An In Vitro Microbial-Caries Model Used to Study the Efficacy of Antibodies to Streptococcus mutans Surface Proteins in Preventing Dental Caries

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The first step for a pathogenic bacterium to initiate infection is via attachment (i.e., through surface determinants) to a suitable receptor. An in vitro microbial artificial-mouth model was used to test the efficacy of polyclonal antibodies to Streptococcus mutans cell surface proteins (CsAb) and a cell surface 59-kDa protein (59Ab) in preventing S. mutans colonization and carious lesion formation. In study 1, groups of 12 human teeth specimens were inoculated with S. mutans, which were incubated with different concentrations of CsAb (A1 [positive control], sterile saline, no antibody; A2, 0.007 mg of antibody protein/ml; and A3, 0.7 mg of antibody protein/ml) for 1 h at 37°C. The negative control group (B1) was not infected and was incubated with Trypticase soy broth (TSB) without dextrose supplemented with 5% sucrose (TSBS). In study 2, the same study design was used except that 59Ab was used instead of CsAb, normal rabbit serum was used in the positive control group (A1), and TSB supplemented with 1% glucose was used as the nutrient to control sucrose-dependent colonization. All groups were exposed for 4 days to circulating cycles of TSBS and TSB (study 1 and study 2, respectively; 30 min each, three times per day) and a mineral washing solution (21 h per day). Prior to each nutrient cycle, 1 ml of the appropriate CsAb or 59Ab solution was administered to each group and allowed to mix for 30 min before cycling was resumed. Data obtained by confocal laser scanning microscopy demonstrated the presence of a significantly smaller (P < 0.05) lesion area and a smaller total lesion fluorescence in group A3 than in group A1 for both studies. In study 1, group A2 had significantly smaller values than A1 for lesion depth and area. There were no significant differences between groups A2 and A3 for lesion area or between groups A1 and A2 for total lesion fluorescence. In study 2, there were no significant differences among groups A1 and A2 for lesion depth or between groups A2 and A3 for all of the parameters studied. In both studies, there were no significant differences between S. mutans plaque CFU numbers among any of the groups. These studies demonstrated the efficacy of CsAb and 59Ab in reducing primary caries development in this model, although the underlying mechanism remains unclear.

Streptococcus mutans has been identified as the major etiological agent in human dental caries and comprises a significant proportion of the oral streptococci in carious lesions (10). It has been suggested that surface antigens such as antigen I/II or P1 participate in sucrose-independent colonization of tooth surfaces (3, 8), while glucosyltransferase and glucan-binding proteins (GBP) may be responsible for the sucrose-dependent colonization. All groups were exposed for 4 days to circulating cycles of TSBS and TSB (study 1 and study 2, respectively; 30 min each, three times per day) and a mineral washing solution (21 h per day). Prior to each nutrient cycle, 1 ml of the appropriate CsAb or 59Ab solution was administered to each group and allowed to mix for 30 min before cycling was resumed. Data obtained by confocal laser scanning microscopy demonstrated the presence of a significantly smaller (P < 0.05) lesion area and a smaller total lesion fluorescence in group A3 than in group A1 for both studies. In study 1, group A2 had significantly smaller values than A1 for lesion depth and area. There were no significant differences between groups A2 and A3 for lesion area or between groups A1 and A2 for total lesion fluorescence. In study 2, there were no significant differences among groups A1 and A2 for lesion depth or between groups A2 and A3 for all of the parameters studied. In both studies, there were no significant differences between S. mutans plaque CFU numbers among any of the groups. These studies demonstrated the efficacy of CsAb and 59Ab in reducing primary caries development in this model, although the underlying mechanism remains unclear.

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59Ab in the prevention of primary dental caries and to measure the extent of the developed primary carious lesions by using the quantifiable and reproducible method of confocal microscopy.

