetylglucosamine, *N*-acetylgalactosamine, and *N*-acetylneuraminic acid) was assayed for inhibition of the binding of PAc to salivary components. No sugar significantly inhibited the binding of PAc to salivary components (data not shown).

Binding of rPAc fragments to salivary components. For determination of the saliva-binding region of the PAc molecule, we prepared 14 PAc fragments by using PCR technology and an expression vector, pAX4a + (Table 1 and Fig. 1). Fusion proteins expressed by recombinant *E. coli* were purified by APTG affinity chromatography and used for the sandwich assay. Figure 3 shows SDS-PAGE patterns of crude and purified fusion proteins. Proteins purified by APTG affinity chromatography showed two bands. The 116-kDa protein in the purified samples was a β -galactosidase nonfusion protein produced by host *E. coli*.

fPAc (residues 39 to 1531) exhibited a positive reaction in the sandwich assay, and β -galactosidase did not bind to salivary components (Fig. 4). The truncated PAc fragments, PAc-14 (residues 39 to 864) and PAc-15 (residues 39 to 1000) bound to salivary components, whereas the shorter fragment PAc-12 (residues 39 to 481) did not. On the other hand,

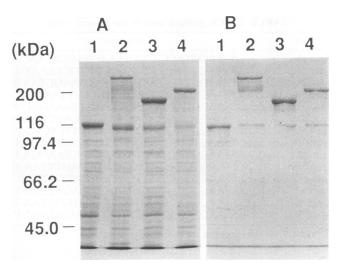


FIG. 3. SDS-PAGE (7.5% acrylamide gel) of *E. coli* crude extracts (A) and affinity-purified fusion proteins (B). Lanes: 1, *E. coli*(pAX4a+); 2, fPAc; 3, PAc-12; 4, PAc-14. We used 5 and 1 μ g of crude extract and purified fusion protein, respectively, per lane.

PAc-27 (residues 200 to 1531), PAc-37 (residues 470 to 1531), and PAc-57 (residues 828 to 1531) exhibited a positive reaction in the sandwich assay with saliva (Fig. 4A). However, biotinylated rPAc bound directly to PAc-37 and PAc-57 in the absence of saliva (Fig. 4B). Collectively, these results suggest that residues 39 to 864 of the PAc molecule are important in the binding of the PAc protein to salivary components and that the C-terminal region of the PAc molecule is capable of binding directly to the PAc protein.

We prepared shorter truncated PAc fragments, PAc-1 to PAc-7 (Fig. 1), and used them for the sandwich assay. None of these fragments, except for PAc-5 (residues 828 to 1000), bound to salivary components (Fig. 5A). PAc-5 bound directly to biotinylated rPAc (Fig. 5B).

The binding of fPAc, PAc-14, PAc-12, and β -galactosidase to saliva from five donors was examined with the sandwich assay (Fig. 6). fPAc and PAc-14 bound to salivary components from all the donors, but PAc-12 and β -galactosidase did not. These results suggest that the salivary components that the region including residues 39 to 864 reacts with are present in all saliva samples. However, the reactivity of the saliva differed with each donor (Fig. 6).

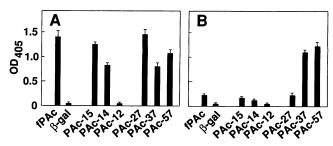


FIG. 4. Sandwich assay with C-terminal and N-terminal truncated PAc fragments. Microtiter plates were coated with fPAc, β -galactosidase (β -gal), PAc-15, PAc-14, PAc-12, PAc-27, PAc-37, or PAc-57 at a final concentration of 50 µg/ml. The sandwich assay was performed with (A) or without (B) human saliva. The OD₄₀₅ was measured after 1 h of incubation with the substrate. Each value represents the mean \pm the standard deviation for triplicate assays.

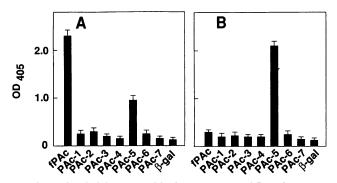


FIG. 5. Sandwich assay with shorter truncated PAc fragments. Microtiter plates were coated with fPAc, PAc-1, PAc-2, PAc-3, PAc-4, PAc-5, PAc-6, PAc-7, or β -galactosidase (β -gal) at a final concentration of 50 µg/ml. The sandwich assay was performed with (A) or without (B) human saliva. The OD₄₀₅ was measured after 1 h of incubation with the substrate. Each value represents the mean ± the standard deviation for triplicate assays.

DISCUSSION

The high-molecular-weight cell surface protein of S. mutans is known to function as a receptor for salivary agglutinin. Brady et al. (3) and Demuth et al. (5) have reported that purified PAc (P1, MSL-1) inhibits the salivary agglutinininduced aggregation of S. mutans.

A 205-kDa cell surface protein of *Streptococcus sanguis*, called SSP-5, is known to be a receptor for salivary agglutinin and exhibits calcium dependency for binding to salivary agglutinin (4). The binding of the SSP-5 protein to salivary agglutinin is inhibited by *N*-acetylneuraminic acid, suggesting that *N*-acetylneuraminic acid residues of salivary agglutinin are ligands for the SSP-5 protein (4, 5). On the other hand, the interaction between the *S. mutans* PAc protein and salivary agglutinin is poorly understood, although the molecular structure of the PAc protein is very similar to that of *S. sanguis* SSP-5 (5, 12, 24).