MATERIALS AND METHODS

CsAb and 59Ab preparation. (i) Cell surface protein preparation. This procedure was previously described by Fontana et al. (3). Briefly, S. mutans TH16 (serotype c) was grown in 9 liters of Todd-Hewitt broth (Difco Laboratories, Detroit, Mich.) supplemented with 1% glucose at 37°C in 5% CO2 and 95% air for 24 h. Cells from 9 liters of culture were harvested by centrifuging at 16,000 × g for 15 min at 4°C; washed once in buffer (20 mM Tris, 1 mM MgCl2, 0.02% NaN3; pH 6.8), and frozen as a pellet at −20°C overnight. A mixture of surface proteins from S. mutans were isolated by using a shearing technique. Frozen cells were thawed, suspended in buffer, and blended in a Waring blender for two 1-min cycles at high speed. Intact cells and cell debris were removed by a slow centrifugation (16,000 × g, 4°C, 10 min), and the supernatant, containing the cell surface protein (Cs protein) preparation, was retained and centrifuged at 110,000 × g for 2 h. The resulting Cs protein pellet was resuspended in the same buffer and centrifuged a second time at 16,000 × g for 10 min to further remove cell debris and aggregated components. The supernatant containing the Cs preparation was divided into aliquots and frozen at −80°C until use.

(ii) 59-kDa protein isolation. In order to separate cell surface protein fractions, preparative gel electrophoresis (Prep Cell model 491; Bio-Rad Laboratories, Richmond, Calif.) was utilized. The resolving and stacking gels were composed of 10% and 3% acrylamide (National Diagnostics, Atlanta, Ga.), respectively. A concentrated Cs protein preparation (2 ml, 1 mg/ml) from S. mutans A32-2 (serotype c) was added to an equal volume of sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) sample buffer, boiled for 7 min, placed on a 6-cm column, and subjected to 12 W of continuous power. The proteins were electrophoresed (150 V for approximately 1 h) by using reducing protein were electrophoresed (150 V for approximately 1 h) by using reducing gel electrophoresis after staining with Coomassie brilliant blue. The fractions of interest were pooled, collected and analyzed for molecular size and purity by gel electrophoresis after approximately 59 to 65 kDa (termed here the 59-kDa protein). The proteins were previously determined by immunoblots of Cs protein to have a molecular size of approximately 59 to 65 kDa (termed here the 59-kDa protein). The proteins were collected and analyzed for molecular size and purity by gel electrophoresis after staining with Coomassie brilliant blue. The fractions of interest were pooled, passed through an affinity column that removes SDS (Extracti-Gel; Pierce, Rockford, Ill.), and stored at −80°C.

(iii) CsAb preparation. Three New Zealand White rabbits were immunized with the prepared S. mutans Cs protein preparation (0.377 mg of protein/ml) by using the RIBI adjuvant system as suggested by the manufacturer (RIBI Immunochrom Research, Inc., Hamilton, Mont.). Injection of the protein preparation and RIBI adjuvant was done on day 0 and boosted on day 28. A total dose per animal of 1.0 ml (0.377 mg of protein/ml) was administered at each time period as follows: 0.3 ml intradermally (0.05 ml in each of six different sites), 0.4 ml intramuscularly (0.2 ml into each hind leg), 0.1 ml subcutaneously (in the neck region), and 0.2 ml intraperitoneally. Blood was collected by cardiac puncture on day 45, and serum was separated from the clot by centrifugation (5,000 × g, 10 min) and then stored at −20°C until use. The sera from the three rabbits were pooled and used as the antibody (CsAb) source for study 1. The ELISA absorbance values of the pooled sera were as follows: for a serum dilution of 1:1,000 the optical density at 490 nm (OD490) was 0.248 ± 0.043, for a serum dilution of 1:5,000 OD490 was 0.095 ± 0.010; and for a serum dilution of 1:10,000 the OD490 = 0.056 ± 0.008. All animal studies received Institutional Animal Care and Use Committee approval.

(iv) 59Ab preparation. Rat antisera to the 59-kDa surface protein were obtained from eight animals, each immunized with 5 mg of protein/ml incorporated into the RIBI adjuvant system (RIBI Immunochrom Research). Preparations were injected with 0.2 ml subcutaneously in each of two sites (in the upper back of the animals) and 0.1 ml intraperitoneally twice, 21 days apart, and blood was collected 7 days after the last injection. The blood was allowed to clot, and serum was obtained and frozen at −20°C until used. The sera from the eight rats were pooled and used as the antibody (59Ab) source for study 2. The ELISA absorbance values of the pooled sera were as follows: for a serum dilution of 1:2 the OD490 was 1.038 ± 0.314; for a serum dilution of 1:4 the OD490 was 0.927 ± 0.002; and for a serum dilution of 1:8 the OD490 was 0.635 ± 0.042.

(v) Electrophoretic techniques. To confirm CsAb and 59Ab antibody specificity to cell surface components, the Cs-enriched preparation and the 59-kDa protein were electrophoresed (150 V for approximately 1 h) by using reducing SDS–10% PAGE (National Diagnostics). Molecular size standards were included in the gel (Rainbow colored protein molecular size markers; Amersham, Arlington Heights, Ill.). Proteins were transferred electrophoretically (70 V for 2 h) to nitrocellulose paper for immunoblotting. The blots were blocked overnight with washing buffer (Trizma base; NaCl; Tween 20, pH 7.4) containing 0.5 ml of glutaraldehyde. The blots were probed with rabbit anti-S. mutans surface protein-enriched serum (CsAb; Fig. 1) and rat anti-59 kDa surface protein (59Ab) at a 1:1,000 dilution. Proteins which reacted with antibody were visualized on nitrocellulose by alkaline phosphatase-labeled anti-rabbit or anti-rat immunoglobulin G (IgG) heavy-chain-specific antibody (Sigma Chemical Company, St. Louis, Mo.) and nitroblue tetrazolium–5-bromo-4-chloro-3-indolylphosphate (NBT-BCIP; Bio-Rad). Molecular size was determined by comparison to protein standards by using an Ultrascan XL laser densitometer and GelScan XL software (Pharmacia LKB Biotechnology, Uppsala, Sweden).

The immunoblot of the S. mutans Cs protein preparation probed with CsAb demonstrated bands at approximately 190, 157, 65, 59, and 40 to 46 kDa (Fig. 1A), a result similar to that described previously (3). The major immunodominant band, whose role is currently being investigated, was at 59 kDa. This has been shown (unpublished data) to be distinct from the 59-kDa GBP (14). The minor bands seen at 190 and 157 kDa have been shown previously to be P1 and GTF, respectively (12). The immunoblot of the 59-kDa protein probed with CsAb demonstrated a single band at 59 to 65 kDa (Fig. 1B), supporting the purity of the isolated protein. Also, the 59Ab immunoblot demonstrated only one band at 59 to 65 kDa and has been identified to have amylase-binding properties (data not shown).
TABLE 1. Confocal microscopy measurements of primary carious lesions after treatment with CsAb (study 1)*

| Group | Inoculum | CsAbab | Area (μm², 10⁻¹) | Total fluorescence (10⁻¹µm) | Depth (µm) |
|-------|----------|--------|----------------|--------------------------|-----------|
| A1    | S. mutans TH16  | - | 20.4 ± 1.0 | 28.1 ± 4.1 | 67.0 ± 1.9 |
| A2    | S. mutans TH16 + (1:100) | 16.9 ± 0.8 | 28.4 ± 2.2 | 58.2 ± 2.8 |
| A3    | S. mutans TH16 + (Und) | 14.2 ± 0.9 | 13.4 ± 2.1 | 44.9 ± 3.2 |
| B1    | TSBS       |     | 0             | 0                        | 0            |
| Baseline |         |     | 0             | 0                        | 0            |