To understand the interaction between the PAc protein and salivary components at the molecular level, we prepared a series of truncated PAc fragments by using PCR technology and an expression vector, pAX4a+. In addition, we used a sandwich assay to determine the binding of the

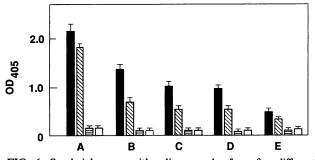


FIG. 6. Sandwich assay with saliva samples from five different human donors. Microtiter plates were coated with fPAc (\blacksquare), PAc-14 (\square), PAc-12 (\blacksquare), or β -galactosidase (\square) at a final concentration of 50 μ g/ml. The coated plates were allowed to react with saliva from each donor (A to E) and then with biotinylated rPAc. The bound biotinylated rPAc was detected with alkaline phosphatase-conjugated streptavidin and the substrate. The OD₄₀₅ was measured after 1 h of incubation with the substrate. Each value represents the mean \pm the standard deviation for triplicate assays.

truncated PAc fragments to salivary components. The present study suggested that residues 39 to 864 of the PAc molecule play an important role in the binding of the protein to salivary components (Fig. 4). Shorter fragments, such as PAc-12 (residues 39 to 481), PAc-3 (residues 470 to 749), and PAc-4 (residues 688 to 864), exhibited no binding activity. This finding may support the idea that a rather large region of the PAc molecule is required for reaction with salivary components. The N terminus of the PAc protein is clearly involved in binding to salivary components. However, it is difficult to rule out the possibility that the C-terminal region also contains a binding site, as the ability of PAc-14 (residues 39 to 864) to bind to salivary components was lower than that of PAc-15 (residues 39 to 1000) (Fig. 4). It is possible that structural changes result from truncation of the PAc peptide or its expression as a fusion protein with β -galactosidase and that such changes can influence the interpretation of results.

Salivary agglutinin-induced aggregation of S. mutans is reported to be inhibited by fucose and lactose (5). On the other hand, Russell and Mansson-Rahemtulla (29) showed that the binding of ¹²⁵I-PAc (I/II) to parotid gland salivacoated hydroxyapatite is inhibited by glucosamine, galactosamine, mannosamine, and N-acetylneuraminic acid. In the sandwich assay, no sugar significantly inhibited the binding of PAc to salivary components. Although glucosamine and galactosamine (0.2 M) weakly inhibited the binding of PAc, the degree of inhibition was not significant.

Brady et al. (3) used an array of 11 anti-PAc (P1) MAbs in a salivary agglutinin-induced aggregation inhibition assay and found that 6 MAbs markedly inhibited the salivary agglutinin-induced aggregation of S. mutans. Among these MAbs, one MAb recognized the truncated PAc fragment of residues 39 to 480, two MAbs recognized a fragment of residues 480 to 612, one MAb recognized a fragment of residues 612 to 1218, and two MAbs recognized a fragment of residues 1307 to 1561. Their findings suggest that the functional domain involved in the agglutinin-induced aggregation of S. mutans is located throughout the entire PAc (P1) molecule. On the other hand, the present study suggested that the saliva-interacting region of the PAc molecule might be the region between residues 39 to 864. We cannot compare simply our results with those reported by Brady et al. (3). We used whole saliva in the present study, but Brady et al. (3) used agglutinin purified from whole saliva. Moreover, the sandwich assay used in this study seems more like adherence than aggregation. It is possible that inhibitory antibodies react with a number of sites that all compose a functional domain when the PAc protein is properly configured. These sites may appear to be far apart on the amino acid sequence but actually may be closely related spatially.

The present study revealed that PAc-5 (residues 828 to 1000), corresponding to the P region of the PAc protein, bound to the whole PAc protein (Fig. 5). Moreover, binding to the P region was observed with PAc-57 (residues 828 to 1531) and PAc-37 (residues 470 to 1531) but not with PAc-27 (residues 200 to 1531) (Fig. 4B). These results suggest that binding to the P region is counteracted by the N-terminal portion of the PAc molecule. The biological significance of PAc protein binding to the P region is unknown. Our preliminary study, however, indicated that the native PAc protein formed aggregates in solution, and the molecular mass of the aggregates was estimated to be more than 1,000 kDa by gel filtration, suggesting that the PAc protein in solution may be a 5-mer or a 6-mer (unpublished data). Therefore, it is possible that the P region contributes to self-aggregation of the PAc protein. In addition, saliva

inhibited self-binding of the PAc-5 fragment (P region) (Fig. 5), suggesting that this region may also possess a functional domain for salivary binding.

S. mutans can be bound and/or agglutinated by various salivary molecules, such as secretory immunoglobulin A (27), β_2 -microglobulin (6), histidine-rich polypeptides (26), a 60-kDa glycoprotein (1), and high-molecular-weight glycoproteins (2, 7, 10). Russell and Mansson-Rahemtulla (29) reported that cell surface protein antigen PAc (I/II) of S. mutans binds to several salivary components, such as basic proline-rich salivary proteins with molecular weights of 28,000 and 38,000, lysozyme, and α -amylase. The present study showed that salivary components bound to a rather large N-terminal region of the PAc molecule. However, it is possible that these salivary components differ from one another with regard to their binding regions on the PAc molecule. Further work is needed to determine the precise region on the PAc molecule that is bound by each salivary component.

As suggested by Russell and Mansson-Rahemtulla (29), the interaction between the PAc protein and salivary components may be of a low affinity and a weak specificity. However, the sandwich assay developed in the present study seemed to be useful and sensitive enough for studying the interaction between the PAc protein and salivary components. In addition, the truncated PAc fragments may be very useful for characterizing the biological and immunological properties of the PAc protein of *S. mutans*.

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ADDENDUM

After submission of this paper, a paper by Crowley et al. (3a) appeared. Their results indicate that the A-region of the PAc (P1) molecule interacts directly with salivary agglutinin.

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