* CsAb, rabbit antisem to S. mutans TH16 Cs protein preparation (the dilution is shown in parentheses [Und, undiluted]). Group A1 received sterile saline. Values for area, total fluorescence, and depth are the mean ± the standard error of the mean. (n = 12 for groups A1, A2, A3, and B1; n = 8 for baseline values). Group A1 received the antibody solution in saline (8.78 g of NaCl/liter of deionized water) was used instead of CsAb in the positive control group in study 1. Normal rabbit serum (undiluted) was used instead of 59Ab in the positive control A1 group in study 2. After inoculation, the specimens were incubated for 2 h at 37°C to allow the control bacteria or antibody-treated bacteria to implant on the teeth. Each group of specimens was then placed in a separate caries-forming vessel and attached to the MW, TSBS or TSBS, and drainage container bottles.

In vitro microbial caries model experiments. (i) General experimental design. For each study, four groups of 12 human teeth specimens per group were treated for a 4-day test period in an in vitro microbial oral-microenvironment model. The groups differed from each other in the presence (A1, A2, and A3) or absence (B1, negative control) of S. mutans (serotype c) and in the concentration of antibody used (Tables 1 and 2). Two studies were conducted. In study 1, the effect of CsAb on bacterial adhesion and caries development was tested. In study 2, the effect of 59Ab was assessed.

(ii) Specimen preparation. Enamel specimens (3 mm in diameter) were drilled from extracted sound, human, lower permanent incisors which had been obtained from oral surgeons and sterilized by soaking in 3% buffered (neutral) formalin since the time of extraction. Each specimen was mounted on a polycrystalline rod by using methyl methacrylate resin. The specimens were first ground by using 600-grade silicon carbide paper to remove approximately 50 µm of the surface and then polished to a high luster with Gamma Alumina (0.05 µm) by standard methods. The specimens were then randomly assigned to test groups, with each group initially composed of 14 specimens. All specimens were sterilized with ethylene oxide gas. Two specimens from each group were randomly chosen before treatment and examined to obtain baseline confocal microscopy data. The 12 specimens that remained in each group were secured in a caries-forming vessel by gluing the ends of their plexiglass rods to a round plexiglass base that fit in the bottom of the vessel.

(iii) Treatment regimen and circulating fluids. Trypsinase soy broth without dextrose (Difco Laboratories) supplemented with 5% sucrose (TSBS) was used as the bacterial nutrient broth for study 1. For each group of specimens, 1 ml of the appropriate antibody solution was administered to each group by injection, followed by washing with 5 ml of sterile saline (8.78 g of NaCl/liter of deionized water). Antibody solutions were allowed to mix with fluid (MW) in the caries vessels by stirring for 30 min before the cycling was resumed.

(iv) Experimental setup. All of the media and model components, except for the enamel specimens, were autoclaved at 121°C for 20 min prior to the initiation of each experiment. For both studies, each group of 12 specimens was placed in a caries-forming vessel (125-ml Pyrex slow speed stirring vessel; Fisher). All caries-forming vessels were placed on an electric stirrer inside an incubator at 37°C under aerobic conditions. Each caries vessel had three inlets, one for TSBS or TSBS, one for MW, and one for injection of the antibody and one outlet for drainage tubing. The drainage tubing ended flush in a drainage container, which was also placed inside the incubator. Drainage of fluid from each caries vessel was maintained at 0.7 ml/min by a peristaltic pump.

(v) Preparation of bacterial inoculum and inoculation procedures. Each specimen in groups A1, A2, and A3 was inoculated by use of a micropipette with 20 ul of washed, overnight (16 h), stationary-phase cells of S. mutans TH16 (serotype c) for study 1 and S. mutans A32-2 (serotype c) for study 2, resuspended in TBS and TSBS (study 1 and study 2, respectively) to an OD₅₆₅ of 0.5. TSBS or TSB only was added to specimens in the negative control B1 groups. Prior to inoculation, filter (0.2 µm [pore size])-sterilized polyclonal antibodies prepared against S. mutans surface proteins (CsAb in study 1; 59Ab in study 2) were incubated in equal amounts (0.5 ml of each), at the appropriate concentration (Tables 1 and 2), with the S. mutans inoculum for 1 h at 37°C. Sterile saline (8.78 g of NaCl/liter of deionized water) was used instead of CsAb in the positive control group in study 1. Normal rabbit serum (undiluted) was used instead of 59Ab in the positive control A1 group in study 2. After inoculation, the specimens were incubated for 2 h at 37°C to allow the control bacteria or antibody-treated bacteria to implant on the teeth. Each group of specimens was then placed in a separate caries-forming vessel and attached to the MW, TSBS or TSBS, and drainage container bottles.

(vi) Monitoring of specimens. The following parameters were measured in the supply and drainage containers fluid at the beginning and at the end of the 4-day test periods to monitor the absence of contamination and the viability of the inoculum: (i) pH; (ii) S. mutans viability (by plating on mitis salivarius agar [Difco] supplemented with 20% sucrose and 200 IU of bacitracin per liter); and (iii) bacterial contamination (by plating on Trypticae soy agar [Difco]). Plates were inoculated at 37°C in 5% CO₂-95% air for 3 days.

In order to quantitate the bacteria adhered to the teeth at the end of the test periods, the specimens (two in study 1 and three in study 2) from each group were isolated individually in 5 ml of each. Each specimen was vortexed (20 s) and sonicated (20 s) until all visible dental plaque was displaced from the surface of the tooth. All samples were double plated on mitis salivarius and Trypticae soy agar.

For both studies, the two MW bottles and all four TSBS bottles maintained a neutral pH (approximately 7.0) at the beginning and at the end of the test period, indicating no contamination of the supply vessels. The pH of the negative control group caries vessel (B1) in both studies remained neutral throughout the experiment indicating a lack of contamination. In addition, control caries-forming vessel, MW bottles, and TSBS bottles remained sterile throughout the treatment periods, and no contamination was observed in the experimental or control groups.

(vii) Evaluation of tooth samples. After termination of each study, a casting resin (Meyer Plastics, Indianapolis, Ind.) was applied to the top of each specimen and allowed to polymerize. This was done in order to protect the surface of the specimen during the cutting procedure. Specimens were sectioned in half by using a Surker-Taylor hard-tissue microtome. One-half of each specimen was stained with a 0.1 mM solution of rhodamine B (Aldrich Chemical Co., Milwaukee, Wis.) overnight, with no further rinsing. The stained surface was analyzed by using a laser scanning confocal microscope (Odyssey; Noran Instruments, Inc., Middleton, Wis.) to determine the extent of the lesion (S). One area, 300 µm in length, was scanned for each specimen. The confocal microscope permits nonimaging imaging of subsurface tissue structures, including the enamel and dentin. Areas were scanned planar parallel to the transversal cut surface of the specimen and perpendicular to the natural surface of the tooth. The areas were not scanned directly at the cut surface of the specimen because of concern regarding the smear layer created during the cutting procedure. Areas were scanned at between 10 and 50 µm below the cut surface with the confocal microscope by using Image I (version 4.14.C) software (Universal Image Corp., West Chester, Pa.). After being brought into focus (with a 10X Nikon objective [numerical aperture, 0.25] and 10X eyepiece), the specimens were illuminated with an argon laser by using a 529-nm excitation wavelength. Confocal slits were set at a 100-nm width with a 550-nm long-pass filter, and the argon laser intensity was set at 100%. The argon laser used had a power intensity of 1.23 mW per scanned point in the specimen. After examination of all specimens, confocal settings (contrast and brightness) were maximized and held constant. For the collection of images, the samples were frame averaged by using 128 frames per image. The parameters that were measured included the area of the fluorescent lesion, the total lesion fluorescence, and the depth of the lesion.

TABLE 2. Confocal microscopy measurements of primary carious lesions after treatment with 59Ab (study 2)*

| Group | Inoculum | 59Ab | Area (µm², 10⁻¹) | Total fluorescence (10⁻¹µm) | Depth (µm) |
|-------|----------|------|----------------|--------------------------|-----------|
| A1    | S. mutans A32-2 | - | 8.0 ± 1.7 | 7.8 ± 2.2 | 32.0 ± 6.5 |
| A2    | S. mutans A32-2 + (1:100) | 4.4 ± 1.3 | 3.3 ± 1.4 | 27.0 ± 8.2 |
| A3    | S. mutans A32-2 + (Und) | 2.1 ± 1.4 | 1.9 ± 1.6 | 20.2 ± 5.5 |
| B1    | TSBS     | 0    | 0             | 0                        | 0            |

* 59Ab, rat antisem to S. mutans A32-2 59-kDa protein (the dilution is shown in parentheses [Und, undiluted]). Group A1 received rabbit normal serum. Values for area, total fluorescence, and depth are the mean ± the standard error of the mean. (n = 7 for group A1; n = 9 for groups A2, A3, and B1). Groups A2 and A3 were not significantly different (P > 0.05), as determined by Tukey's procedure.
RESULTS

Data analysis. Means and variances were calculated for each measured parameter (area of the lesion, total lesion fluorescence, lesion depth, and numbers of plaque S. mutans CFU) and treatment group. These data were analyzed by using a single factor analysis of variance model. Where a significant effect was detected (α=0.05), multiple comparisons were conducted by using Tukey’s procedure.

DISCUSSION

Our laboratory has been extensively involved in establishing the role of S. mutans fimbrial components in the adherence and colonization of the tooth surface by this bacterium and in testing whether antibodies against S. mutans surface proteins, enriched in fimbria components, reduce the adherence of S. mutans to the tooth surface, thereby inhibiting the development of primary dental caries (3, 12, 13). Caries-free (CF) adult individuals have higher levels of salivary IgA antibodies to an enriched-fimbriae preparation of S. mutans than caries-active (CA) individuals (3). These results suggest that CF subjects may be protected immunologically from dental caries in part by salivary IgA antibody against S. mutans fimbrial-cell surface antigens. Perrone et al. (12) demonstrated by using immunoblot analysis and ELISA techniques with antibody to the cell surface, fimbria-enriched preparations, GTF, and P1 antigen that the levels of fimbria components, GTF, and P1 antigens were higher in cell surface fimbria-enriched preparations from S. mutans isolates from CA subjects than in preparations from CF individuals. These results suggest that the differences between the composition of S. mutans cell surface fimbria-enriched preparations in isolates from CA and CF subjects may play an important role in the virulence of this microorganism in dental caries. While S. mutans TH16 was the laboratory strain chosen for study 1, S. mutans A32-2 (a CA isolate) was chosen as the inoculum for study 2 based partly on the findings of Perrone et al. (12). Our laboratory has also reported that a 52-kDa salivary protein, identified as amylase, is the major S. mutans fimbria-enriched cell surface preparation binding protein in whole saliva (13). Preliminary studies
were conducted to address the need to add a salivary pellicle to the teeth prior to bacterial inoculation in the microbial-caries model. Results showed no significant differences in bacterial attachment and caries scores between surfaces with and those without a salivary pellicle (4). Therefore, to minimize the number of variables in these initial studies, it was decided not to use a pellicle. Of course, the fact that we did not use a pure preparation in study 1, but rather a mixture of most major cell surface associated antigens of \textit{S. mutans}, indicated that although most of the CsAb was directed against a cell surface component (59 kDa), there was some small reactivity with other surface antigens such as P1 and GTF that could in part be responsible for the elicited caries protection. Therefore, a follow-up study (study 2) was conducted with this same model and polyclonal antibodies elicited in rats to the 59-kDa protein to study the degree of protection obtained by targeting the 59-kDa surface protein, and the results demonstrated a similar degree of caries protection, supporting the importance of the 59-kDa protein in caries development. Studies to clone this protein are currently under way (6).

The goals in the prevention of colonization of specific pathogenic bacteria include long-lasting protection conferred by an appropriate vaccine. An intranasal vaccination study was conducted with the same cell surface preparation as the one used to prepare CsAb as the immunogen (2). The results demonstrated that the elicited antibodies were mainly elicited in both saliva and serum against a 59-kDa protein and protected against smooth surface caries. However, there were no differences in the ability of the antibody to inhibit bacterial colonization. The in vitro caries model described here is a fast, simple, economical, and novel approach for testing the effect of \textit{S. mutans} anti-surface protein antibodies or other antimicrobials in caries prevention. Bacterial caries systems, where the flora is controlled by in vitro environmental and nutritional conditions, provide a controlled means for studying complex ecosystems, such as dental plaque and its effect on the development of dental caries. The use of a bacterial artificial caries system permits more clinically relevant in vitro investigations of primary caries etiology and prevention, since it links bacteria with the resulting demineralization of the tooth. By providing CsAb or 59Ab in the system, the production of salivary anti-bodies that occurs in caries-free subjects or that would result as a consequence of orally immunizing an animal or individual with \textit{S. mutans} cell surface fimbria-enriched preparations was simulated. Since normal rabbit serum had no enzyme-linked immunosorbent assay reactivity to the enriched cell surface preparation, a saline solution was used as the negative control for study 1. However, the possibility that nonantibody components (i.e., albumin) may interfere with \textit{S. mutans} colonization or metabolic activity was directly addressed in study 2, in which normal rabbit (preimmune) serum was used as a negative control. Unfortunately, CsAb and 59Ab could not be delivered continuously to the system as would occur by saliva in vivo. Also, since the drainage of liquid occurred from the top of the caries vessels and since the antibodies caused bacterial agglutination, the bacteria were not completely eliminated (they would be swallowed in vivo) but rather partially accumulated in the bottom of the vessel and may have maintained metabolic activity. This may have contributed to the low pH and relatively higher caries scores (compared to B1) observed in the A2 and A3 groups. This issue will be addressed in follow-up studies, since the caries vessels have been modified so that the outlet is located in the bottom of the vessel.

Our hypothesis was that the smaller lesions observed in the antibody-treated specimens might be due in part to a reduction in \textit{S. mutans} adherence to the tooth surfaces and to reduced plaque formation. However, based on the data of these in vitro studies as well as results from the rat immunization study (2), the mechanism for the elicited caries protection is not clear. The data demonstrated that although CsAb-treated specimens in group A3 (study 1) showed a trend toward a decreased number of \textit{S. mutans} adhered to the teeth surface, the results were not statistically significant, a result probably due to the small sample sizes used for bacterial analysis. However, these CsAb antibodies could have also affected bacterial metabolism, leading to a less-cariogenic \textit{S. mutans} plaque. It must be noted that the results presented here for study 1 are representative of two identical experiments; therefore, the studies are reproducible. Also, the fact that sucrose was provided in the diet probably supported the action of GTF in inducing, mainly, a glucan-dependent plaque. This may partially explain why significant differences in bacterial numbers were not observed among the treatment groups in study 1, in which an effect of antibody on sucrose-independent attachment was anticipated. Therefore, study 2 was conducted without the use of sucrose in the nutrient medium. This change decreased the amount of caries obtained in study 2 compared to study 1, but it still did not result in significant differences in the numbers of plaque bacteria. Despite this, however, our results demonstrated the application of an in vitro caries model in testing the efficacy of antibodies to \textit{S. mutans} surface proteins in decreasing the level of caries development.

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